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Cloudera Data Science Workbench is a secure, self-service enterprise data science platform that lets data scientists manage their own analytics pipelines, thus accelerating machine learning projects from exploration to production. It allows data scientists to bring their existing skills and tools, such as R, Python, and Scala, to securely run computations on data in Hadoop clusters. It enables data science teams to use their preferred data science packages to run experiments with on-demand access to compute resources. Models can be trained, deployed, and managed centrally for increased agility and compliance.

Built for the enterprise, Cloudera Data Science Workbench includes direct integration with the Cloudera platform for a complete machine learning workflow that supports collaborative development, and can run both in the public cloud and on-premises.

**Demo** - Watch this video for a quick 3 minute demo of Cloudera Data Science Workbench: [CDSW Quickstart](#)

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### Typical Machine Learning Project Workflow

Machine learning is a discipline that uses computer algorithms to extract useful knowledge from data. There are many different types of machine learning algorithms, and each one works differently. In general however, machine learning algorithms begin with an initial hypothetical model, determine how well this model fits a set of data, and then work on improving the model iteratively. This training process continues until the algorithm can find no additional improvements, or until the user stops the process.

A typical machine learning project will include the following high-level steps that will transform a loose data hypothesis into a model that serves predictions.

1. Explore and experiment with and display findings of data
2. Deploy automated pipelines of analytics workloads
3. Train and evaluate models
4. Deploy models as REST APIs to serve predictions

With Cloudera Data Science Workbench, you can deploy the complete lifecycle of a machine learning project from research to deployment.
Core Capabilities of Cloudera Data Science Workbench

<table>
<thead>
<tr>
<th>For Data Scientists</th>
</tr>
</thead>
<tbody>
<tr>
<td>Projects</td>
</tr>
<tr>
<td>Organize your data science efforts as isolated projects, which might include reusable code, configuration, artifacts, and libraries. Projects can also be connected to GitHub repositories for integrated version control and collaboration.</td>
</tr>
<tr>
<td>Workbench</td>
</tr>
<tr>
<td>A workbench for data scientists and data engineers that includes support for:</td>
</tr>
<tr>
<td>• Interactive user sessions with Python, R, and Scala through flexible and extensible engines.</td>
</tr>
<tr>
<td>• Project workspaces powered by Docker containers for control over environment configuration. You can install new packages or run command-line scripts directly from the built-in terminal.</td>
</tr>
<tr>
<td>• Distributing computations to your Cloudera Manager cluster using CDS 2.x Powered by Apache Spark and Apache Impala.</td>
</tr>
<tr>
<td>• Sharing, publishing, and collaboration of projects and results.</td>
</tr>
<tr>
<td>Jobs</td>
</tr>
<tr>
<td>Automate analytics workloads with a lightweight job and pipeline scheduling system that supports real-time monitoring, job history, and email alerts.</td>
</tr>
<tr>
<td>Batch Experiments</td>
</tr>
<tr>
<td>Demo - Experiments</td>
</tr>
<tr>
<td>Use batch jobs to train and compare versioned, reproducible models. With experiments, data scientists can:</td>
</tr>
<tr>
<td>• Create versioned snapshots of model code, dependencies, and any configuration parameters required to train the model.</td>
</tr>
<tr>
<td>• Build and execute each training run in an isolated container.</td>
</tr>
<tr>
<td>• Track model metrics, performance, and model artifacts as required.</td>
</tr>
<tr>
<td>Models</td>
</tr>
<tr>
<td>Demo - Model Deployment</td>
</tr>
<tr>
<td>Deploy and serve models as REST APIs. Data scientists can select a specific Python or R function within a project file to be deployed as a model, and Cloudera Data Science Workbench will:</td>
</tr>
<tr>
<td>• Create a snapshot of the model code, saved model parameters, and dependencies.</td>
</tr>
<tr>
<td>• Build an immutable executable container with the trained model and serving code.</td>
</tr>
<tr>
<td>• Deploy the model as a REST API along with a specified number of replicas, automatically load balanced.</td>
</tr>
<tr>
<td>• Save the built model container, along with metadata such as who built or deployed it.</td>
</tr>
<tr>
<td>• Allow data scientists to test and share the model</td>
</tr>
</tbody>
</table>
For IT Administrators

Native Support for the Cloudera Enterprise Data Hub

Direct integration with the Cloudera Enterprise Data Hub makes it easy for end users to interact with existing clusters, without having to bother IT or compromise on security. No additional setup is required. They can just start coding.

Enterprise Security

Cloudera Data Science Workbench can leverage your existing authentication systems such as SAML or LDAP/Active Directory. It also includes native support for Kerberized Hadoop clusters.

Native Spark 2 Support

Cloudera Data Science Workbench connects to existing Spark-on-YARN clusters with no setup required.

Flexible Deployment

Deploy on-premises or in the cloud (on IaaS) and scale capacity as workloads change.

Multitenancy Support

A single Cloudera Data Science Workbench deployment can support different business groups sharing common infrastructure without interfering with one another, or placing additional demands on IT.

Architecture Overview

Important: The rest of this documentation assumes you are familiar with CDH and Cloudera Manager. If not, make sure you read the documentation for CDH and Cloudera Manager before you proceed.

Cloudera Manager

Cloudera Manager is an end-to-end application used for managing CDH clusters. When a CDH service (such as Impala, Spark, etc.) is added to the cluster, Cloudera Manager configures cluster hosts with one or more functions, called roles. In a Cloudera Manager cluster, a gateway role is one that designates that a host should receive client configuration for a CDH service even though the host does not have any role instances for that service running on it. Gateway roles provide the configuration required for clients that want to access the CDH cluster. Hosts that are designated with gateway roles for CDH services are referred to as gateway hosts.

Cloudera Data Science Workbench runs on one or more dedicated gateway hosts on CDH clusters. Each of these hosts has the Cloudera Manager Agent installed on them. The Cloudera Management Agent ensures that Cloudera Data Science Workbench has the libraries and configuration necessary to securely access the CDH cluster.

Cloudera Data Science Workbench does not support running any other services on these gateway hosts. Each gateway host must be dedicated solely to Cloudera Data Science Workbench. This is because user workloads require dedicated CPU and memory, which might conflict with other services running on these hosts. Any workloads that you run on Cloudera Data Science Workbench hosts will have immediate secure access to the CDH cluster.

From the assigned gateway hosts, one will serve as the master host while others will serve as worker hosts.
Master Host

The master host keeps track of all critical persistent and stateful data within Cloudera Data Science Workbench. This data is stored at /var/lib/cdsw.

- **Project Files**

  Cloudera Data Science Workbench uses an NFS server to store project files. Project files can include user code, any libraries you install, and small data files. The master host provides a persistent filesystem which is exported to worker hosts using NFS. This filesystem allows users to install packages interactively and have their dependencies and code available on all Cloudera Data Science Workbench nodes without any need for synchronization. The files for all the projects are stored on the master host at /var/lib/cdsw/current/projects. When a job or session is launched, the project’s filesystem is mounted into an isolated Docker container at /home/cdsw.

- **Relational Database**

  The Cloudera Data Science Workbench uses a PostgreSQL database that runs within a container on the master host at /var/lib/cdsw/current/postgres-data.

- **Livelog**

  Cloudera Data Science Workbench allows users to work interactively with R, Python, and Scala from their browser and display results in realtime. This realtime state is stored in an internal database called Livelog, which stores data on the master host at /var/lib/cdsw/current/livelog. Users do not need to be connected to the server for results to be tracked or jobs to run.

Worker Hosts

While the master host stores the stateful components of the Cloudera Data Science Workbench, the worker hosts are transient. These can be added or removed as needed, which gives you flexibility with scaling the deployment. As the number of users and workloads increases, you can add more worker hosts to Cloudera Data Science Workbench over time.

**Note:** Worker hosts are not required for a fully-functional Cloudera Data Science Workbench deployment. For proof-of-concept deployments, you can deploy a 1-host cluster with just a Master host. The Master host can run user workloads just as a worker host can.

Even on multi-host deployments, the Master host doubles up to perform both functions: those of the Master outlined above, and those of a worker. Starting with version 1.4.3, multi-host deployments can be customized to reserve the Master only for internal processes while user workloads are run exclusively on workers. For details, see Reserving the Master Host for Internal CDSW Components on page 209.

Engines

Cloudera Data Science Workbench engines are responsible for running R, Python, and Scala code written by users and intermediating access to the CDH cluster. You can think of an engine as a virtual machine, customized to have all the necessary dependencies to access the CDH cluster while keeping each project’s environment entirely isolated. To ensure that every engine has access to the parcels and client configuration managed by the Cloudera Manager Agent, a number of folders are mounted from the host into the container environment. This includes the parcel path /opt/cloudera, client configuration, as well as the host’s JAVA_HOME.

For more details on basic concepts and terminology related to engines in Cloudera Data Science Workbench, see Cloudera Data Science Workbench Engines on page 169.

Docker and Kubernetes

Cloudera Data Science Workbench uses Docker containers to deliver application components and run isolated user workloads. On a per project basis, users can run R, Python, and Scala workloads with different versions of libraries and
system packages. CPU and memory are also isolated, ensuring reliable, scalable execution in a multi-tenant setting. Each Docker container running user workloads, also referred to as an engine, provides a visualized gateway with secure access to CDH cluster services such as HDFS, Spark 2, Hive, and Impala. CDH dependencies and client configuration, managed by Cloudera Manager, are mounted from the underlying gateway host. Workloads that leverage CDH services such as HDFS, Spark, Hive, and Impala are executed across the full CDH cluster.

To enable multiple users and concurrent access, Cloudera Data Science Workbench transparently subdivides and schedules containers across multiple hosts dedicated as gateway hosts. This scheduling is done using Kubernetes, a container orchestration system used internally by Cloudera Data Science Workbench. Neither Docker nor Kubernetes are directly exposed to end users, with users interacting with Cloudera Data Science Workbench through a web application.

Cloudera Data Science Workbench Web Application

The Cloudera Data Science Workbench web application is typically hosted on the master host, at http://cdsw.<your_domain>.com. The web application provides a rich GUI that allows you to create projects, collaborate with your team, run data science workloads, and easily share the results with your team. For a quick demonstration, either watch this video or read the Quickstart Guide.

You can log in to the web application either as a site administrator or a regular user. See the Administration and User Guides respectively for more details on what you can accomplish using the web application.

CDS 2.x Powered by Apache Spark

Important: The rest of this topic assumes you are familiar with Apache Spark and CDS 2.x Powered by Apache Spark. If not, make sure you read the CDS 2.x documentation before you proceed.

Apache Spark is a general purpose framework for distributed computing that offers high performance for both batch and stream processing. It exposes APIs for Java, Python, R, and Scala, as well as an interactive shell for you to run jobs.

Cloudera Data Science Workbench provides interactive and batch access to Spark 2. Connections are fully secure without additional configuration, with each user accessing Spark using their Kerberos principal. With a few extra lines of code, you can do anything in Cloudera Data Science Workbench that you might do in the Spark shell, as well as leverage all the benefits of the workbench. Your Spark applications will run in an isolated project workspace.

Cloudera Data Science Workbench’s interactive mode allows you to launch a Spark application and work iteratively in R, Python, or Scala, rather than the standard workflow of launching an application and waiting for it to complete to view the results. Because of its interactive nature, Cloudera Data Science Workbench works with Spark on YARN’s client mode, where the driver persists through the lifetime of the job and runs executors with full access to the CDH cluster resources. This architecture is illustrated the following figure:
More resources:

- [Documentation for CDS 2.x Powered by Apache Spark](#)
- [Apache Spark 2 upstream documentation](#)
Cloudera Data Science Workbench Release Notes

These release notes provide information on new features, fixed issues and incompatible changes for all generally-available (GA) versions of Cloudera Data Science Workbench. For the current known issues and limitations, see Known Issues and Limitations in Cloudera Data Science Workbench 1.5.x on page 42.

Cloudera Data Science Workbench 1.5.0

This section lists the release notes for Cloudera Data Science Workbench 1.5.0.

New Features and Changes in Cloudera Data Science Workbench 1.5.0

- **Cloudera Enterprise 6.1 Support**
  Cloudera Data Science Workbench is now supported with Cloudera Manager 6.1.x (and higher) and CDH 6.1.x (and higher). For details, see Cloudera Manager and CDH Requirements on page 52.

- **Cloudera Data Science Workbench on Hortonworks Data Platform (HDP)**
  Cloudera Data Science Workbench can now be deployed on HDP 2.6.5 and HDP 3.1.0. For an architecture overview and installation instructions, see Deploying Cloudera Data Science Workbench 1.5.x on Hortonworks Data Platform on page 85.

- **Security Enhancements**
  - **Allow Site Administrators to Enable/Disable Project Uploads and Downloads** - By default, all Cloudera Data Science Workbench users are allowed to upload and download files to/from a project. Version 1.5 introduces a new feature flag that allows site administrators to hide the UI features that let users upload and download project files.
    
    Note that this feature flag only removes the relevant features from the Cloudera Data Science Workbench UI. It does not disable the ability to upload and download files through the backend web API.
    
    For details on how to enable this feature, see Disabling Project File Uploads and Downloads on page 110.

- **OpenJDK Support**
  Cloudera Data Science Workbench now supports Open JDK 8 on Cloudera Enterprise 5.16.1 (and higher). For details, see Product Compatibility Matrix - Supported JDK Versions.

- **Engines**
  - **Base engine upgraded with a new version of R - 3.5.1 (Base Image v7)**
  - **Debugging Improvements** - Previously, engines and their associated logs were deleted immediately after an exit or a crash. With version 1.5, engines are now available for about 5 minutes after they have ended to allow you to collect the relevant logs.

    Additionally, when an engine exits with a non-zero status code, the last 50 lines from the engine's logs are now printed to the Workbench console. Note that a non-zero exit code and the presence of engine logs in the Workbench does not always imply a problem with the code. Events such as session timeouts and out-of-memory issues are also assigned non-zero exit codes and will display engine logs.

- **Installation and Upgrade**
  - **New Configuration Parameters** - Version 1.5 includes three new configuration parameters that can be used to specify the type of distribution you are running, the directory for the installed packages/parcels, and the path where Anaconda is installed (for HDP only).
    
    - DISTRO
Details and sample values for these properties have been added to the relevant installation topics for CDH and HDP.

- **DOCKER_TMPDIR changed to /var/lib/cdsw/tmp/docker** - Previously the Cloudera Data Science Workbench installer would temporarily decompress the base engine image file to the /var/lib/docker/tmp directory. Starting with version 1.5, the installer will use the /var/lib/cdsw/tmp/docker directory instead. Make sure you have an Application block device mounted to /var/lib/cdsw as recommended so that installation/upgrade can proceed without issues.

- **Improved Validation Checks** - Improved the validation checks run by the installer and the error messages that are displayed during the installation process. Cloudera Data Science Workbench now:
  - Checks that space is available on the root directory, the Application Block Device and the Docker Block Device(s).
  - Checks that DNS forward and reverse lookup works for the Cloudera Data Science Workbench Domain and Master IP address provided.
  - Displays better error messages for the `cdsw status` and `cdsw validate` commands for easier debugging.

- **Command Line**
  - `cdsw logs` - Previously, the `cdsw logs` command generated two log bundles - one in plaintext and one with sensitive information redacted. With version 1.5, the command now generates only a single bundle that has all the sensitive information redacted by default.

  To turn off redaction of log files for internal use, you can use the new `--skip-redaction` option as follows:

  ```
  cdsw logs --skip-redaction
  ```

- **Networking**
  - Cloudera Data Science Workbench now uses DNS hostnames (not IP addresses) for internal communication between components. As a result, the wildcard DNS hostname configured for Cloudera Data Science Workbench must now be resolvable from both, the CDSW cluster, and your browser.

  - Cloudera Data Science Workbench now enables IPv4 forwarding (`net.ipv4.conf.default.forwarding`) during the installation process.

### Engine Upgrade

Cloudera Data Science Workbench 1.5.0 (and later) ships **version 7** of the base engine image which includes the following versions of R and Python:

- **R**  - 3.5.1
- **Python**  - 2.7.11, 3.6.1

**Pre-installed Packages in Engine 7** - For details about the packages included in the base engine, see [Cloudera Data Science Workbench Engine Versions and Packaging](#) on page 187.

**Upgrade Projects to Use the Latest Base Engine Images** - Make sure you test and upgrade existing projects to Base Image v7 (Project Settings > Engine) to take advantage of the latest fixes.

**Note that this is a required step if you are upgrading to using Cloudera Data Science Workbench on CDH 6.**

The base engine image you use must be compatible with the version of CDH you are running. This is especially important if you are running workloads on Spark. Older base engines (v6 and lower) cannot support the latest versions of CDH 6. That is because these engines were configured to point to the Spark 2 parcel. However, on CDH 6 clusters, Spark
now packaged as a part of CDH 6 and the separate add-on Spark 2 parcel is no longer supported. If you want to run Spark workloads on CDH 6, you must upgrade your projects to base engine 7 (or higher).

Table 1: CDSW Base Engine Compatibility for Spark Workloads on CDH 5 and CDH 6

<table>
<thead>
<tr>
<th>Base Engine Versions</th>
<th>CDH 5</th>
<th>CDH 6</th>
</tr>
</thead>
<tbody>
<tr>
<td>Base engines 6 (and lower)</td>
<td>Yes</td>
<td>No</td>
</tr>
<tr>
<td>Base engines 7 (and higher)</td>
<td>Yes</td>
<td>Yes</td>
</tr>
</tbody>
</table>

Incompatible Changes in Cloudera Data Science Workbench 1.5.0

Deprecated Property - CDH Parcel Directory

The CDH parcel directory property is no longer available in the Site Administration panel at Admin > Engines. Depending on your deployment, use one of the following ways to configure this property:

- **CSD deployments**: If you are using the default parcel directory, /opt/cloudera/parcels, no action is required. If you want to use a custom location for the parcel directory, configure this in Cloudera Manager as documented here.
- **RPM deployments**: If you are using the default parcel directory, /opt/cloudera/parcels, no action is required. If you want to specify a custom location for the parcel directory, configure the DISTRO_DIR property in the cdsw.conf file on both master and worker hosts. Run cdsw restart after you make this change.

Issues Fixed in Cloudera Data Science Workbench 1.5.0

- Fixed an issue with RPM installations where NO_PROXY settings were being ignored.
  Cloudera Bug: DSE-4444
- Fixed an issue where CDSW would not start because of IP issues with web pods. Version 1.5 fixes this by enabling IPv4 forwarding at startup.
  Cloudera Bug: DSE-4609
- Fixed an issue where engines would get deleted immediately after an exit/crash and engine logs did not persist which made it difficult to debug issues with crashes or auto-restarts.
  Cloudera Bug: DSE-4008, DSE-4417
- Fixed intermittent issues with starting and stopping Cloudera Data Science Workbench on CSD deployments.
  Cloudera Bug: DSE-4426, DSE-4829
- Fixed an issue where Cloudera Data Science Workbench was reporting incorrect file sizes for files larger than 2 MB.
  Cloudera Bug: DSE-4531, DSE-4532
- Fixed an issue where the Run New Experiment dialog box did not include the file selector and the Script name had to be typed in manually.
  Cloudera Bug: DSE-3650
- Fixed an issue where underlying Kubernetes processes were running out of resources leading to Out of Memory (OOM) errors. Cloudera Data Science Workbench now reserves compute resources for Kubernetes components.
  Cloudera Bug: DSE-4896, DSE-5001
- Fixed an issue where the PYSPARK3_PYTHON environment variable was not working as expected for Python 3 workloads.
  Cloudera Bug: DSE-4329
• Fixed an issue where Docker commands would fail on Cloudera Data Science Workbench engines that are not available locally (such as custom engine images) when an HTTP/HTTPS proxy was in use.
  Cloudera Bug: DSE-4427

• Fixed an issue where installation of the XML package would fail in the R kernel.
  Cloudera Bug: DSE-2201

Known Issues and Limitations in Cloudera Data Science Workbench 1.5.0

For a complete list of the current known issues and limitations in Cloudera Data Science Workbench 1.5.x, see Known Issues and Limitations in Cloudera Data Science Workbench 1.5.x on page 42.

Cloudera Data Science Workbench 1.4.3

This section lists the release notes for Cloudera Data Science Workbench 1.4.3.

New Features and Changes in Cloudera Data Science Workbench 1.4.3

• Reserve Master Host for Internal Application Components

  Cloudera Data Science Workbench now allows you to reserve the master host for running internal application components and services such as Livelog, the PostgreSQL database, and so on, while user workloads run exclusively on worker hosts.

  By default, the master host runs both, user workloads as well as the application's internal services. However, depending on the size of your CDSW deployment and the number of workloads running at any given time, it's possible that user workloads might dominate resources on the master host. Enabling this feature will ensure that CDSW's application components always have access to the resources they need on the master host and are not adversely affected by user workloads.

  **Important:** This feature only applies to deployments with more than one Cloudera Data Science Workbench host. Enabling this feature on single-host deployments will leave Cloudera Data Science Workbench incapable of scheduling any workloads.

  For details on how to enable this feature, see Reserving the Master Host for Internal CDSW Components on page 209.

• Allow Only Session Creators to Execute Commands in Active Sessions

  By default, project contributors, project administrators, and site administrators have the ability to execute commands within your actively running sessions in the Workbench. Cloudera Data Science Workbench 1.4.3 introduces a new feature that allows site administrators to restrict this ability. When this feature is enabled, only the user that creates the session will be able to execute commands in that session. No other users, regardless of their permissions in the team or as project collaborators/administrators, will be able to execute commands on active sessions that were not created by them.

  For details on how to enable this feature, see Restricting Collaborator and Administrator Access to Active Sessions on page 128.

Issues Fixed in Cloudera Data Science Workbench 1.4.3

**TSB-349: SQL Injection Vulnerability in Cloudera Data Science Workbench**

An SQL injection vulnerability was found in Cloudera Data Science Workbench. This would allow any authenticated user to run arbitrary queries against CDSW’s internal database. The database contains user contact information, bcrypt-hashed CDSW passwords (in the case of local authentication), API keys, and stored Kerberos keytabs.
**Products affected:** Cloudera Data Science Workbench (CDSW)

**Releases affected:** CDSW 1.4.0, 1.4.1, 1.4.2

**Users affected:** All

**Date/time of detection:** 2018-10-18

**Detected by:** Milan Magyar (Cloudera)

**Severity (Low/Medium/High):** Critical (9.9): **CVSS:3.0/AV:N/AC:L/PR:L/UI:N/S:C/C:H/I:H/A:H**

**Impact:** An authenticated CDSW user can arbitrarily access and modify the CDSW internal database. This allows privilege escalation in CDSW, Kubernetes, and the Linux host; creation, deletion, modification, and exfiltration of data, code, and credentials; denial of service; and data loss.

**CVE:** CVE-2018-20091

**Immediate action required:**

1. Strongly consider performing a backup before beginning. We advise you to have a backup before performing any upgrade and before beginning this remediation work.

2. Upgrade to Cloudera Data Science Workbench 1.4.3 (or higher).

3. In an abundance of caution Cloudera recommends that you revoke credentials and secrets stored by CDSW. To revoke these credentials:
   a. Change the password for any account with a keytab or kerberos credential that has been stored in CDSW. This includes the Kerberos principals for the associated CDH cluster if entered on the CDSW “Hadoop Authentication” user settings page.
   b. With Cloudera Data Science Workbench 1.4.3 running, run the following remediation script on each CDSW host, including the master and all workers: Remediation Script for TSB-349
      
      **Note:** Cloudera Data Science Workbench will become unavailable during this time.
   c. The script performs the following actions:
      a. If using local user authentication, logs out every user and resets their CDSW password.
      b. Regenerates or deletes various keys for every user.
      c. Resets secrets used for internal communications.
   d. Fully stop and start Cloudera Data Science Workbench (a restart is not sufficient).
      • For CSD-based deployments, restart the CDSW service in Cloudera Manager.
      
      **OR**
      • For RPM-based deployments, run cdsw stop followed by cdsw start on the CDSW master host.
   e. If using internal TLS termination: revoke and regenerate the CDSW TLS certificate and key.
   f. For each user, revoke the previous CDSW-generated SSH public key for git integration on the git side (the private key in CDSW has already been deleted). A new SSH key pair has already been generated and should be installed in the old key’s place.
   g. Revoke and regenerate any credential stored within a CDSW project, including any passwords stored in projects’ environment variables.

4. Verify all CDSW settings to ensure they are unchanged (e.g. SMTP server, authentication settings, custom docker images, host mounts, etc).

5. Treat all CDSW hosts as potentially compromised with root access. Remediate per your policy.
Addressed in release/refresh/patch: Cloudera Data Science Workbench 1.4.3
For the latest update on this issue see the corresponding Knowledge article: TSB 2019-349: CDSW SQL Injection Vulnerability

TSB-350: Risk of Data Loss During Cloudera Data Science Workbench (CDSW) Shutdown and Restart
Stopping Cloudera Data Science Workbench involves unmounting the NFS volumes that store CDSW project directories and then cleaning up a folder where CDSW stores its temporary state. However, due to a race condition, this NFS unmount process can take too long or fail altogether. If this happens, any CDSW projects that remain mounted will be deleted.

TSB-2018-346 was released in the time-frame of CDSW 1.4.2 to fix this issue, but it only turned out to be a partial fix. With CDSW 1.4.3, we have fixed the issue permanently. However, the script that was provided with TSB-2018-346 still ensures that data loss is prevented and must be used to shutdown/restart all the affected CDSW released listed below. The same script is also available under the Immediate Action Required section below.

Products affected: Cloudera Data Science Workbench
Releases affected: Cloudera Data Science Workbench versions
- 1.0.x
- 1.1.x
- 1.2.x
- 1.3.0, 1.3.1
- 1.4.0, 1.4.1, 1.4.2

Users affected: This potentially affects all CDSW users.
Detected by: Nehmé Tohmé (Cloudera)
Severity (Low/Medium/High): High
Impact: Potential data loss.
CVE: N/A

Immediate action required: If you are running any of the affected Cloudera Data Science Workbench versions, you must run the following script on the CDSW master host every time before you stop or restart Cloudera Data Science Workbench. Failure to do so can result in data loss.

This script should also be run before initiating a Cloudera Data Science Workbench upgrade. As always, we recommend creating a full backup prior to beginning an upgrade.

cds wcharkstop restart.sh - Available for download at: cdsn protect_stop restart.sh.

#!/bin/bash
set -e
cat << EXPLANATION
This script is a workaround for Cloudera TSB-346 and TSB-350. It protects your CDSW projects from a rare race condition that can result in data loss.
Run this script before stopping the CDSW service, irrespective of whether the stop precedes a restart, upgrade, or any other task.
Run this script only on the master node of your CDSW cluster.
You will be asked to specify a target folder on the master node where the script will save a backup of all your project files. Make sure the target folder has enough free space to accommodate all of your project files. To
determine how much space is required, run 'du -hs /var/lib/cdsw/current/projects' on the CDSW master node.

This script will first back up your project files to the specified target folder. It will then temporarily move your project files aside to protect against the data loss condition. At that point, it is safe to stop the CDSW service. After CDSW has stopped, the script will move the project files back into place.

Note: This workaround is not required for CDSW 1.4.3 and higher.

EXPLANATION

read -p "Enter target folder for backups: " backup_target

echo "Backing up to $backup_target..."

rsync -azp /var/lib/cdsw/current/projects "$backup_target"

read -n 1 -p "Backup complete. Press enter when you are ready to stop CDSW: "

echo "Deleting all Kubernetes resources..."

kubectl delete configmaps,deployments,daemonsets,replicasets,services,ingress,secrets,persistentvolumes,persistentvolumeclaims,jobs --all

kubectl delete pods --all

echo "Temporarily saving project files to /var/lib/cdsw/current/projects_tmp..."

mkdir /var/lib/cdsw/current/projects_tmp

mv /var/lib/cdsw/current/projects/* /var/lib/cdsw/current/projects_tmp

echo -e "Please stop the CDSW service."

read -n 1 -p "Press enter when CDSW has stopped: "

echo "Moving projects back into place..."

mv /var/lib/cdsw/current/projects_tmp/* /var/lib/cdsw/current/projects

rm -rf /var/lib/cdsw/current/projects_tmp

echo -e "Done. You may now upgrade or start the CDSW service."

echo -e "When CDSW is running, if desired, you may delete the backup data at $backup_target"

Addressed in release/refresh/patch: This issue is fixed in Cloudera Data Science Workbench 1.4.3.

Note that you are required to run the workaround script above when you upgrade from an affected version to a release with the fix. This helps guard against data loss when the affected version needs to be shut down during the upgrade process.

TSB-351: Unauthorized Project Access in Cloudera Data Science Workbench

Malicious CDSW users can bypass project permission checks and gain read-write access to any project folder in CDSW.

Products affected: Cloudera Data Science Workbench

Releases affected: Cloudera Data Science Workbench 1.4.0, 1.4.1, 1.4.2

Users affected: All CDSW Users

Date/time of detection: 10/29/2018

Detected by: Che-Yuan Liang (Cloudera)


Impact: Project data can be read or written (changed, destroyed) by any Cloudera Data Science Workbench user.

CVE: CVE-2018-20090

Immediate action required:
Upgrade to a version of Cloudera Data Science Workbench with the fix (version 1.4.3, 1.5.0, or higher).

**Addressed in release/refresh/patch:** Cloudera Data Science Workbench 1.4.3 (and higher)

*For the latest update on this issue see the corresponding Knowledge article:*

TSB 2019-351: Unauthorized Project Access in Cloudera Data Science Workbench

**Other Notable Fixed Issues in Cloudera Data Science Workbench 1.4.3**

- Fixed an issue where malicious Cloudera Data Science Workbench users were able to bypass project permission checks and gain read-write access to any project folder in Cloudera Data Science Workbench.
  
  Cloudera Bug: DSE-5138

- Fixed an issue where Cloudera Data Science Workbench would become unresponsive because the web application was making too many simultaneous requests to the Kubernetes API server. CDSW now caches calls to the API and refreshes the cache periodically.
  
  Cloudera Bug: DSE-5265, DSE-5269

- Fixed an issue where Cloudera Data Science Workbench workloads would intermittently crash with Exit Code 2: Misuse of Shell builtins.
  
  Cloudera Bug: DSE-4709

- Fixed an issue where Cloudera Data Science Workbench would not start when internal TLS termination was enabled and the TLS private key/certificate pair in use did not include a trailing newline character.
  
  Cloudera Bug: DSE-4853

**Known Issues and Limitations in Cloudera Data Science Workbench 1.4.3**

For a complete list of the current known issues and limitations in Cloudera Data Science Workbench 1.4.x, see [Known Issues and Limitations in Cloudera Data Science Workbench 1.5.x](#) on page 42.

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**Cloudera Data Science Workbench 1.4.2**

![Note: Cloudera Data Science Workbench 1.4.2 is the next official maintenance release after Cloudera Data Science Workbench 1.4.0. Version 1.4.1 is no longer publicly available.]

This section lists the release notes for Cloudera Data Science Workbench 1.4.2.

**New Features and Changes in Cloudera Data Science Workbench 1.4.2**

- **Operating System:** Added support for RHEL / CentOS / Oracle Linux RHCK 7.5.

- **Engines**
  - Mounts - By default, host mounts (specified at Admin > Engines > Mounts) are loaded into engine containers with read-only permissions. With version 1.4.2, a new checkbox allows you to make these mounted directories available in engine containers with read-write permissions instead.
  
  - Engine upgrade (Base Image v6)

- **Models**
  - In Cloudera Data Science Workbench 1.4.0, model request sizes were limited to 100 KB. In version 1.4.2, this limit has now been increased to 5 MB. To take advantage of this higher threshold, you will need to upgrade to Cloudera Data Science Workbench 1.4.2 and rebuild your existing models.

- **Security**
Added three new properties to the Admin > Security page that allow you to customize HTTP headers accepted by Cloudera Data Science Workbench.

- Enable HTTP security headers
- Enable cross-origin resource sharing (CORS)
- Enable HTTP Strict Transport Security (HSTS)

For details, see Configuring HTTP Headers for Cloudera Data Science Workbench on page 244.

Engine Upgrade

Cloudera Data Science Workbench 1.4.2 ships version 6 of the base engine image which includes the following versions of R and Python:

- R - 3.4.1
- Python - 2.7.11, 3.6.1

Pre-installed Packages in Engine 6 - For details about the packages included in the base engine, see Cloudera Data Science Workbench Engine Versions and Packaging on page 187.

Additionally, Cloudera Data Science Workbench will now alert you when a new engine version is available. Make sure you test and upgrade existing projects to Base Image v6 (Project Settings > Engine) to take advantage of the latest fixes.

Issues Fixed in Cloudera Data Science Workbench 1.4.2

TSB-346: Risk of Data Loss During Cloudera Data Science Workbench (CDSW) Shutdown and Restart

Stopping Cloudera Data Science Workbench involves unmounting the NFS volumes that store CDSW project directories and then cleaning up a folder where the kubelet stores its temporary state. However, due to a race condition, this NFS unmount process can take too long or fail altogether. If this happens, CDSW projects that remain mounted will be deleted by the cleanup step.

Products affected: Cloudera Data Science Workbench

Releases affected: Cloudera Data Science Workbench versions -

- 1.0.x
- 1.1.x
- 1.2.x
- 1.3.0, 1.3.1
- 1.4.0, 1.4.1

Users affected: This potentially affects all CDSW users.

Detected by: Nehmé Tohmé (Cloudera)

Severity (Low/Medium/High): High

Impact: If the NFS unmount fails during shutdown, data loss can occur. All CDSW project files might be deleted.

CVE: N/A

Immediate action required: If you are running any of the affected Cloudera Data Science Workbench versions, you must run the following script on the CDSW master host every time before you stop or restart Cloudera Data Science Workbench. Failure to do so can result in data loss.

This script should also be run before initiating a Cloudera Data Science Workbench upgrade. As always, we recommend creating a full backup prior to beginning an upgrade.
This script is a workaround for Cloudera TSB-346. It protects your CDSW projects from a rare race condition that can result in data loss. Run this script before stopping the CDSW service, irrespective of whether the stop precedes a restart, upgrade, or any other task.

Run this script only on the master node of your CDSW cluster.

You will be asked to specify a target folder on the master node where the script will save a backup of all your project files. Make sure the target folder has enough free space to accommodate all of your project files. To determine how much space is required, run 'du -hs /var/lib/cdsw/current/projects' on the CDSW master node.

This script will first back up your project files to the specified target folder. It will then temporarily move your project files aside to protect against the data loss condition. At that point, it is safe to stop the CDSW service. After CDSW has stopped, the script will move the project files back into place.

Note: This workaround is not required for CDSW 1.4.2 and higher.

Addressed in release/refresh/patch: This issue is fixed in Cloudera Data Science Workbench 1.4.2.

Note that you are required to run the workaround script above when you upgrade from an affected version to a release with the fix. This helps guard against data loss when the affected version needs to be shut down during the upgrade process.

For the latest update on this issue see the corresponding Knowledge article:

TSB 2018-346: Risk of Data Loss During Cloudera Data Science Workbench (CDSW) Shutdown and Restart
TSB-328: Unauthenticated User Enumeration in Cloudera Data Science Workbench

Unauthenticated users can get a list of user accounts of Cloudera Data Science Workbench.

**Products affected:** Cloudera Data Science Workbench

**Releases affected:** Cloudera Data Science Workbench 1.4.0 (and lower)

**Users affected:** All users of Cloudera Data Science Workbench 1.4.0 (and lower)

**Date/time of detection:** June 11, 2018

**Severity (Low/Medium/High):** 5.3 (Medium) **CVSS:** 3.0/AV:N/AC:L/PR:N/UI:N/S:U/C:L/I:N/A:N

**Impact:** Unauthenticated user enumeration in Cloudera Data Science Workbench.

**CVE:** CVE-2018-15665

**Immediate action required:** Upgrade to the latest version of Cloudera Data Science Workbench (1.4.2 or higher).

Note that Cloudera Data Science Workbench 1.4.1 is no longer publicly available due to **TSB 2018-346: Risk of Data Loss During Cloudera Data Science Workbench (CDSW) Shutdown and Restart.**

**Addressed in release/refresh/patch:** Cloudera Data Science Workbench 1.4.2 (and higher)

For the latest update on this issue see the corresponding Knowledge article:

**TSB 2018-318: Unauthenticated User Enumeration in Cloudera Data Science Workbench**

**Other Notable Fixed Issues in Cloudera Data Science Workbench 1.4.2**

- Fixed an issue where attempting to fork a large project would result in unexpected 'out of memory' errors.
  Cloudera Bug: DSE-4464

- Fixed an issue in version 1.4.0 where Cloudera Data Science Workbench workloads would intermittently get stuck in the Scheduling state due to a Red Hat kernel slab leak.
  Cloudera Bug: DSE-4098

- Fixed an issue in version 1.4.0 where the Hadoop username on non-kerberized clusters defaulted to cdsw. This was a known issue and has been fixed in version 1.4.2. The Hadoop username will now once again default to your Cloudera Data Science Workbench username.
  Cloudera Bug: DSE-4240

- Fixed an issue in version 1.4.0 where creating a project using Git via SSH did not work.
  Cloudera Bug: DSE-4278

- Fixed an issue in version 1.4.0 where environmental variables set in the Admin panel were not being propagated to projects (experiments, sessions, jobs) as expected.
  Cloudera Bug: DSE-4422

- Fixed an issue in version 1.4.0 where Cloudera Data Science Workbench would not start when external TLS termination was enabled.
  Cloudera Bug: DSE-4640

- Fixed an issue in version 1.4.0 where HTTP/HTTPS proxy settings in Cloudera Manager were erroneously escaped when propagated to Cloudera Data Science Workbench engines.
  Cloudera Bug: DSE-4421

- Fixed an issue in version 1.4.0 where **SSH tunnels** did not work as expected.
  Cloudera Bug: DSE-4741
• Fixed an issue in version 1.4.0 where copying multiple files into a folder resulted in unexpected behavior such as overwritten files and incorrect UI messages.

Cloudera Bug: DSE-4831

• Fixed an issue in version 1.4.0 where workers (in engines) and collection of usage metrics failed on TLS-enabled clusters.

Cloudera Bug: DSE-4293, DSE-4572

• Fixed an issue in version 1.4.0 where the Files > New Folder dialog box did not work.

Cloudera Bug: DSE-4807

• Fixed an issue in version 1.4.0 where deleting an experiment did not work from certain dashboards. Consequently, deleting the parent project would also fail in such cases.

Cloudera Bug: DSE-4227

Known Issues and Limitations in Cloudera Data Science Workbench 1.4.2

For a complete list of the current known issues and limitations in Cloudera Data Science Workbench 1.4.x, see Known Issues and Limitations in Cloudera Data Science Workbench 1.5.x on page 42.

Cloudera Data Science Workbench 1.4.0

This section lists the release notes for Cloudera Data Science Workbench 1.4.0.

New Features in Cloudera Data Science Workbench 1.4.0

• Models and Experiments - Cloudera Data Science Workbench 1.4 extends the machine learning platform experience from research to production. Now you can use Cloudera Data Science Workbench to build, train, and deploy models in a unified workflow.
  • Experiments - Train and compare versioned, reproducible models
  • Models - Deploy and manage models as REST APIs to serve predictions

• External Authentication
  – LDAP/SAML users can now restrict access to Cloudera Data Science Workbench to specific LDAP/SAML groups. Additionally, you can now specify groups that should automatically be granted site administrator privileges when they log in to Cloudera Data Science Workbench. For details, see Configuring External Authentication with LDAP and SAML on page 237.
  – Cloudera Data Science Workbench now supports multiple identity provider signing certificates for SAML 2.0 authentication.
  – Cloudera Data Science Workbench now supports SAML 2.0 Errata 05 E43 for SAML 2.0 authentication.

• Projects and Workbench
  – Site administrators can now disable individual built-in template projects by using a checkbox in the Project Templates table at Admin > Settings. Only enabled project templates will be displayed in the dropdown menu when creating a new project.
  – The default .gitignore file that is created with each new project has been updated to:

```
R
node_modules
*.pyc
```
• **Networking**
  - Cloudera Data Science Workbench now supports DNS resolution of localhost to non-local IP address (not 127.0.0.1).
  - Cloudera Data Science Workbench now appends the following default values to the `NO_PROXY` parameter if any of the following properties are configured: `HTTP_PROXY`, `HTTPS_PROXY`, or `ALL_PROXY`.

```bash
*127.0.0.1,localhost,100.66.0.1,100.66.0.2,100.66.0.3,
100.66.0.4,100.66.0.5,100.66.0.6,100.66.0.7,100.66.0.8,100.66.0.9,
100.66.0.10,100.66.0.11,100.66.0.12,100.66.0.13,100.66.0.14,
100.66.0.15,100.66.0.16,100.66.0.17,100.66.0.18,100.66.0.19,
100.66.0.20,100.66.0.21,100.66.0.22,100.66.0.23,100.66.0.24,
100.66.0.25,100.66.0.26,100.66.0.27,100.66.0.28,100.66.0.29,
100.66.0.30,100.66.0.31,100.66.0.32,100.66.0.33,100.66.0.34,
100.66.0.35,100.66.0.36,100.66.0.37,100.66.0.38,100.66.0.39,
100.66.0.40,100.66.0.41,100.66.0.42,100.66.0.43,100.66.0.44,
100.66.0.45,100.66.0.46,100.66.0.47,100.66.0.48,100.66.0.49,
100.66.0.50,100.77.0.10,100.77.0.128,100.77.0.129,100.77.0.130,
100.77.0.131,100.77.0.132,100.77.0.133,100.77.0.134,100.77.0.135,
100.77.0.136,100.77.0.137,100.77.0.138,100.77.0.139
```

• **Installation Validation Checks** - Improved validation checks run during the installation process. Cloudera Data Science Workbench now:
  - Verifies that the wildcard DNS subdomain has been configured.
  - Verifies that `resolv.conf` is not pointing to 127.0.0.1.
  - Validates iptables chains to ensure there are no custom rules being set.
  - Throws a warning if you are using a self-signed TLS certificate, an expired certificate, or if the certificate is not valid for the wildcard domain used for Cloudera Data Science Workbench.

• **Command Line** - Added a verbose option to the `cdsw status` command.

  `cdsw status [-v|--verbose]`

• Kubernetes has been upgraded to version 1.8.12.

**Engine Upgrade**

Cloudera Data Science Workbench 1.4.0 (and later) ships version 5 of the base engine image which includes the following versions of R and Python:

- R - 3.4.1
- Python - 2.7.11, 3.6.1

**Pre-installed Packages in Engine 5** - For details about the packages included in the base engine, see [Cloudera Data Science Workbench Engine Versions and Packaging](#) on page 187.

Additionally, Cloudera Data Science Workbench will now alert you when a new engine version is available. Make sure you test and upgrade existing projects to **Base Image v5 (Project Settings > Engine)** to take advantage of the latest fixes.

**Incompatible Changes in Cloudera Data Science Workbench 1.4.0**

**Host Mounts are now Read-Only in Engines** - Previously, mounts (specified at **Admin > Engines > Mounts**) were loaded into engine containers with read-write permissions.
Starting with version 1.4.0, mount points are now loaded into engines with **read-only** permissions.

**Issues Fixed in Cloudera Data Science Workbench 1.4.0**

- Fixed an issue where Git would timeout when cloning a project took too long. The timeout has now been increased to 60 seconds when creating a new project from Git.
  Cloudera Bug: DSE-3363
- Fixed an issue where manual parcel deployments could not detect parcel hash files with a `.sha1` extension.
  Cloudera Bug: DSE-3301
- Fixed several usability issues (file create, save, and so on) with Internet Explorer 11.
  Cloudera Bug: DSE-3426, DSE-3434
- Fixed an issue where CSD installations would fail to recognize Oracle Linux 7.3 as a supported operating system.
  Cloudera Bug: DSE-3257
- Fixed an issue where Cloudera Data Science Workbench would hang with 100 percent CPU utilization.
  Cloudera Bug: DSE-3450
- Fixed a SAML 2.0 configuration issue where uploading the identity provider metadata XML file did not update identity provider signing certificate and/or SSO URL on Cloudera Data Science Workbench correctly.
  Cloudera Bug: DSE-3076
- Fixed an issue with SAML 2.0 authentication where the identity provider’s signature was not being validated correctly.
  Cloudera Bug: DSE-3694
- Fixed the **Save As** functionality in the project Workbench.
  Cloudera Bug: DSE-3870
- Fixed an issue where if a user had some files opened in the Workbench in a previous session, and those files no longer existed in the project filesystem, a File Not Found error would occur when opening the Workbench.
  Cloudera Bug: DSE-3835

**Known Issues and Limitations in Cloudera Data Science Workbench 1.4.0**

For a complete list of the current known issues and limitations in Cloudera Data Science Workbench 1.4.x, see [Known Issues and Limitations in Cloudera Data Science Workbench 1.5.x](#) on page 42.

**Cloudera Data Science Workbench 1.3.1**

This section lists the release notes for Cloudera Data Science Workbench 1.3.1.

**New Features in Cloudera Data Science Workbench 1.3.1**

- **Operating System**: Added support for RHEL / CentOS / Oracle Linux RHCK 7.5.
- **SAML**
  - Cloudera Data Science Workbench now supports multiple identity provider signing certificates for SAML 2.0 authentication.
  - Cloudera Data Science Workbench now supports SAML 2.0 Errata OS E43 for SAML 2.0 authentication.
Issues Fixed in Cloudera Data Science Workbench 1.3.1

Remote Command Execution and Information Disclosure in Cloudera Data Science Workbench

A configuration issue in Kubernetes used by Cloudera Data Science Workbench can allow remote command execution and privilege escalation in CDSW. A separate information permissions issue can cause the LDAP bind password to be exposed to authenticated CDSW users when LDAP bind search is enabled.

Products affected: Cloudera Data Science Workbench

Releases affected: Cloudera Data Science Workbench 1.3.0 (and lower)

Users affected: All users of Cloudera Data Science Workbench 1.3.0 (and lower)

Date/time of detection: May 16, 2018

Severity (Low/Medium/High): High

Impact: Remote command execution and information disclosure

CVE: CVE-2018-11215

Immediate action required: Upgrade to the latest version of Cloudera Data Science Workbench (1.3.1 or higher) and change the LDAP bind password if previously configured in Cloudera Data Science Workbench.

Addressed in release/refresh/patch: Cloudera Data Science Workbench 1.3.1 (and higher)

For the latest update on this issue see the corresponding Knowledge Base article:

TSB: 2018-313: Remote Command Execution and Information Disclosure

Other Notable Fixed Issues in Cloudera Data Science Workbench 1.3.1

• Fixed an issue where CSD installations would fail to recognize Oracle Linux 7.3 as a supported operating system.
  Cloudera Bug: DSE-3257

• Fixed several usability issues (file create, save, and so on) with Internet Explorer 11.
  Cloudera Bug: DSE-3426, DSE-3434

• Fixed a SAML 2.0 configuration issue where uploading the identity provider metadata XML file did not update identity provider signing certificate and/or SSO URL on Cloudera Data Science Workbench correctly.
  Cloudera Bug: DSE-3265

• Fixed an issue where the owner of a console output could not view their own shared consoles from sessions /job runs when sharing with Specific user/team.
  Cloudera Bug: DSE-3143

• Fixed issue with missing connectors in Jobs dependency chart.
  Cloudera Bug: DSE-3185

Known Issues and Limitations in Cloudera Data Science Workbench 1.3.1

For a list of the current known issues and limitations, refer the documentation for version 1.3.x at Cloudera Data Science Workbench 1.3.x.

Cloudera Data Science Workbench 1.3.0

This section lists the release notes for Cloudera Data Science Workbench 1.3.0.
New Features and Changes in Cloudera Data Science Workbench 1.3.0

- Added support for SUSE Linux Enterprise Server 12 SP3.
- Site administrators can now add template projects that are customized for their organization's use-cases.
- Version 1.3 introduces a new environment variable for Python 3 sessions called `PYSPARK3_PYTHON`. Python 2 sessions will continue to use the default `PYSPARK_PYTHON` variable. This will allow you to configure distinct variables for Python 2 and Python 3 applications.
- In the Cloudera Manager CDSW service, the **Wildcard DNS Domain** property has been renamed to **Cloudera Data Science Workbench Domain**.
- Output for the `cdsw version` command now includes the type of deployment you are running – RPM or CSD.
- Added `log4j` and `spark-defaults` sample configuration to the PySpark and Scala template projects.

Issues Fixed in Cloudera Data Science Workbench 1.3.0

- Fixed an issue where the `cdsw status` command failed to run all the required system checks.
  Cloudera Bug: DSE-3070
- Session lists now include additional metadata to help distinguish between different sessions.
  Cloudera Bug: DSE-2814
- Pre-install validation checks have been improved to detect issues with `iptables` modules and Java settings.
  Cloudera Bug: DSE-2293
- Fixed an issue with the `cdsw status` command output when TLS is enabled.
  Cloudera Bug: DSE-3182
- CDS 2.2 Release 2 fixes the issue where a PySpark application could only be run once per active Workbench session.
  Cloudera Bug: CDH-58475
- Fixed an issue that prevented Bokeh plots from rendering.
  Cloudera Bug: DSE-3134
- Fixed an issue in Cloudera Data Science Workbench 1.2.2 that prevented WebSocket re-connections and caused console hangs.
  Cloudera Bug: DSE-3085
- Improved CDSW service restart performance for CSD deployments.
  Cloudera Bug: DSE-2937

Incompatible Changes in Cloudera Data Science Workbench 1.3.0

Deploying Cloudera Data Science Workbench with Cloudera Director 2.7

While this is not a Cloudera Data Science Workbench change, you should note that Cloudera Director 2.7 includes a new instance-level setting that sets the `mountAllUnmountedDisks` property to false:

```yaml
normalizationConfig {
  mountAllUnmountedDisks: false
}
```
This means Cloudera Director 2.7 (and higher) users no longer need to set `lp.normalization.mountAllUnmountedDisksRequired` to `false` in the Cloudera Director server's `application.properties` file. Note that Cloudera Director 2.6 still requires this setting.

Known Issues and Limitations in Cloudera Data Science Workbench 1.3.0

For a list of the current known issues and limitations, refer the documentation for version 1.3.x at Cloudera Data Science Workbench 1.3.x.

Cloudera Data Science Workbench 1.2.2

This section lists the release notes for Cloudera Data Science Workbench 1.2.2. The documentation for version 1.2.x can be found at Cloudera Data Science Workbench 1.2.x.

New Features and Changes in Cloudera Data Science Workbench 1.2.2

- Added support for SUSE Linux Enterprise Server 12 SP2.
- Added support for multi-homed networks.
- Cloudera Director now allows you to deploy CSD-based Cloudera Data Science Workbench 1.2.x deployments on AWS. For more specifics on supported platforms, see Cloudera Altus Director Support (AWS and Azure Only) on page 56.
- Added a new environment variable called `MAX_TEXT_LENGTH` that allows you to set the maximum number of characters that can be displayed in a single text cell. By default, this value is set to 800,000 and any more characters will be truncated.

Engine Upgrade

Cloudera Data Science Workbench 1.2.2 (and later) ships version 4 of the base engine image which includes bug fixes related to Python development and Kerberos authentication. Engine 4 ships the following versions of R and Python:

- R - 3.4.1
- Python - 2.7.11, 3.6.1

For details about the packages included in the base engine, see Cloudera Data Science Workbench Engine Versions and Packaging on page 187.

Make sure you upgrade existing projects to Base Image v4 (Project Settings > Engine) to take advantage of these fixes.

The new engine also changes how you configure and use Conda in Python sessions and extended engines. For more details, see Using Conda with Cloudera Data Science Workbench on page 182.

Issues Fixed In Cloudera Data Science Workbench 1.2.2

- Fixed an issue where Conda environmental variables were not being propagated to the Terminal correctly.
  Cloudera Bug: DSE-2256
- Fixed an issue where GPUs were not being detected by Cloudera Data Science Workbench due to incorrect mount settings.
  Cloudera Bug: DSE-2957
- Fixed an issue where jobs were failing due to Kerberos TGT renewal issues.
  Cloudera Bug: DSE-1007
- Fixed an issue on Internet Explorer 10 and 11 where the browser would fail to render console output after launching too many interactive sessions.
  Cloudera Bug: DSE-2998, DSE-2979
- Cloudera Data Science Workbench now correctly renders HTML that contains iFrames with the `srcdoc` attribute.
Cloudera Data Science Workbench Release Notes

Known Issues and Limitations in Cloudera Data Science Workbench 1.2.2

For a list of known issues and limitations, refer the documentation for version 1.2.x at Cloudera Data Science Workbench 1.2.x.

Cloudera Data Science Workbench 1.2.1

This section lists the release notes for Cloudera Data Science Workbench 1.2.1. The documentation for version 1.2.x can be found at Cloudera Data Science Workbench 1.2.x.

Issues Fixed In Cloudera Data Science Workbench 1.2.1

- The Master Node IPv4 Address parameter has been added to Cloudera Manager’s Add Service wizard and is now a required parameter for installation on AWS. This should fix any related installation issues for deployments on AWS.

  Cloudera Bug: DSE-2879

- Fixed an issue with CSD-based deployments where certain operations would fail because the Prepare Node command was not installing all the required packages during First Run of the service. To see the updated list of packages that are now being installed by the Prepare Node command, refer the CSD install guide.

  Cloudera Bug: DSE-2869

- Fixed an issue where the LD_LIBRARY_PATH environmental variable was not getting propagated to CUDA engines.

  Cloudera Bug: DSE-2828

- Fixed an issue where stopping Cloudera Data Science Workbench on worker hosts resulted in the application hanging indefinitely.

  Cloudera Bug: DSE-2880

Incompatible Changes in Cloudera Data Science Workbench 1.2.1

Upgrading from Cloudera Data Science Workbench 1.2.0 to 1.2.1 on CSD-based deployments

After upgrading from Cloudera Data Science Workbench 1.2.0 to 1.2.1 on a CSD-based deployment, CLI commands might not work as expected due to missing binaries in the environment. Note that this issue does not affect fresh installs.
Known Issues and Limitations in Cloudera Data Science Workbench 1.2.1

For a list of known issues and limitations, refer the documentation for version 1.2.x at Cloudera Data Science Workbench 1.2.x.

Cloudera Data Science Workbench 1.2.0

This section lists the release notes for Cloudera Data Science Workbench 1.2.0. The documentation for version 1.2.x can be found at Cloudera Data Science Workbench 1.2.x.

New Features and Changes in Cloudera Data Science Workbench 1.2.0

- Cloudera Data Science Workbench is now available as an add-on service for Cloudera Manager. To this end, Cloudera Data Science Workbench is now distributed in a parcel that integrates with Cloudera Manager using a Custom Service Descriptor (CSD). You can use Cloudera Manager to install, upgrade, and monitor Cloudera Data Science Workbench. Diagnostic data bundles can be generated and submitted to Cloudera through Cloudera Manager.
- Cloudera Data Science Workbench now enables secure sharing of job and session consoles. Additionally, site administrators can disable anonymous sharing from the Site Administrator dashboard (Admin > Security). See Sharing Job and Session Console Outputs on page 129.
- The Admin > Usage page now includes graphs for monitoring usage activity such as number of CPUs or GPUs used, memory usage, and total session runs, over customizable periods of time.
- Cloudera Data Science Workbench now lets you configure session, job, and idle timeouts. These can be configured using environmental variables either for the entire deployment or per-project.
- The csys enable and disable commands are no longer needed. The master host will now automatically detect the IP addresses of worker hosts joining or leaving Cloudera Data Science Workbench. See the revised Cloudera Data Science Workbench Command Line Reference on page 254.
- The Kudu Python client is now included in the Cloudera Data Science Workbench base engine image.
- Interactive session names can now be modified by project contributors and admins. By default, session names are set to 'Untitled Session'.
- All-numeric usernames are now accepted.
- Kubernetes has been upgraded to version 1.6.11.

Engine Upgrade

- Cloudera Data Science Workbench 1.2.0 ships version 3 of the base engine image which includes matplotlib improvements and the Kudu client libraries. Engine 3 ships the following versions of R and Python:
  - R - 3.4.1
  - Python - 2.7.11, 3.6.1

Make sure you upgrade existing projects to Base Image v3 (Project Settings > Engine) to take advantage of the new features and bug fixes included in the new engine.

Issues Fixed in Cloudera Data Science Workbench 1.2.0

Privilege Escalation and Database Exposure in Cloudera Data Science Workbench

Several web application vulnerabilities allowed malicious authenticated Cloudera Data Science Workbench (CDSW) users to escalate privileges in CDSW. In combination, such users could exploit these vulnerabilities to gain root access to CDSW hosts, gain access to the CDSW database which includes Kerberos keytabs of CDSW users and bcrypt hashed passwords, and obtain other privileged information such as session tokens, invitations tokens, and environmental variables.

Products affected: Cloudera Data Science Workbench

Releases affected: Cloudera Data Science Workbench 1.0.0, 1.0.1, 1.1.0, 1.1.1
Users affected: All users of Cloudera Data Science Workbench 1.0.0, 1.0.1, 1.1.0, 1.1.1

Date/time of detection: September 1, 2017

Detected by: NCC Group

Severity (Low/Medium/High): High

Impact: Privilege escalation and database exposure.

CVE: CVE-2017-15536

Addressed in release/refresh/patch: Cloudera Data Science Workbench 1.2.0 or higher.

Immediate action required: Upgrade to the latest version of Cloudera Data Science Workbench.

Other Notable Fixed Issues in Cloudera Data Science Workbench 1.2.0

- Fixed an issue where the Workbench editor screen jumps unexpectedly when typing or scrolling.
- Fixed auto-scroll behavior in the Workbench console. This was a browser compatibility issue that affected Chrome and Firefox, but not Safari.
- Fixed an issue where if a user logged out of Cloudera Data Science Workbench, and logged back in as a different user, they may see a `SecurityError` message in the Workbench.
- Fixed an issue that was preventing site administrators from uploading the SAML metadata file.
- Fixed several issues related to plotting with `matplotlib`. If you have previously used any workarounds for plotting, you might consider removing them now.
- Engines now use the same build of Kerberos utilities (`ktutil`, `kinit`, and `klist`) as the rest of Cloudera Data Science Workbench. This will improve logs obtained from `kinit` and make debugging Kerberos issues easier.
- `KRB5_TRACE` is now included in the error logs obtained when you `kinit`.
- Fixed an issue that was affecting health checks in deployments using AWS elastic load balancing.

Incompatible Changes in Cloudera Data Science Workbench 1.2.0

Proxy Configuration Change: If you are using a proxy server, you must ensure that the IP addresses for the web and Livelog services are skipped from the proxy.

Depending on your deployment (parcel or package), append the following IP addresses to either the `No Proxy` property in the Cloudera Manager CDSW service, or to the `NO_PROXY` parameter in `cdsw.conf`.

```
100.77.0.129
100.77.0.130
```

These have also been added to the installation instructions.

Known Issues and Limitations in Cloudera Data Science Workbench 1.2.0

For a list of known issues and limitations, refer the documentation for version 1.2.x at Cloudera Data Science Workbench 1.2.x.

Cloudera Data Science Workbench 1.1.1

This section lists the release notes for Cloudera Data Science Workbench 1.1.1. The documentation for version 1.1.x can be found at Cloudera Data Science Workbench 1.1.x.

New Features in Cloudera Data Science Workbench 1.1.1

- **Keytab Authentication** - With version 1.1.1, you can now authenticate yourself to the CDH cluster by uploading your Kerberos keytab to Cloudera Data Science Workbench. To use this feature, go to the top-right dropdown menu, click `Account settings > Hadoop Authentication`, enter your Kerberos principal and click `Upload Keytab`.
Issues Fixed in Cloudera Data Science Workbench 1.1.0

- Fixed support for dynamic data visualizations in Python, including Bokeh.
- Fixed issues with the Python template project. The project now supports offline mode and will therefore work on airgapped clusters.

Engine Upgrade

Cloudera Data Science Workbench 1.1.0 ships version 2 of the base engine image that includes new versions of Pandas, seaborn, and assorted bug fixes. Engine 2 ships the following versions of R and Python:

- R - 3.3.0
- Python - 2.7.11, 3.6.1

Make sure you upgrade existing projects to Base Image v2 (Project Settings > Engine) to take advantage of the new features and bug fixes included in the new engine.

Issues Fixed in Cloudera Data Science Workbench 1.1.0

- Improved support for dynamic data visualizations in Python, including Bokeh.
- Fixed issues with the Python template project. The project now supports offline mode and will therefore work on airgapped clusters.
• Fixed issues related to cached responses in Internet Explorer 11.
• Fixed issues with Java symlinks outside of JAVA_HOME.
• The cdsw status command can now be run on worker hosts.
• Removed unauthenticated localhost access to Kubernetes.
• Fixed Kerberos authentication issues with specific enc-types and Active Directory.
• Removed restrictions on usernames with special characters for better compatibility with external authentication systems such as Active Directory.
• Fixed issues with LDAP configuration validation that caused application crashes.
• Improved LDAP test configuration form to avoid confusion on parameters being sent.

Incompatible Changes in Cloudera Data Science Workbench 1.1.0

• Upgrading from version 1.0.x to 1.1.x
During the upgrade process, you will encounter incompatibilities between the two versions of cdsw.conf. This is because even though you are installing the latest RPM, your previous configuration settings in cdsw.conf will remain unchanged. Depending on the release you are upgrading from, you will need to modify cdsw.conf to ensure it passes the validation checks run by the 1.1.x release.

Key changes to note:
• JAVA_HOME is now a required parameter. Make sure you add JAVA_HOME to cdsw.conf before you start Cloudera Data Science Workbench.
• Previous versions allowed MASTER_IP to be set to a DNS hostname. If you are still using a DNS hostname, switch to an IP address.
• Python engine updated in version 1.1.x
Version 1.1.x includes an updated base engine image for Python which no longer uses the deprecated pylab mode in Jupyter to import the numpy and matplotlib functions into the global scope. With version 1.1.x, engines will now use built-in functions like any rather than the pylab counterpart, numpy.any. As a result of this change, you might see certain behavioral changes and differences in results between the two versions.

Also note that Python projects originally created with engine 1 will be running pandas version 0.19, and will not auto-upgrade to version 0.20 by simply selecting engine 2. You will also need to manually install version 0.20.1 of pandas when you launch a project session.

Known Issues and Limitations in Cloudera Data Science Workbench 1.1.0
For a list of known issues and limitations, refer the documentation for version 1.1.x at Cloudera Data Science Workbench 1.1.x.

Cloudera Data Science Workbench 1.0.1
This section lists the release notes for Cloudera Data Science Workbench 1.0.1. The documentation for version 1.0.x can be found at Cloudera Data Science Workbench 1.0.x.

Issues Fixed in Cloudera Data Science Workbench 1.0.1
• Fixed a random port conflict that could prevent Scala engines from running.
• Improved formatting of validation, and visibility of some errors.
• Fixed an issue with Firefox that was resulting in duplicate jobs on job creation.
• Removed the Mathjax external dependency on CDN.
• Improved PATH and JAVA_HOME handling that previously broke Hadoop CLIs.
• Fixed an issue with Java security policy files that caused Kerberos issues.
• Fixed an issue that caused git clone to fail on some repositories.
• Fixed an issue where updating LDAP admin settings deactivated the local fallback login.
• Fixed an issue where bad LDAP configuration crashed the application.
• Fixed an issue where job environmental variable settings did not persist.

Known Issues and Limitations in Cloudera Data Science Workbench 1.0.x

For a list of known issues and limitations, refer the documentation for version 1.0.x at Cloudera Data Science Workbench 1.0.x.

Cloudera Data Science Workbench 1.0.0

Version 1.0 represents the first generally available (GA) release of Cloudera Data Science Workbench. For information about the main features and benefits of Cloudera Data Science Workbench, as well as an architectural overview of the product, see Cloudera Data Science Workbench Overview on page 14.

Known Issues and Limitations in Cloudera Data Science Workbench 1.5.x

This topic lists the current known issues and limitations in Cloudera Data Science Workbench 1.5.x. For previous versions, see:

- Known Issues in Cloudera Data Science Workbench 1.4.x
- Known Issues in Cloudera Data Science Workbench 1.3.x
- Known Issues in Cloudera Data Science Workbench 1.2.x
- Known Issues in Cloudera Data Science Workbench 1.1.x

Upgrades

Please read the following upgrade issues before you being the upgrade process:

TSB-350: Permanent Fix for Data Loss Risk During Cloudera Data Science Workbench (CDSW) Shutdown and Restart

TSB-346 was released in the time-frame of CDSW 1.4.2 to fix this issue, but it only turned out to be a partial fix. With CDSW 1.4.3, we have fixed the issue permanently and released TSB-350 to address this fix. Note that the script that was provided with TSB-346 still ensures that data loss is prevented and must be used to shutdown/restart all the affected CDSW released listed below.

Affected Versions: Cloudera Data Science Workbench 1.0.x, 1.1.x, 1.2.x, 1.3.x, 1.4.0, 1.4.1, 1.4.2

Fixed Version: Cloudera Data Science Workbench 1.4.3 (and higher)

Cloudera Bug: DSE-5108

The complete text for TSB-350 is available in the 1.4.3 release notes and in the Cloudera Security Bulletins: TSB-350: Risk of Data Loss During Cloudera Data Science Workbench (CDSW) Shutdown and Restart.

TSB-346: Risk of Data Loss During Cloudera Data Science Workbench (CDSW) Shutdown and Restart

Stopping Cloudera Data Science Workbench involves unmounting the NFS volumes that store CDSW project directories and then cleaning up a folder where the kubelet stores its temporary state. However, due to a race condition, this NFS
unmount process can take too long or fail altogether. If this happens, CDSW projects that remain mounted will be deleted by the cleanup step.

**Products affected:** Cloudera Data Science Workbench

**Releases affected:** Cloudera Data Science Workbench versions -
- 1.0.x
- 1.1.x
- 1.2.x
- 1.3.0, 1.3.1
- 1.4.0, 1.4.1

**Users affected:** This potentially affects all CDSW users.

**Detected by:** Nehmé Tohmé (Cloudera)

**Severity (Low/Medium/High):** High

**Impact:** If the NFS unmount fails during shutdown, data loss can occur. All CDSW project files might be deleted.

**CVE:** N/A

**Immediate action required:** If you are running any of the affected Cloudera Data Science Workbench versions, you must run the following script on the CDSW master host every time before you stop or restart Cloudera Data Science Workbench. Failure to do so can result in data loss.

This script should also be run before initiating a Cloudera Data Science Workbench upgrade. As always, we recommend creating a full backup prior to beginning an upgrade.

**cdsw_protect_stop_restart.sh** - Available for download at: [cdsw_protect_stop_restart.sh](#)

```
#!/bin/bash
set -e
cat << EXPLANATION
This script is a workaround for Cloudera TSB-346. It protects your CDSW projects from a rare race condition that can result in data loss. Run this script before stopping the CDSW service, irrespective of whether the stop precedes a restart, upgrade, or any other task.

Run this script only on the master node of your CDSW cluster.

You will be asked to specify a target folder on the master node where the script will save a backup of all your project files. Make sure the target folder has enough free space to accommodate all of your project files. To determine how much space is required, run 'du -hs /var/lib/cdsw/current/projects' on the CDSW master node.

This script will first back up your project files to the specified target folder. It will then temporarily move your project files aside to protect against the data loss condition. At that point, it is safe to stop the CDSW service. After CDSW has stopped, the script will move the project files back into place.

Note: This workaround is not required for CDSW 1.4.2 and higher.

EXPLANATION
read -p "Enter target folder for backups: " backup_target
echo "Backing up to $backup_target..."
```
rsync -azp /var/lib/cdsw/current/projects "$backup_target"
read -n 1 -p "Backup complete. Press enter when you are ready to stop CDSW: "
    echo "Deleting all Kubernetes resources..."
    kubectl delete
    configmaps,deployments,daemonsets,replicasets,services,ingress,secrets,persistentvolumes,persistentvolumesclaims,jobs
    --all
    kubectl delete pods --all
    echo "Temporarily saving project files to /var/lib/cdsw/current/projects_tmp..."
    mkdir /var/lib/cdsw/current/projects_tmp
    mv /var/lib/cdsw/current/projects/* /var/lib/cdsw/current/projects_tmp
    echo -e "Please stop the CDSW service."
    read -n 1 -p "Press enter when CDSW has stopped: "
    echo "Moving projects back into place..."
    mv /var/lib/cdsw/current/projects_tmp/* /var/lib/cdsw/current/projects
    rm -rf /var/lib/cdsw/current/projects_tmp
    echo -e "Done. You may now upgrade or start the CDSW service."
    echo -e "When CDSW is running, if desired, you may delete the backup data at
    $backup_target"

**Addressed in release/refresh/patch:** This issue is fixed in Cloudera Data Science Workbench 1.4.2.

Note that you are required to run the workaround script above when you upgrade from an affected version to a release with the fix. This helps guard against data loss when the affected version needs to be shut down during the upgrade process.

For the latest update on this issue see the corresponding Knowledge article:

**TSB 2018-346: Risk of Data Loss During Cloudera Data Science Workbench (CDSW) Shutdown and Restart**

(Red Hat Only) Host Reboot Required for Upgrades from Cloudera Data Science Workbench 1.4.0

Cloudera Data Science Workbench 1.4.2 includes a fix for a Red Hat kernel slab leak issue that was found in Cloudera Data Science Workbench 1.4.0. However, to have this fix go into effect, Red Hat users must reboot all Cloudera Data Science Workbench hosts before proceeding with an upgrade from CDSW 1.4.0 to CDSW 1.4.2 (or higher).

Therefore, if you are a Red Hat user upgrading from Cloudera Data Science Workbench 1.4.0, your upgrade path will require the following steps:

1. Run the `cdsw_protect_stop_restart.sh` script to safely stop CDSW.
2. Backup all your application data.
3. Reboot all Cloudera Data Science Workbench hosts. As a precaution, you should consult your cluster/IT administrator before you start rebooting hosts.
4. Proceed with the upgrade to Cloudera Data Science Workbench 1.4.2 (or higher).

These steps have also been added to the upgrade documentation here:

- **CSD Upgrades**
- **RPM Upgrades**

**Cloudera Bug:** DSE-4098

**CDH Integration**

**CDH client configuration changes require a full Cloudera Data Science Workbench reset**

Cloudera Data Science Workbench does not automatically detect configuration changes on the CDH cluster. Therefore, any changes made to CDH services, ranging from updates to service configuration properties to complete CDH or CDS parcel upgrades, must be followed by a full reset of Cloudera Data Science Workbench.
**Workaround:** Depending on your deployment, use one of the following sets of steps to perform a full reset of Cloudera Data Science Workbench. Note that this reset does not impact your data in any way.

- **CSD Deployments** - To reset Cloudera Data Science Workbench using Cloudera Manager:
  1. Log into the Cloudera Manager Admin Console.
  2. On the Cloudera Manager homepage, click to the right of the CDSW service and select **Restart**. Confirm your choice on the next screen and wait for the action to complete.

  **OR**

- **RPM Deployments** - Run the following steps on the Cloudera Data Science Workbench master host:

  ```
cdsw reset
cdsw init
  ```

**Cloudera Manager Integration**

CSD distribution/activation fails on mixed-OS clusters when there are third-party parcels running on OSs that are not supported by Cloudera Data Science Workbench

For example, adding a new CDSW gateway host on a RHEL 6 cluster running RHEL-6 compatible parcels will fail. This is because Cloudera Manager will not allow distribution of the RHEL 6 parcels on the new host which will likely be running a CDSW-compatible operating system such as RHEL 7.

**Workaround:** To ensure adding a new CDSW gateway host is successful, you must create a copy of the 'incompatible' third-party parcel files and give them the corresponding RHEL 7 names so that Cloudera Manager allows them to be distributed on the new gateway host. Use the following sample instructions to do so:

1. SSH to the Cloudera Manager Server host.
2. Navigate to the directory that contains all the parcels. By default, this is `/opt/cloudera/parcels`.

```
cd /opt/cloudera/parcels
```

3. Make a copy of the incompatible third-party parcel with the new name. For example, if you have a RHEL 6 parcel that cannot be distributed on a RHEL 7 CDSW host:

```
cp <PARCELNAME.cdh5.x.x.p0.123>-el6.parcel <PARCELNAME.cdh5.x.x.p0.123>-el7.parcel
```

4. Repeat the previous step for parcel's SHA file.

```
cp <PARCELNAME.cdh5.x.x.p0.123>-el6.parcel.sha <PARCELNAME.cdh5.x.x.p0.123>-el7.parcel.sha
```

5. Update the new files' owner and permissions to match those of existing parcels in the `/opt/cloudera/parcels` directory.

```
chown cloudera-scm:cloudera-scm <PARCELNAME.cdh5.x.x.p0.123>-el7.parcel
chown cloudera-scm:cloudera-scm <PARCELNAME.cdh5.x.x.p0.123>-el7.parcel.sha
chmod 640 <PARCELNAME.cdh5.x.x.p0.123>-el7.parcel
chmod 640 <PARCELNAME.cdh5.x.x.p0.123>-el7.parcel.sha
```

You should now be able to add new gateway hosts for Cloudera Data Science Workbench to your cluster.

**Cloudera Bug:** OPSAPS-42130, OPSAPS-31880
CDSW Service health status after a restart does not match the actual state of the application

After a restart, the Cloudera Data Science Workbench service in Cloudera Manager will display **Good** health even though the Cloudera Data Science Workbench web application might need a few more minutes to get ready to serve requests.

Cloudera Data Science Workbench diagnostics data might be missing from Cloudera Manager diagnostic bundles.

This occurs because the default timeout for Cloudera Manager data collection is currently set to 3 minutes. However, in the case of Cloudera Data Science Workbench, collecting metrics and logs using the `cdsw logs` command can take longer than 3 minutes.

**Workaround:** Use the following steps to modify the default timeout for Cloudera Data Science Workbench data collection:

1. Login to the Cloudera Manager Admin Console.
2. Go to the CDSW service.
3. Click **Configuration**.
4. Search for the **Docker Daemon Diagnostics Collection Timeout** property and set it to 5 minutes.
5. Click **Save Changes**.

Alternatively, you can generate a diagnostic bundle by running the `cdsw logs` command directly on the Master host.

Cloudera Bug: OPSAPS-44016, DSE-3160

CDS Powered By Apache Spark

Spark lineage collection is not supported with Cloudera Data Science Workbench

Lineage collection is enabled by default in Spark 2.3. This feature does not work with Cloudera Data Science Workbench because the lineage log directory is not automatically mounted into CDSW engines when a session/job is started.

**Affected Versions:** CDS 2.3 release 2 (and higher) Powered By Apache Spark

**With Spark 2.3 release 3**, if Spark cannot find the lineage log directory, it will automatically disable lineage collection for that application. Spark jobs will continue to execute in Cloudera Data Science Workbench, but lineage information will not be collected.

**With Spark 2.3 release 2**, Spark jobs will fail in Cloudera Data Science Workbench. Either upgrade to Spark 2.3 release 3 which includes a partial fix (as described above) or use one of the following workarounds to disable Spark lineage:

**Workaround 1: Disable Spark Lineage Per-Project in Cloudera Data Science Workbench**

To do this, set `spark.lineage.enabled` to `false` in a `spark-defaults.conf` file in your Cloudera Data Science Workbench project. This will need to be done individually for each project as required.

**Workaround 2: Disable Spark Lineage for the Cluster**

1. Log in to Cloudera Manager and go to the Spark 2 service.
2. Click **Configuration**.
3. Search for the **Enable Lineage Collection** property and uncheck the checkbox to disable lineage collection.
4. Click **Save Changes**.
5. Go back to the Cloudera Manager homepage and **restart the CDSW service** for this change to go into effect.

Cloudera Bug: DSE-3720, CDH-67643

Crashes and Hangs

- High I/O utilization on the application block device can cause the application to stall or become unresponsive. Users should read and write data directly from HDFS rather than staging it in their project directories.

- Installing ipywidgets or a Jupyter notebook into a project can cause Python engines to hang due to an unexpected configuration. The issue can be resolved by deleting the installed libraries from the R engine terminal.
Engines

- Configuring duplicate mount points in the site admin panel (Admin > Engines > Mounts) results in sessions crashing in the workbench.

  **Cloudera Bug:** DSE-3308

- Spawning remote workers fails in R when the env parameter is not set. For more details, see [Spawning Workers](#) on page 126.

  **Cloudera Bug:** DSE-3384

- Autofs mounts are not supported with Cloudera Data Science Workbench.

  **Cloudera Bug:** DSE-2238

- When using Conda to install Python packages, you must specify the Python version to match the Python versions shipped in the engine image (2.7.11 and 3.6.1). If not specified, the conda-installed Python version will not be used within a project. Pip (pip and pip3) does not face this issue.

Custom Engine Images

- Cloudera Data Science Workbench only supports customized engines that are based on the Cloudera Data Science Workbench base image.

- Cloudera Data Science Workbench does not support creation of custom engines larger than 10 GB.

  **Cloudera Bug:** DSE-4420

- Cloudera Data Science Workbench does not support pulling images from registries that require Docker credentials.

  **Cloudera Bug:** DSE-1521

- The contents of certain pre-existing standard directories such as /home/cdsw, /tmp, /opt/cloudera, and so on, cannot be modified while creating customized engines. This means any files saved in these directories will not be accessible from sessions that are running on customized engines.

  **Workaround:** Create a new custom directory in the Dockerfile used to create the customized engine, and save your files to that directory. Or, create a new custom directory on all the Cloudera Data Science Workbench gateway hosts and save your files to those directories. Then, mount this directory to the custom engine.

Experiments

- Experiments do not store snapshots of project files. You cannot automatically restore code that was run as part of an experiment.

- Experiments will fail if your project filesystem is too large for the Git snapshot process. As a general rule, any project files (code, generated model artifacts, dependencies, etc.) larger than 50 MB must be part of your project's .gitignore file so that they are not included in snapshots for experiment builds.

- Experiments cannot be deleted. As a result, be conscious of how you use the track_metrics and track_file functions.

  - Do not track files larger than 50MB.
  - Do not track more than 100 metrics per experiment. Excessive metric calls from an experiment may cause Cloudera Data Science Workbench to hang.

- The Experiments table will allow you to display only three metrics at a time. You can select which metrics are displayed from the metrics dropdown. If you are tracking a large number of metrics (100 or more), you might notice some performance lag in the UI.

- Arguments are not supported with Scala experiments.

- The track_metrics and track_file functions are not supported with Scala experiments.
The UI does not display a confirmation when you start an experiment or any alerts when experiments fail.

**GPU Support**

**Only CUDA-enabled NVIDIA GPU hardware is supported**

Cloudera Data Science Workbench only supports CUDA-enabled NVIDIA GPU cards.

**Heterogeneous GPU hardware is not supported**

You must use the same GPU hardware across a single Cloudera Data Science Workbench deployment.

**GPUs are not detected after a machine reboot**

This issue occurs because certain NVIDIA modules do not load automatically after a reboot.

**Workaround:** To work around this issue, use the following steps to manually load the required modules before Cloudera Data Science Workbench services start. The following commands load the `nvidia.ko` module, create the `/dev/nvidiactl` device, and create the list of devices at `/dev/nvidia0`. They will also create the `/dev/nvidia-uvm` and `/dev/nvidia-uvm-tools` devices, and assign execute privileges to `/etc/rc.modules`. Run these commands once on all the machines that have GPU hardware.

Manually load the required NVIDIA modules:

```bash
sudo cat >> /etc/rc.modules <<EOMSG
/usr/bin/nvidia-smi
/usr/bin/nvidia-modprobe -u -c=0
EOMSG
```

Set execute permission for `/etc/rc.modules`:

```bash
sudo chmod +x /etc/rc.modules
```

**Cloudera Bug:** DSE-2847

**Jobs API**

- Cloudera Data Science Workbench does not support changing your API key, or having multiple API keys.
- Currently, you cannot create a job, stop a job, or get the status of a job using the Jobs API.

**Models**

- **Known Issues with Model Builds and Deployed Models**
  - Re-deploying or re-building models results in model downtime (usually brief).
  - Re-starting Cloudera Data Science Workbench does not automatically restart active models. These models must be manually restarted so they can serve requests again.
  - Model deployment will fail if your project filesystem is too large for the Git snapshot process. As a general rule, any project files (code, generated model artifacts, dependencies, etc.) larger than 50 MB must be part of your project’s .gitignore file so that they are not included in snapshots for model builds.
  - Model builds will fail if your project filesystem includes a .git directory (likely hidden or nested). Typical build stage errors include:

```
Error: 2 UNKNOWN: Unable to schedule build: [Unable to create a checkpoint of current source: [Unable to push sources to git server: ...
```
To work around this, rename the .git directory (for example, NO.git) and re-build the model.

Cloudera Bug: DSE-4657

- JSON requests made to active models should not be more than 5 MB in size. This is because JSON is not suitable for very large requests and has high overhead for binary objects such as images or video. Call the model with a reference to the image or video, such as a URL, instead of the object itself.

- Any external connections, for example, a database connection or a Spark context, must be managed by the model’s code. Models that require such connections are responsible for their own setup, teardown, and refresh.

- Model logs and statistics are only preserved so long as the individual replica is active. Cloudera Data Science Workbench may restart a replica at any time it is deemed necessary (such as bad input to the model).

- **(Affects version 1.4.x, 1.5.x)** The model deployment example (predict.py) in the in-built Python template project does not work anymore due to a change in dependencies in the sklearn package. A working replacement for the predict.py file has been provided here: [Deploy the Model - Iris Dataset](#).

  Cloudera Bug: DSE-5314

**Limitations**

- Scala models are not supported.

- **Spawning worker threads** is not supported with models.

- Models deployed using Cloudera Data Science Workbench are not highly-available.

- Dynamic scaling and auto-scaling are not currently supported. To change the number of replicas in service, you will have to re-deploy the build.

**Networking**

- Custom /etc/hosts entries on Cloudera Data Science Workbench hosts do not propagate to sessions and jobs running in containers.

  **Cloudera Bug:** DSE-2598

- Initialisation of Cloudera Data Science Workbench (cdsw init) will fail if localhost does not resolve to 127.0.0.1.

- Cloudera Data Science Workbench does not support DNS servers running on 127.0.0.1:53. This IP address resolves to the container localhost within Cloudera Data Science Workbench containers. As a workaround, use either a non-loopback address or a remote DNS server.

- Kubernetes throws the following error when /etc/resolv.conf lists more than three domains:

  ```
  Resolv.conf file '/etc/resolv.conf' contains search line consisting of more than 3 domains!
  ```

  Due to a limitation in the libc resolver, only two DNS servers are supported in /etc/resolv.conf. Kubernetes uses one additional entry for the cluster DNS.

**Security**

**SSH access to Cloudera Data Science Workbench hosts must be disabled**

The container runtime and application data storage is not fully secure from untrusted users who have SSH access to the gateway hosts. Therefore, SSH access to the gateway hosts for untrusted users should be disabled for security and resource utilization reasons.
LDAP

- LDAP group search fails when Active Directory returns escape characters as part of the distinguished name (DN).

  **Cloudera Bug:** DSE-4898

TLS/SSL

- Self-signed certificates where the Certificate Authority is not part of the user's trust store are not supported for TLS termination. For more details, see [Enabling TLS/SSL - Limitations](#).

- Cloudera Data Science Workbench does not support the use of encrypted private keys for TLS.

  **Cloudera Bug:** DSE-1708

- A "certificate has expired" error displays when you log in to the Cloudera Data Science Workbench web UI. This issue can occur if Cloudera Data Science Workbench exceeds 365 days of continuous uptime because the internal certificate for Kubernetes expires after 1 year.

  **Workaround:** Restart the Cloudera Data Science Workbench deployment.
  - For CSD installations, restart the Cloudera Data Science Workbench service in Cloudera Manager.
  - For RPM installations, run the following commands on the Master host:

```bash
# restart Cloudera Data Science Workbench
cdsw reset
# generate a new certificate for Kubernetes
cdsw init
```

Kerberos

- Using Kerberos plugin modules in `krb5.conf` is not supported.

- Modifying the `default_ccache_name` parameter in `krb5.conf` does not work in Cloudera Data Science Workbench. Only the default path for this parameter, `/tmp/krb5cc_${uid}`, is supported.

- PowerBroker-equipped Active Directory is not supported.

  **Cloudera Bug:** DSE-1838

- Cloudera Data Science Workbench does not support the use of a FreeIPA KDC.

  **Cloudera Bug:** DSE-1482

- When you upload a Kerberos keytab to authenticate yourself to the CDH cluster, Cloudera Data Science Workbench might display a fleeting error message ('cancelled') in the bottom right corner of the screen, even if authentication was successful. This error message can be ignored.

  **Cloudera Bug:** DSE-2344

Usability

- iFrame visualizations do not render in the workbench. Cloudera Data Science Workbench versions 1.4.2 (and higher) added a new feature that allowed users to [enable HTTP security headers](#) for responses to Cloudera Data Science Workbench. This setting is enabled by default. However, the X-Frame-Options header added as part of this feature blocks rendering of iFrames injected by third-party data visualization libraries.

  **Workaround:** To work around this issue, a site administrator can go to the **Admin > Security** page and disable the **Enable HTTP security headers** property. Restart Cloudera Data Science Workbench for this change to take effect.

  **Affected Version:** Cloudera Data Science Workbench 1.4.2 (and higher)

  **Cloudera Bug:** DSE-5274
• In a scenario where 100s of users are logged in and creating processes, the `nproc` and `nofile` limits of the system may be reached. Use `ulimits` or other methods to increase the maximum number of processes and open files that can be created by a user on the system.

• When rebooting, Cloudera Data Science Workbench hosts can take a significant amount of time (about 30 minutes) to become ready.

• Long-running operations such as `fork` and `clone` can time out when projects are large or connections outlast the HTTP timeouts of reverse proxies.

• The Scala kernel does not support autocomplete features in the editor.

• Scala and R code can sometimes indent incorrectly in the workbench editor.

  **Cloudera Bug:** DSE-1218
Cloudera Data Science Workbench 1.5.x Requirements and Supported Platforms

Important: For requirements and supported platforms specific to Hortonworks Data Platform, see Deploying Cloudera Data Science Workbench 1.5.x on Hortonworks Data Platform on page 85.

This topic lists the software and hardware configuration required to successfully install and run Cloudera Data Science Workbench. Cloudera Data Science Workbench does not support hosts or clusters that do not conform to the requirements listed on this page.

Cloudera Manager and CDH Requirements

Cloudera Data Science Workbench 1.5.x is supported on the following versions of CDH and Cloudera Manager:

<table>
<thead>
<tr>
<th>Type</th>
<th>CDH</th>
<th>Cloudera Manager</th>
</tr>
</thead>
<tbody>
<tr>
<td>CSD Deployments</td>
<td>• CDH 5.7 or higher</td>
<td>• Cloudera Manager 5.13 or higher</td>
</tr>
<tr>
<td></td>
<td>• CDH 6.1.x or higher</td>
<td>• Cloudera Manager 6.1.x or higher</td>
</tr>
<tr>
<td>RPM Deployments</td>
<td>• CDH 5.7 or higher</td>
<td>• Cloudera Manager 5.11 or higher</td>
</tr>
<tr>
<td></td>
<td>• CDH 6.1.x or higher</td>
<td>• Cloudera Manager 6.1.x or higher</td>
</tr>
</tbody>
</table>

All cluster hosts must be managed by Cloudera Manager. Note that all Cloudera Data Science Workbench administrative tasks require root access to the cluster’s gateway hosts where Cloudera Data Science Workbench is installed. Therefore, Cloudera Data Science Workbench does not support single-user mode installations.

Apache Spark Requirements

<table>
<thead>
<tr>
<th>CDH Version</th>
<th>Spark 2 Compatibility</th>
</tr>
</thead>
<tbody>
<tr>
<td>CDH 5</td>
<td>CDS 2.1.x Powered by Apache Spark (and higher)</td>
</tr>
<tr>
<td>CDH 6</td>
<td>On CDH 6 clusters, Apache Spark 2 is packaged with CDH and can no longer be installed separately. To find out which version of Spark 2 ships with your version of CDH 6, refer the CDH 6 Packaging Information guide.</td>
</tr>
</tbody>
</table>

Operating System Requirements

Cloudera Data Science Workbench 1.5.x is supported on the following operating systems. A gateway host that is dedicated to running Cloudera Data Science Workbench must use one of the following supported versions even if the remaining CDH hosts in your cluster are running any of the other operating systems supported by Cloudera Enterprise 5 or 6.

<table>
<thead>
<tr>
<th>Operating System</th>
<th>Versions</th>
<th>Notes</th>
</tr>
</thead>
<tbody>
<tr>
<td>RHEL / CentOS / Oracle Linux RHCK</td>
<td>7.2, 7.3, 7.4, 7.5</td>
<td>• When IPv6 is disabled, CDSW installations on RHEL/CentOS 7.3 fail due to an issue in kernel versions 3.10.0-514 - 3.10.0-693. For details, see <a href="https://access.redhat.com/solutions/3039771">https://access.redhat.com/solutions/3039771</a>.</td>
</tr>
</tbody>
</table>
Notes

• CDSW installations on RHEL/CentOS 7.2 might fail due to an issue with certain versions of the nfs-utils package. To fix the issue, either downgrade the nfs-utils package or upgrade to a version with the fix. View the complete Red Hat bug report here.

<table>
<thead>
<tr>
<th>Operating System</th>
<th>Versions</th>
<th>Notes</th>
</tr>
</thead>
<tbody>
<tr>
<td>Oracle Linux (UEK - default)</td>
<td>7.3</td>
<td>-</td>
</tr>
<tr>
<td>SUSE Linux Enterprise Server (SLES)</td>
<td>12 SP2, 12 SP3</td>
<td>Only supported with CSD deployments.</td>
</tr>
</tbody>
</table>

Cloudera Data Science Workbench publishes placeholder parcels for other operating systems as well. However, note that these do not work and have only been included to support mixed-OS clusters.

Additional OS-level Settings

- Enable memory cgroups on your operating system.
- Disable swap for optimum stability and performance. For instructions, see Setting the vm.swappiness Linux Kernel Parameter.
- Cloudera Data Science Workbench uses uid 8536 for an internal service account. Make sure that this user ID is not assigned to any other service or user account.
- Cloudera recommends that all users have the max-user-processes ulimit set to at least 65536.
- Cloudera recommends that all users have the max-open-files ulimit set to 1048576.

JDK Requirements

The entire CDH cluster, including Cloudera Data Science Workbench gateway hosts, must use the same version of JDK. Points to remember:

- Oracle JDK 7 is supported across all versions of Cloudera Manager 5 and CDH 5. Oracle JDK 8 is supported in Cloudera Enterprise 5.3.x and higher. Note the JDK 8 Requirement for Spark 2.2 (or higher) on page 53.
- OpenJDK 8 is supported in Cloudera Enterprise 5.16.1 and higher. OpenJDK 7 is not supported.
- For Red Hat/CentOS deployments in particular, Java Cryptography Extension (JCE) Unlimited Strength Jurisdiction must be enabled on the Cloudera Data Science Workbench gateway hosts.

For more specifics on the versions of Oracle JDK and OpenJDK recommended for CDH and Cloudera Manager clusters, and instructions on how to install the Java Cryptography Extension, see the Cloudera Product Compatibility Matrix - Supported JDK Versions.

JDK 8 Requirement for Spark 2.2 (or higher)

CSD-based deployments:

On CSD-based deployments, Cloudera Manager automatically detects the path and version of Java installed on Cloudera Data Science Workbench gateway hosts. You do not need to explicitly set the value for JAVA_HOME unless you want to use a custom location, use JRE, or (in the case of Spark 2) force Cloudera Manager to use JDK 1.8 as explained below.

To upgrade your entire CDH cluster to JDK 1.8, see Upgrading to Oracle JDK 1.8.

Package-based deployments:

Set JAVA_HOME to the JDK 8 path in cdsw.conf during the installation process. If you need to modify JAVA_HOME after the fact, restart the master and worker hosts to have the changes go into effect.
Networking and Security Requirements

**Important:** Make sure that your networking/security settings on Cloudera Data Science Workbench gateway hosts are not being overwritten behind-the-scenes, either by any automated scripts, or by other high-priority configuration such as `/etc/sysctl.conf`, `/etc/krb5.conf`, or `/etc/hosts.deny`.

- All Cloudera Data Science Workbench gateway hosts must be part of the same datacenter and use the same network. Hosts from different data-centers or networks can result in unreliable performance.
- A wildcard subdomain such as `*.cdsw.company.com` must be configured. Wildcard subdomains are used to provide isolation for user-generated content.

Starting with version 1.5, the wildcard DNS hostname configured for Cloudera Data Science Workbench must now be resolvable from both, the CDSW cluster, and your browser.

- Disable all pre-existing `iptables` rules. While Kubernetes makes extensive use of `iptables`, it’s difficult to predict how pre-existing `iptables` rules will interact with the rules inserted by Kubernetes. Therefore, Cloudera recommends you use the following commands to disable all pre-existing rules before you proceed with the installation.

```bash
sudo iptables -P INPUT ACCEPT
sudo iptables -P FORWARD ACCEPT
sudo iptables -P OUTPUT ACCEPT
sudo iptables -t nat -F
sudo iptables -t mangle -F
sudo iptables -F
sudo iptables -X
```

- Cloudera Data Science Workbench sets the following `sysctl` options in `/etc/sysctl.d/k8s.conf`:
  - `net.bridge.bridge-nf-call-iptables=1`
  - `net.bridge.bridge-nf-call-ip6tables=1`
  - `net.ipv4.ip_forward=1`
  - `net.ipv4.conf.default.forwarding=1`

  Underlying components of Cloudera Data Science Workbench (Docker, Kubernetes, and NFS) require these options to work correctly. Make sure they are not overridden by high-priority configuration such as `/etc/sysctl.conf`.

- SELinux must either be disabled or run in _permissive_ mode.
- Multi-homed networks are supported with Cloudera Data Science Workbench 1.2.2 (and higher).
- Firewall restrictions must be disabled across Cloudera Data Science Workbench and CDH hosts. Internally, the Cloudera Data Science Workbench master and worker hosts require full connectivity with no firewalls. Externally, end users connect to Cloudera Data Science Workbench exclusively through a web server running on the master host, and therefore do not need direct access to any other internal Cloudera Data Science Workbench or CDH services.

Review the complete list of ports required by Cloudera Data Science Workbench at [Ports Used By Cloudera Data Science Workbench](#) on page 216.

- Non-root SSH access is not allowed on Cloudera Data Science Workbench hosts.
- `localhost` must resolve to `127.0.0.1`.
- Cloudera Data Science Workbench does not support DNS servers running on `127.0.0.1:53`. This IP address resolves to the container localhost within Cloudera Data Science Workbench containers. As a workaround, use either a non-loopback address or a remote DNS server.

Cloudera Data Science Workbench does not support hosts or clusters that do not conform to these restrictions.
Recommended Hardware Configuration

**Important:**
- Allocate separate CDH gateway hosts for Cloudera Data Science Workbench. Do not reuse existing hosts that are already running other CDH services. Doing this can lead to port conflicts, unreliable execution of user workloads, and out-of-memory errors.
- All Cloudera Data Science Workbench gateway hosts must be part of the same datacenter and use the same network. Hosts from different data-centers or networks can result in unreliable performance.
- Starting with version 1.4.3, multi-host CDSW deployments can be customized to reserve the Master only for internal processes while user workloads are run exclusively on workers. For details, see Reserving the Master Host for Internal CDSW Components on page 209.

Cloudera Data Science Workbench hosts are added to your CDH cluster as gateway hosts. The recommended minimum hardware configuration for Cloudera Data Science Workbench gateway hosts is:

<table>
<thead>
<tr>
<th>Resource Type</th>
<th>Master</th>
<th>Workers</th>
<th>Notes</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>CPU</strong></td>
<td>16+ CPU (vCPU) cores</td>
<td>16+ CPU (vCPU) cores</td>
<td></td>
</tr>
<tr>
<td><strong>RAM</strong></td>
<td>32+ GB</td>
<td>32+ GB</td>
<td>If you are going to partition the root volume, make sure you allocate at least 20 GB to / so that the installer can proceed without running out of space.</td>
</tr>
<tr>
<td><strong>Disk Space</strong></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Root Volume</td>
<td>100+ GB</td>
<td>100+ GB</td>
<td></td>
</tr>
<tr>
<td>Application Block Device</td>
<td>1 TB</td>
<td>-</td>
<td>The Application Block Device is only required on the Master where it is mounted to /var/lib/cdsw. You will be asked to create a /var/lib/cdsw directory on all the Worker hosts during the installation process. However, they do not need to be mounted to a block device. It is only used to store client configuration for HDP cluster services on Workers.</td>
</tr>
<tr>
<td>Docker Block Device</td>
<td>1 TB</td>
<td>1 TB</td>
<td>The Docker Block Device is required on all Master and Worker hosts.</td>
</tr>
</tbody>
</table>

**Scaling Guidelines**

New hosts can be added and removed from a Cloudera Data Science Workbench deployment without interrupting any jobs already scheduled on existing hosts. Therefore, it is rather straightforward to increase capacity based on observed usage. At a minimum, Cloudera recommends you allocate at least 1 CPU core and 2 GB of RAM per concurrent session or job. CPU can burst above a 1 CPU core share when spare resources are available. Therefore, a 1 CPU core allocation is often adequate for light workloads. Allocating less than 2 GB of RAM can lead to out-of-memory errors for many applications.

As a general guideline, Cloudera recommends hosts with RAM between 60GB and 256GB, and between 16 and 48 cores. This provides a useful range of options for end users. Note that SSDs are strongly recommended for application data storage. Using standard HDDs can sometimes result in poor application performance.
For some data science and machine learning applications, users can collect a significant amount of data in memory within a single R or Python process, or use a significant amount of CPU resources that cannot be easily distributed into the CDH cluster. If individual users frequently run larger workloads or run workloads in parallel over long durations, increase the total resources accordingly. Understanding your users’ concurrent workload requirements or observing actual usage is the best approach to scaling Cloudera Data Science Workbench.

**Python Supported Versions**

The default Cloudera Data Science Workbench engine currently includes **Python 2.7.11** and **Python 3.6.1**. To use PySpark with lambda functions that run within the CDH cluster, the Spark executors must have access to a matching version of Python. For many common operating systems, the default system Python will not match the minor release of Python included in Data Science Workbench.

To ensure that the Python versions match, Python can either be installed on every CDH host or made available per job run using Spark’s ability to distribute dependencies. Given the size of a typical isolated Python environment and the desire to avoid repeated uploads from gateway hosts, Cloudera recommends installing Python 2.7 and 3.6 on the cluster if you are using PySpark with lambda functions.

You can install Python 2.7 and 3.6 on the cluster using any method and set the corresponding `PYSPARK_PYTHON` environment variable in your project. Cloudera Data Science Workbench 1.3 (and higher) include a separate environment variable for Python 3 sessions called `PYSPARK3_PYTHON`. Python 2 sessions continue to use the default `PYSPARK_PYTHON` variable. This will allow you to run Python 2 and Python 3 sessions in parallel without either variable being overridden by the other.

For an example on distributing Python dependencies dynamically, see **Example: Distributing Dependencies on a PySpark Cluster** on page 202.

**Anaconda**

Continuum Analytics and Cloudera have partnered to create an Anaconda parcel for CDH to enable simple distribution, installation, and management of popular Python packages and their dependencies. Note that this parcel is not directly supported by Cloudera.

**Docker and Kubernetes Support**

Cloudera Data Science Workbench only supports the versions of **Docker** and **Kubernetes** that are shipped with each release. Upgrading Docker or Kubernetes, or running on third-party Kubernetes clusters is not supported.

**Supported Browsers**

- Chrome (latest stable version)
- Firefox (latest released version and latest ESR version)
- Safari 9+
- Internet Explorer (IE) 11+
  - IE’s Compatibility View mode is not supported.

**Cloudera Altus Director Support (AWS and Azure Only)**

Altus Director support for Cloudera Data Science Workbench is available for the following platforms:

- **Amazon Web Services (AWS)** - Cloudera Altus Director 2.6.0 (and higher)
  - **Microsoft Azure** - Cloudera Altus Director 2.7 (and higher)
Deploying Cloudera Data Science Workbench with Altus Director

Points to note when using Altus Director to install Cloudera Data Science Workbench:

- (Required for Director 2.6) Before you run the command to bootstrap a new cluster, set the `lp.normalization.mountAllUnmountedDisksRequired` property to `false` in the Altus Director server’s `application.properties` file, and then restart Altus Director.

  Higher versions of Altus Director do not require this step. Altus Director 2.7 (and higher) include an instance-level setting called `mountAllUnmountedDisks` that must be set to `false` as demonstrated in the following sample configuration files.

- Depending on your cloud platform, you can use one of the following sample configuration files to deploy a Cloudera Manager cluster with Cloudera Data Science Workbench.
  - AWS - `aws.cdsw.conf`
  - Azure - `azure.cdsw.conf`

  Note that these sample files are tailored to Altus Director 2.7 (and higher) and they install a very limited CDH cluster with just the following services: HDFS, YARN, and Spark 2. You can extend them as needed to match your use case.

Related Topics:
- Using Products outside CDH with Altus Director
- Altus Director CLI
- The Altus Director Configuration File
- Setting Altus Director Properties

Recommended Configuration on Amazon Web Services (AWS)

On AWS, Cloudera Data Science Workbench must be used with persistent/long-running Apache Hadoop clusters only.

CDH and Cloudera Manager Hosts

- For instructions on deploying CDH and Cloudera Manager on AWS, refer the Cloudera Reference Architecture for AWS deployments.

Cloudera Data Science Workbench Hosts

- Operations
  - Use Cloudera Director to orchestrate operations. Use Cloudera Manager to monitor the cluster.

- Networking
  - No security group or network restrictions between hosts.
  - HTTP connectivity to the corporate network for browser access. Do not use proxies or manual SSH tunnels.

- Recommended Instance Types
  - m4.4xlarge-m4.16xlarge

    In this case, bigger is better. That is, one m4.16large is better than four m4.4xlarge hosts. AWS pricing scales linearly, and larger instances have more EBS bandwidth.

- Storage
  - 100 GB root volume block device (gp2) on all hosts
  - 500 GB Docker block devices (gp2) on all hosts
Recommended Configuration on Microsoft Azure

**CDH and Cloudera Manager Hosts**
- For instructions on deploying CDH and Cloudera Manager on Azure, refer the Cloudera Reference Architecture for Azure deployments.

**Cloudera Data Science Workbench Hosts**
- **Operations**
  - Use Cloudera Director to orchestrate operations. Use Cloudera Manager to monitor the cluster.
- **Networking**
  - No security group or network restrictions between hosts.
  - HTTP connectivity to the corporate network for browser access. Do not use proxies or manual SSH tunnels.
- **Recommended Instance Types**
  - DS13-DS14 v2 instances on all hosts.
- **Storage**
  - P30 premium storage for the Application and Docker block devices.
  
  Cloudera Data Science Workbench *requires* premium disks for its block devices on Azure. Standard disks can lead to unacceptable performance even on small clusters.
Installing Cloudera Data Science Workbench 1.5.x on CDH

This topic walks you through the installation paths available for Cloudera Data Science Workbench 1.5.x. It also describes the steps needed to configure your cluster gateway hosts and block devices before you can begin installing the Cloudera Data Science Workbench parcel/package.

Installing Cloudera Data Science Workbench 1.5.x

You can use one of the following ways to install Cloudera Data Science Workbench 1.5.x:

- **Using a Custom Service Descriptor (CSD) and Parcel** - Starting with version 1.2.x, Cloudera Data Science Workbench is available as an add-on service for Cloudera Manager. Two files are required for this type of installation: a CSD JAR file that contains all the configuration needed to describe and manage the new Cloudera Data Science Workbench service, and the Cloudera Data Science Workbench parcel. To install this service, first download and copy the CSD file to the Cloudera Manager Server host. Then use Cloudera Manager to distribute the Cloudera Data Science Workbench parcel to the relevant gateway hosts.

  Note that this installation mode does not apply to CDSW-on-HDP deployments.

  or

- **Using a Package (RPM)** - You can install the Cloudera Data Science Workbench package directly on your cluster’s gateway or edge hosts. In this case, you will not be able to manage the Cloudera Data Science Workbench service from a cluster manager such as Cloudera Manager or Ambari.

To begin the installation process, continue reading Required Pre-Installation Steps on page 59.

Airgapped Installations

Sometimes organizations choose to restrict parts of their network from the Internet for security reasons. Isolating segments of a network can provide assurance that valuable data is not being compromised by individuals out of maliciousness or for personal gain. However, in such cases isolated hosts are unable to access Cloudera repositories for new installations or upgrades. Effective version 1.1.1, Cloudera Data Science Workbench supports installation on CDH clusters that are not connected to the Internet.

For CSD-based installs in an airgapped environment, put the Cloudera Data Science Workbench parcel into a new hosted or local parcel repository, and then configure the Cloudera Manager Server to target this newly-created repository.

Required Pre-Installation Steps

The rest of this topic describes the steps you should take to review your platforms and configure your gateway hosts before you begin to install Cloudera Data Science Workbench.

Note: Worker hosts are not required for a fully-functional Cloudera Data Science Workbench deployment. For proof-of-concept deployments, you can deploy a 1-host cluster with just a Master host. The Master host can run user workloads just as a worker host can.

Even on multi-host deployments, the Master host doubles up to perform both functions: those of the Master outlined here, and those of a worker.
Review Requirements and Supported Platforms

Review the complete list of Cloudera Data Science Workbench 1.5.x Requirements and Supported Platforms on page 52 before you proceed with the installation.

Set Up a Wildcard DNS Subdomain

Cloudera Data Science Workbench uses DNS to route HTTP requests to specific engines and services. Wildcard subdomains (such as *.cdsw.<your_domain>.com) are required in order to provide isolation for user-generated content. In particular, wildcard subdomains help:

- Securely expose interactive session services, such as visualizations, the terminal, and web UIs such as TensorBoard, Shiny, Plotly, and so on.
- Securely isolate user-generated content from the application.

To set up subdomains for Cloudera Data Science Workbench, configure your DNS server with an A record for a wildcard DNS name such as *.cdsw.<your_domain>.com for the master host, and a second A record for the root entry of cdsw.<your_domain>.com.

For example, if your master IP address is 172.46.47.48, you'd configure two A records as follows:

```
  cdsw.<your_domain>.com   IN A 172.46.47.48
  *.cdsw.<your_domain>.com  IN A 172.46.47.48
```

You can also use a wildcard CNAME record if it is supported by your DNS provider.

Starting with version 1.5, the wildcard DNS hostname configured for Cloudera Data Science Workbench must now be resolvable from both, the CDSW cluster, and your browser.

Disable Untrusted SSH Access

Cloudera Data Science Workbench assumes that users only access the gateway hosts through the web application. Untrusted users with SSH access to a Cloudera Data Science Workbench host can gain full access to the cluster, including access to other users' workloads. Therefore, untrusted (non-sudo) SSH access to Cloudera Data Science Workbench hosts must be disabled to ensure a secure deployment.

For more information on the security capabilities of Cloudera Data Science Workbench, see the Cloudera Data Science Workbench Security Guide on page 224.

Configure Block Devices

Docker Block Device

The Cloudera Data Science Workbench installer will format and mount Docker on each gateway host. Make sure there is no important data stored on these devices. Do not mount these block devices prior to installation.

Every Cloudera Data Science Workbench gateway host must have one or more block devices with at least 1 TB dedicated to storage of Docker images. The Docker block devices store the Cloudera Data Science Workbench Docker images including the Python, R, and Scala engines. Each engine image can occupy 15GB.

Application Block Device or Mount Point

The master host on Cloudera Data Science Workbench requires at least 1 TB for database and project storage. This recommended capacity is contingent on the expected number of users and projects on the cluster. While large data files should be stored on HDFS, it is not uncommon to find gigabytes of data or libraries in individual projects. Running out of storage will cause the application to fail. Cloudera recommends allocating at least 5 GB per project and at least 1 TB of storage in total. Make sure you continue to carefully monitor disk space usage and I/O using Cloudera Manager.

Cloudera Data Science Workbench stores all application data at /var/lib/cdsw. On a CSD-based deployment, this location is not configurable. Cloudera Data Science Workbench will assume the system administrator has formatted
and mounted one or more block devices to /var/lib/cdsw on the master host. Note that Application Block Device mounts are not required on worker hosts.

Regardless of the application data storage configuration you choose, /var/lib/cdsw must be stored on a separate block device. Given typical database and user access patterns, an SSD is strongly recommended.

By default, data in /var/lib/cdsw is not backed up or replicated to HDFS or other hosts. Reliable storage and backup strategy is critical for production installations. For more information, see Backup and Disaster Recovery for Cloudera Data Science Workbench on page 215.

Install Cloudera Data Science Workbench

To use the Cloudera Manager CSD and parcel to install Cloudera Data Science Workbench, follow the steps at Installation and Upgrade Using Cloudera Manager.

OR

To install the Cloudera Data Science Workbench package on the cluster gateway hosts, follow the steps at Installation and Upgrade Using Packages.

Installing Cloudera Data Science Workbench 1.5.x Using Cloudera Manager

**Note:** Instructions on how to deploy Cloudera Data Science Workbench on Hortonworks Data Platform clusters are available here: Deploying Cloudera Data Science Workbench 1.5.x on Hortonworks Data Platform on page 85.

Use the following steps to install Cloudera Data Science Workbench using Cloudera Manager.

**Prerequisites**

Before you begin installing Cloudera Data Science Workbench, make sure you have completed the steps to secure your hosts, set up DNS subdomains, and configure block devices.

**Configure Apache Spark 2**

1. **(CDH 5 Only)** Install and configure the CDS 2.x Powered by Apache Spark parcel and CSD. For instructions, see Installing CDS 2.x Powered by Apache Spark.

   ![Important](https://cloudera.com/marketplace/assets/images/icons/important.png) **Important:** Do not install CDS 2.x if you are using CDH 6. Spark 2 ships as part of the CDH 6 parcel; the add-on parcel is no longer required. To see which version of Spark 2 ships with CDH, refer the CDH 6 Packaging documentation.

2. **(Required for CDH 5 and CDH 6)** To be able to use Spark 2, each user must have their own /home directory in HDFS. If you sign in to Hue first, these directories will automatically be created for you. Alternatively, you can have cluster administrators create these directories.

   ```bash
   hdfs dfs -mkdir /user/<username>
   hdfs dfs -chown <username>:<username> /user/<username>
   ```

   If you are using CDS 2.3 release 2 (or higher), review the associated known issues here: CDS Powered By Apache Spark on page 46.

3. Test Spark 2 integration on the gateway hosts.
   1. SSH to a gateway host.
   2. If your cluster is kerberized, run kinit to authenticate to the CDH cluster’s Kerberos Key Distribution Center. The Kerberos ticket you create is not visible to Cloudera Data Science Workbench users.
   3. Submit a test job to Spark by executing the following command:
CDH 5

```bash
spark2-submit --class org.apache.spark.examples.SparkPi --master yarn 
--deploy-mode client /opt/cloudera/parcels/SPARK2/lib/spark2/examples/jars/spark-example*.jar 100
```

To view a sample command, click

```bash
spark2-submit --class org.apache.spark.examples.SparkPi --master yarn 
--deploy-mode client /opt/cloudera/parcels/SPARK2/lib/spark2/examples/jars/spark-examples_2.11-2.3.0.cloudera5-SNAPSHOT.jar 100
```

CDH 6

```bash
spark-submit --class org.apache.spark.examples.SparkPi --master yarn 
--deploy-mode client SPARK_HOME/lib/spark-examples*.jar 100
```

To view a sample command, click

```bash
spark-submit --class org.apache.spark.examples.SparkPi --master yarn 
--deploy-mode client /opt/cloudera/parcels/CDH/lib/spark/examples/jars/spark-examples_2.11-2.4.0-cdh6.1.0.jar 100
```

4. View the status of the job in the CLI output or in the Spark web UI to confirm that the host you want to use for the Cloudera Data Science Workbench master functions properly as a Spark gateway.

To view sample CLI output, click

```bash
19/02/15 09:37:39 INFO spark.SparkContext: Running Spark version 2.4.0-cdh6.1.0
19/02/15 09:37:39 INFO spark.SparkContext: Submitted application: Spark Pi
... 19/02/15 09:37:40 INFO util.Utils: Successfully started service 'sparkDriver' on port 37050.
... 19/02/15 09:38:06 INFO scheduler.DAGScheduler: Job 0 finished: reduce at SparkPi.scala:38, took 18.659033 s
```

Configure JAVA_HOME

On CSD-based deployments, Cloudera Manager automatically detects the path and version of Java installed on Cloudera Data Science Workbench gateway hosts. You do not need to explicitly set the value for JAVA_HOME unless you want to use a custom location, use JRE, or (in the case of Spark 2) force Cloudera Manager to use JDK 1.8 as explained below.

Setting a value for JAVA_HOME - The value for JAVA_HOME depends on whether you are using JDK or JRE. For example, if you're using JDK 1.8_162, set JAVA_HOME to /usr/java/jdk1.8.0_162. If you are only using JRE, set it to /usr/java/jdk1.8.0_162/jre.

Issues with Spark 2.2 and higher - Spark 2.2 (and higher) requires JDK 1.8. However, if a host has both JDK 1.7 and JDK 1.8 installed, Cloudera Manager might choose to use JDK 1.7 over JDK 1.8. If you are using Spark 2.2 (or higher), this will create a problem during the first run of the service because Spark will not work with JDK 1.7. To work around this, explicitly configure Cloudera Manager to use JDK 1.8 on the gateway hosts that are running Cloudera Data Science Workbench.

For instructions on how to set JAVA_HOME, see Configuring a Custom Java Home Location in Cloudera Manager.

To upgrade the whole CDH cluster to JDK 1.8, see Upgrading to JDK 1.8.

Download and Install the Cloudera Data Science Workbench CSD

1. Download the Cloudera Data Science Workbench CSD. Make sure you download the CSD that corresponds to the version of CDH you are using.
Link to CSDVersion
CDH 6 - CLOUDERA_DATA SCIENCE WORKBENCH CDH6 1.5.0.jar
CDH 5 - CLOUDERA_DATA SCIENCE WORKBENCH CDH5 1.5.0.jar

2. Log on to the Cloudera Manager Server host, and place the CSD file under /opt/cloudera/csd, which is the default location for CSD files. To configure a custom location for CSD files, refer to the Cloudera Manager documentation at Configuring the Location of Custom Service Descriptor Files.

3. Set the file ownership to cloudera-scm:cloudera-scm with permission 644.

   Set the file ownership:

   chown cloudera-scm:cloudera-scm CLOUDERA_DATA SCIENCE WORKBENCH-CDH<X>-1.5.<Y>.jar

   Set the file permissions:

   chmod 644 CLOUDERA_DATA SCIENCE WORKBENCH-CDH<X>-1.5.<Y>.jar

4. Restart the Cloudera Manager Server:

   service cloudera-scm-server restart

5. Log into the Cloudera Manager Admin Console and restart the Cloudera Management Service:

   a. Select Clusters > Cloudera Management Service.
   b. Select Actions > Restart.

Install the Cloudera Data Science Workbench Parcel

1. Log into the Cloudera Manager Admin Console.
2. Click Hosts > Parcels in the main navigation bar.
3. If you placed the Cloudera Data Science Workbench CSD in the default CSD directory, /opt/cloudera/csd, Cloudera Manager auto-detects and adds the parcel repository URL. Therefore, the parcel should already be present and ready for downloading.

   If for some reason the parcel is not available, add the remote parcel repository URL to Cloudera Manager. For detailed steps, click

   1. On the Parcels page, click Configuration.
   2. In the Remote Parcel Repository URLs list, click the addition symbol to open an additional row.
   3. Enter the path to the repository. Cloudera Data Science Workbench publishes placeholder parcels for other operating systems as well. However, note that these do not work and have only been included to support mixed-OS clusters.

<table>
<thead>
<tr>
<th>Version</th>
<th>Remote Parcel Repository URL</th>
</tr>
</thead>
<tbody>
<tr>
<td>Cloudera Data Science Workbench 1.5.0</td>
<td><a href="https://archive.cloudera.com/cdsw1/1.5.0/parcels/">https://archive.cloudera.com/cdsw1/1.5.0/parcels/</a></td>
</tr>
</tbody>
</table>

4. Click Save Changes.

5. Go to the Hosts > Parcels page. The external parcel should now appear in the set of parcels available for download.

4. Click Download. Once the download is complete, click Distribute to distribute the parcel to all the CDH hosts in your cluster. Then click Activate. For more detailed information on each of these tasks, see Managing Parcels.
For airgapped installations, create your own local repository, put the Cloudera Data Science Workbench parcel there, and then configure the Cloudera Manager Server to target this newly-created repository.

Add the Cloudera Data Science Workbench Service

Perform the following steps to add the Cloudera Data Science Workbench service to your cluster:

1. Log into the Cloudera Manager Admin Console.
2. On the Home > Status tab, click to the right of the cluster name and select Add a Service to launch the wizard. A list of services will be displayed.
3. Select the Cloudera Data Science Workbench service and click Continue.
4. Select the services which the new CDSW service should depend on. At a minimum, the HDFS, Spark 2, and YARN services are required for the CDSW service to run successfully. Click Continue.
   (Required for CDH 6) If you want to run SparkSQL workloads, you must also add the Hive service as a dependency.
5. Assign Cloudera Data Science Workbench roles to gateway hosts.
   Master
   Assign the Master role to a gateway host that is the designated Master host. This is the host that should have the Application Block Device mounted to it.
   Worker
   Assign the Worker role to any other gateway hosts that will be used for Cloudera Data Science Workbench. Note that Worker hosts are not required for a fully-functional Cloudera Data Science Workbench deployment. For proof-of-concept deployments, you can deploy a 1-host cluster with just a Master host. The Master host can run user workloads just as a worker host can.
   Even if you are setting up a multi-host deployment, do not assign the Master and Worker roles to the same host. By default, the Master host doubles up to perform both functions: those of the Master, outlined here, and those of a worker.

Docker Daemon

This role runs underlying Docker processes on all Cloudera Data Science Workbench hosts. The Docker Daemon role must be assigned to every Cloudera Data Science Workbench gateway host.

On First Run, Cloudera Manager will automatically assign this role to each Cloudera Data Science Workbench gateway host. However, if any more hosts are added or reassigned to Cloudera Data Science Workbench, you must explicitly assign the Docker Daemon role to them.

Application

This role runs the Cloudera Data Science Workbench application. This role runs only on the CDSW Master host.

On First Run, Cloudera Manager will assign the Application role to the host running the Cloudera Data Science Workbench Master role. The Application role is always assigned to the same host as the Master. Consequently, this role must never be assigned to a Worker host.

The following image shows the role assignments for a Cloudera Data Science Workbench Master host and Worker host:
6. Configure the following parameters and click **Continue**.

<table>
<thead>
<tr>
<th>Properties</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Cloudera Data Science Workbench Domain</strong></td>
<td>DNS domain configured to point to the master host. If the previously configured DNS subdomain entries are <code>cdsw.&lt;your_domain&gt;.com</code> and <code>*.cdsw.&lt;your_domain&gt;.com</code>, then this parameter should be set to <code>cdsw.&lt;your_domain&gt;.com</code>. Users' browsers contact the Cloudera Data Science Workbench web application at <code>http://cdsw.&lt;your_domain&gt;.com</code>. This domain for DNS only and is unrelated to Kerberos or LDAP domains.</td>
</tr>
<tr>
<td><strong>Master Node IPv4 Address</strong></td>
<td>IPv4 address for the master host that is reachable from the worker host. <strong>By default, this field is left blank and Cloudera Manager uses the IPv4 address of the Master host.</strong> Within an AWS VPC, set this parameter to the internal IP address of the master host; for instance, if your hostname is <code>ip-10-251-50-12.ec2.internal</code>, set this property to the corresponding IP address, 10.251.50.12.</td>
</tr>
<tr>
<td><strong>Install Required Packages</strong></td>
<td>When this parameter is enabled, the <strong>Prepare Node</strong> command will install all the required package dependencies on First Run. If you choose to disable this property,</td>
</tr>
</tbody>
</table>
you must manually install the following packages on all gateway hosts running Cloudera Data Science Workbench roles:

- nfs-utils
- libseccomp
- lvm2
- bridge-utils
- libtool-ltdl
- iptables
- rsync
- policycoreutils-python
- selinux-policy-base
- selinux-policy-targeted
- ntp
- ebtables
- bind-utils
- nmap-ncat
- openssl
- e2fsprogs
- redhat-lsb-core
- socat

**Docker Block Device**

Block device(s) for Docker images. Use the full path to specify the image(s), for instance, /dev/xvde.

The Cloudera Data Science Workbench installer will format and mount Docker on each gateway host that is assigned the Docker Daemon role. Do not mount these block devices prior to installation.

7. The wizard will now begin a First Run of the Cloudera Data Science Workbench service. This includes deploying client configuration for HDFS, YARN and Spark 2, installing the package dependencies on all hosts, and formatting the Docker block device. The wizard will also assign the Application role to the host running Master and the Docker Daemon role to all the gateway hosts running Cloudera Data Science Workbench.

8. Once the First Run command has completed successfully, click Finish to go back to the Cloudera Manager home page.

**Create the Administrator Account**

After your installation is complete, set up the initial administrator account. Go to the Cloudera Data Science Workbench web application at http://cdsw.<your_domain>.com.

You must access Cloudera Data Science Workbench from the Cloudera Data Science Workbench Domain configured when setting up the service, and not the hostname of the master host. Visiting the hostname of the master host will result in a 404 error.

The first account that you create becomes the site administrator. You may now use this account to create a new project and start using the workbench to run data science workloads. For a brief example, see Getting Started with the Cloudera Data Science Workbench.

**Next Steps**

As a site administrator, you can invite new users, monitor resource utilization, secure the deployment, and upload a license key for the product. For more details on these tasks, see the Administration and Security guides.

You can also start using the product by configuring your personal account and creating a new project. For a quickstart that walks you through creating and running a simple template project, see Getting Started with Cloudera Data Science Workbench on page 98. For more details on collaborating with teams, working on projects, and sharing results, see the Managing Cloudera Data Science Workbench Users on page 102.
Installing Cloudera Data Science Workbench 1.5.x Using Packages

**Note:** Instructions on how to deploy Cloudera Data Science Workbench on Hortonworks Data Platform clusters are available here: Deploying Cloudera Data Science Workbench 1.5.x on Hortonworks Data Platform on page 85.

Use the following steps to install the latest Cloudera Data Science Workbench 1.5.x using RPM packages.

**Prerequisites**

Before you begin installing Cloudera Data Science Workbench, make sure you have completed the steps to configure your hosts and block devices.

**Configure Gateway Hosts Using Cloudera Manager**

Cloudera Data Science Workbench hosts must be added to your CDH cluster as gateway hosts, with gateway roles properly configured. To configure gateway hosts:

1. If you have not already done so and plan to use PySpark, install either the Anaconda parcel or Python (versions 2.7.11 and 3.6.1) on your CDH cluster. For more information see, Python Supported Versions on page 56.

2. Configure Apache Spark on your gateway hosts.
   
   a. **(CDH 5 only)** Install and configure the CDS 2.x Powered by Apache Spark parcel and CSD. For instructions, see Installing CDS 2.x Powered by Apache Spark.

   **Important:** Do not install CDS 2.x if you are using CDH 6. Spark 2 ships as part of the CDH 6 package; the add-on parcel is no longer required. To see which version of Spark 2 ships with CDH, refer the CDH 6 Packaging documentation.

   b. **(Required for CDH 5 and CDH 6)** To be able to use Spark 2, each user must have their own /home directory in HDFS. If you sign in to Hue first, these directories will automatically be created for you. Alternatively, you can have cluster administrators create these directories.

   ```bash
   hdfs dfs -mkdir /user/<username>
   hdfs dfs -chown <username>:<username> /user/<username>
   ```

   If you are using CDS 2.3 release 2 (or higher), review the associated known issues here: CDS Powered By Apache Spark on page 46.

3. Use Cloudera Manager to create add gateway hosts to your CDH cluster.
   
   1. Create a new host template that includes gateway roles for HDFS, YARN, and Spark 2.

   **(Required for CDH 6)** If you want to run SparkSQL workloads, you must also add the Hive gateway role to the template.

   2. Use the instructions at Adding a Host to the Cluster to add gateway hosts to the cluster. Apply the template created in the previous step to these gateway hosts. If your cluster is kerberized, confirm that the krb5.conf file on your gateway hosts is correct.

4. Test Spark 2 integration on the gateway hosts.
   
   1. SSH to a gateway host.
   2. If your cluster is kerberized, run kinit to authenticate to the CDH cluster’s Kerberos Key Distribution Center.

   The Kerberos ticket you create is not visible to Cloudera Data Science Workbench users.

   3. Submit a test job to Spark by executing the following command:
Install Cloudera Data Science Workbench on the Master Host

Use the following steps to install Cloudera Data Science Workbench on the master host. Note that airgapped clusters and non-airgapped clusters use different files for installation.

1. Download the Cloudera Data Science Workbench repo file and/or RPM.

<table>
<thead>
<tr>
<th>Red Hat 7 Repository File</th>
<th>Link to RPM</th>
</tr>
</thead>
<tbody>
<tr>
<td>For non-airgapped installations, download this file and save it to <code>/etc/yum.repos.d/</code>: cloudera-cdsworkbench.repo</td>
<td>For airgapped installations, download the Cloudera Data Science Workbench RPM file: Cloudera Data Science Workbench 1.5.0</td>
</tr>
</tbody>
</table>

Important: Make sure all Cloudera Data Science Workbench hosts (master and worker) are running the same version of Cloudera Data Science Workbench.
2. Skip this step for airgapped installations. Add the Cloudera Public GPG repository key. This key verifies that you are downloading genuine packages.

```sh
sudo rpm --import https://archive.cloudera.com/cdsw1/1.5.0/redhat7/yum/RPM-GPG-KEY-cloudera
```

3. **Non-airgapped Installation** - Install the latest RPM with the following command:

```sh
sudo yum install cloudera-data-science-workbench
```

**Airgapped Installation** - Copy the RPM downloaded in the previous step to the appropriate gateway host. Then, use the complete filename to install the package. For example:

```sh
sudo yum install cloudera-data-science-workbench-1.5.0.12345.rpm
```

For guidance on any warnings displayed during the installation process, see [Understanding Installation Warnings](#) on page 247.

4. Edit the configuration file at `/etc/cdsw/config/cdsw.conf`. The following table lists the configuration properties that can be configured in `cdsw.conf`.

<table>
<thead>
<tr>
<th>Properties</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Required Configuration</strong></td>
<td></td>
</tr>
<tr>
<td>DOMAIN</td>
<td>Wildcard DNS domain configured to point to the master host.</td>
</tr>
<tr>
<td></td>
<td>If the wildcard DNS entries are configured as <code>cdsw.&lt;company&gt;.com</code> and</td>
</tr>
<tr>
<td></td>
<td><code>*.cdsw.&lt;company&gt;.com</code>, then <code>DOMAIN</code> should be set to <code>cdsw.&lt;company&gt;.com</code>.</td>
</tr>
<tr>
<td></td>
<td>Users’ browsers should then contact the Cloudera Data Science Workbench web</td>
</tr>
<tr>
<td></td>
<td>application at <code>http://cdsw.&lt;company&gt;.com</code>. This domain is for DNS and is</td>
</tr>
<tr>
<td></td>
<td>unrelated to Kerberos or LDAP domains.</td>
</tr>
<tr>
<td>MASTER_IP</td>
<td>IPv4 address for the master host that is reachable from the worker hosts.</td>
</tr>
<tr>
<td></td>
<td>Within an AWS VPC, <code>MASTER_IP</code> should be set to the internal IP address of</td>
</tr>
<tr>
<td></td>
<td>the master host; for instance, if your hostname is <code>ip-10-251-50-12.ec2.internal</code>, set <code>MASTER_IP</code> to the corresponding IP address, <code>10.251.50.12</code>.</td>
</tr>
<tr>
<td>DISTRO</td>
<td>The Hadoop distribution installed on the cluster. Set this property to CDH.</td>
</tr>
<tr>
<td>DOCKER_BLOCK_DEVICES</td>
<td>Block device(s) for Docker images (space separated if there are multiple).</td>
</tr>
<tr>
<td></td>
<td>Use the full path to specify the image(s), for instance, <code>/dev/xvde</code>.</td>
</tr>
<tr>
<td>JAVA_HOME</td>
<td>Path where Java is installed on the Cloudera Data Science Workbench hosts.</td>
</tr>
<tr>
<td></td>
<td>The value for <code>JAVA_HOME</code> depends on whether you are using JDK or JRE. For</td>
</tr>
<tr>
<td></td>
<td>example, if you’re using JDK 1.8.162, set <code>JAVA_HOME</code> to <code>/usr/java/jdk1.8.0_162</code>. If you are only using JRE, set it to <code>/usr/java/jdk1.8.0_162/jre</code>.</td>
</tr>
<tr>
<td></td>
<td>Note that Spark 2.2 (and higher) requires JDK 1.8. For more details on the</td>
</tr>
<tr>
<td></td>
<td>specific versions of Oracle JDK recommended for CDH and Cloudera Manager</td>
</tr>
<tr>
<td></td>
<td>clusters, see the Cloudera Product Compatibility Matrix - Supported JDK</td>
</tr>
<tr>
<td></td>
<td>Versions.</td>
</tr>
</tbody>
</table>

<p>| Optional Configuration      |                                                                             |
| APPLICATION_BLOCK_DEVICE    | <strong>(Master Host Only)</strong> Configure a block device for application state.     |
|                             | If this property is left blank, the filesystem mounted at <code>/var/lib/cdsw</code> on  |
|                             | the master host will be used to store all user data. For production         |
|                             | deployments, Cloudera                                                      |</p>
<table>
<thead>
<tr>
<th>Properties</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>strongly recommends you use this option with a dedicated SSD block device for the /var/lib/cdsw mount.</td>
</tr>
<tr>
<td></td>
<td><em>(Not recommended)</em> If set, Cloudera Data Science Workbench will format the provided block device as ext4, mount it to /var/lib/cdsw, and store all user data on it. This option has only been provided for proof-of-concept setups, and Cloudera is not responsible for any data loss. Use the full path to specify the mount point, for instance, /dev/xvdf.</td>
</tr>
<tr>
<td>RESERVE_MASTER</td>
<td>Set this property to <em>true</em> to reserve the master host for Cloudera Data Science Workbench’s internal components and services such as Livelog, the PostgreSQL database, and so on. User workloads will now run exclusively on worker hosts, while the master is reserved for internal application services. Note that <em>this property is not yet available as a configuration property in Cloudera Manager</em>. However, you can use an Advanced Configuration Snippet (Safety Valve) to configure this as described here: Reserving the Master Host for Internal CDSW Components on page 209.</td>
</tr>
<tr>
<td>DISTRO_DIR</td>
<td>Path where the Hadoop distribution is installed on the Cloudera Data Science Workbench hosts. For CDH clusters, the default location of the parcel directory is /opt/cloudera/parcels. Specify this property only if you are using a non-default location.</td>
</tr>
<tr>
<td>ANACONDA_DIR</td>
<td>Path where the Anaconda package is installed. On CDH clusters, Anaconda is installed as a parcel in Cloudera Manager. Therefore, this parameter does not apply and must be left blank.</td>
</tr>
<tr>
<td>TLS_ENABLE</td>
<td>Enable and enforce HTTPS (TLS/SSL) for web access. Set to <em>true</em> to enable and enforce HTTPS access to the web application. You can also set this property to <em>true</em> to enable external TLS termination. For more details on TLS termination, see Enabling TLS/SSL for Cloudera Data Science Workbench on page 231.</td>
</tr>
<tr>
<td>TLS_CERT</td>
<td>Certificate and private key for internal TLS termination. Setting <code>TLS_CERT</code> and <code>TLS_KEY</code> will enable internal TLS termination. You must also set <code>TLS_ENABLE</code> to <em>true</em> above to enable and enforce internal termination. Set these only if you are not terminating TLS externally. Make sure you specify the full path to the certificate and key files, which must be in PEM format. For details on certificate requirements and enabling TLS termination, see Enabling TLS/SSL for Cloudera Data Science Workbench on page 231.</td>
</tr>
<tr>
<td>Properties</td>
<td>Description</td>
</tr>
<tr>
<td>---------------------</td>
<td>-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------</td>
</tr>
<tr>
<td>HTTP_PROXY</td>
<td>If your deployment is behind an HTTP or HTTPS proxy, set the respective HTTP_PROXY or HTTPS_PROXY property in <code>/etc/cdsw/config/cdsw.conf</code> to the hostname of the proxy you are using.</td>
</tr>
<tr>
<td>HTTPS_PROXY</td>
<td></td>
</tr>
<tr>
<td></td>
<td><code>HTTP_PROXY=&quot;&lt;http://proxy_host&gt;:&lt;proxy-port&gt;&quot;</code></td>
</tr>
<tr>
<td>HTTPS_PROXY</td>
<td></td>
</tr>
<tr>
<td></td>
<td><code>HTTPS_PROXY=&quot;&lt;http://proxy_host&gt;:&lt;proxy_port&gt;&quot;</code></td>
</tr>
<tr>
<td></td>
<td>If you are using an intermediate proxy such as Cntlm to handle NTLM authentication, add the Cntlm proxy address to the HTTP_PROXY or HTTPS_PROXY fields in cdsw.conf.</td>
</tr>
<tr>
<td></td>
<td><code>HTTP_PROXY=&quot;http://localhost:3128&quot;</code></td>
</tr>
<tr>
<td></td>
<td><code>HTTPS_PROXY=&quot;http://localhost:3128&quot;</code></td>
</tr>
<tr>
<td></td>
<td>If the proxy server uses TLS encryption to handle connection requests, you will need to add the proxy’s root CA certificate to your host’s store of trusted certificates. This is because proxy servers typically sign their server certificate with their own root certificate. Therefore, any connection attempts will fail until the Cloudera Data Science Workbench host trusts the proxy's root CA certificate. If you do not have access to your proxy’s root certificate, contact your Network / IT administrator. To enable trust, copy the proxy’s root certificate to the trusted CA certificate store (ca-trust) on the Cloudera Data Science Workbench host.</td>
</tr>
<tr>
<td></td>
<td><code>cp /tmp/&lt;proxy-root-certificate&gt;.crt /etc/pki/ca-trust/source/anchors/</code></td>
</tr>
<tr>
<td></td>
<td>Use the following command to rebuild the trusted certificate store.</td>
</tr>
<tr>
<td></td>
<td><code>update-ca-trust extract</code></td>
</tr>
<tr>
<td>ALL_PROXY</td>
<td>If a SOCKS proxy is in use, set to socks5://&lt;host&gt;:&lt;port&gt;/.</td>
</tr>
<tr>
<td>NO_PROXY</td>
<td>Comma-separated list of hostnames that should be skipped from the proxy.</td>
</tr>
<tr>
<td></td>
<td>Starting with version 1.4, if you have defined a proxy in the HTTP_PROXY(S) or ALL_PROXY properties, Cloudera Data Science Workbench automatically appends the following list of IP addresses to the NO_PROXY configuration. Note that this is the minimum required configuration for this field.</td>
</tr>
<tr>
<td></td>
<td>This list includes 127.0.0.1, localhost, and any private Docker registries and HTTP services inside the firewall that Cloudera Data Science Workbench users might want to access from the engines.</td>
</tr>
<tr>
<td></td>
<td>&quot;127.0.0.1,localhost,100.66.0.1,100.66.0.2,100.66.0.3,100.66.0.4,100.66.0.5,100.66.0.6,100.66.0.7,100.66.0.8,100.66.0.9,100.66.0.10,100.66.0.11,100.66.0.12,100.66.0.13,100.66.0.14,100.66.0.15,100.66.0.16,100.66.0.17,100.66.0.18,100.66.0.19,100.66.0.20,100.66.0.21,100.66.0.22,100.66.0.23,100.66.0.24,100.66.0.25,100.66.0.26,100.66.0.27,100.66.0.28,100.66.0.29,100.66.0.30,100.66.0.31,100.66.0.32,100.66.0.33,100.66.0.34,100.66.0.35,100.66.0.36,100.66.0.37,100.66.0.38,100.66.0.39,100.66.0.40,100.66.0.41,100.66.0.42,100.66.0.43,100.66.0.44,100.66.0.45,100.66.0.46,100.66.0.47,100.66.0.48,100.66.0.49,100.66.0.50,100.77.0.10,100.77.0.128,100.77.0.129,100.77.0.130,100.77.0.131,100.77.0.132,100.77.0.133,100.77.0.134,100.77.0.135,100.77.0.136,100.77.0.137,100.77.0.138,100.77.0.139&quot;</td>
</tr>
<tr>
<td>NVIDIA_GPU_ENABLE</td>
<td>Set this property to true to enable GPU support for Cloudera Data Science Workbench workloads. When this property is enabled on a host is equipped with</td>
</tr>
</tbody>
</table>
### Properties

<table>
<thead>
<tr>
<th>Properties</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>GPU hardware, the GPU(s) will be available for use by Cloudera Data Science Workbench hosts. If this property is set to <code>true</code> on a host that does not have GPU support, there will be no effect. By default, this property is set to <code>false</code>. For detailed instructions on how to enable GPU-based workloads on Cloudera Data Science Workbench, see Using NVIDIA GPUs for Cloudera Data Science Workbench Projects on page 117.</td>
</tr>
<tr>
<td>NVIDIA_LIBRARY_PATH</td>
<td>Complete path to the NVIDIA driver libraries.</td>
</tr>
</tbody>
</table>

5. Initialize and start Cloudera Data Science Workbench.

```bash
cdsw init
```

The application will take a few minutes to bootstrap. You can watch the status of application installation and startup with `watch cdsw status`.

(Optional) Install Cloudera Data Science Workbench on Worker Hosts

**Note:** Worker hosts are not required for a fully-functional Cloudera Data Science Workbench deployment. For proof-of-concept deployments, you can deploy a 1-host cluster with just a Master host. The Master host can run user workloads just as a worker host can.

Cloudera Data Science Workbench supports adding and removing additional worker hosts at any time. Worker hosts allow you to transparently scale the number of concurrent workloads users can run.

Use the following steps to add worker hosts to Cloudera Data Science Workbench. **Note that airgapped clusters and non-airgapped clusters use different files for installation.**

1. Download the Cloudera Data Science Workbench repo file and/or RPM.

   **Red Hat 7 Repository File**

   **Link to RPM**

   For non-airgapped installations, download this file and save it to `/etc/yum.repos.d/:

   ```
   cloudera-cdsw.repo
   ```

   For airgapped installations, download the Cloudera Data Science Workbench RPM file:

   ```
   Cloudera Data Science Workbench 1.5.0
   ```

   **Important:** Make sure all Cloudera Data Science Workbench hosts (master and worker) are running the same version of Cloudera Data Science Workbench.

2. Skip this step for airgapped installations. Add the Cloudera Public GPG repository key. This key verifies that you are downloading genuine packages.

   ```bash
   sudo rpm --import https://archive.cloudera.com/cdsw1/1.5.0/redhat7/yum/RPM-GPG-KEY-cloudera
   ```

3. **Non-airgapped Installation** - Install the latest RPM with the following command:

   ```bash
   sudo yum install cloudera-data-science-workbench
   ```
**Airgapped Installation** - Copy the RPM downloaded in the previous step to the appropriate gateway host. Then, use the complete filename to install the package. For example:

```
sudo yum install cloudera-data-science-workbench-1.5.0.12345.rpm
```

For guidance on any warnings displayed during the installation process, see Understanding Installation Warnings on page 247.

4. Copy `cdsw.conf` file from the master host:

```
scp root@cdsw-host-1.<company>.com:/etc/cdsw/config/cdsw.conf /etc/cdsw/config/cdsw.conf
```

After initialization, the `cdsw.conf` file includes a generated bootstrap token that allows worker hosts to securely join the cluster. You can get this token by copying the configuration file from master and ensuring it has 600 permissions.

If your hosts have heterogeneous block device configurations, modify the Docker block device settings in the worker host configuration file after you copy it. Worker hosts do not need application block devices, which store the project files and database state, and this configuration option is ignored.

5. Create `/var/lib/cdsw` on the worker host. This directory must exist on all worker hosts. Without it, the next step that registers the worker host with the master will fail.

   Unlike the master host, the `/var/lib/cdsw` directory on worker hosts does not need to be mounted to an Application Block Device. It is only used to store CDH client configuration on workers.

6. On the `worker` host, run the following command to add the host to the cluster:

```
cdsw join
```

   This causes the worker hosts to register themselves with the Cloudera Data Science Workbench master host and increase the available pool of resources for workloads.

7. Return to the master host and verify the host is registered with this command:

```
cdsw status
```

---

**Create the Administrator Account**

Installation typically takes 30 minutes, although it might take an additional 60 minutes for the R, Python, and Scala engine to be available on all hosts.

After your installation is complete, set up the initial administrator account. Go to the Cloudera Data Science Workbench web application at `http://cdsw.<company>.com`.

**Note:** You must access Cloudera Data Science Workbench from the `DOMAIN` configured in `cdsw.conf`, and not the hostname of the master host. Visiting the hostname of the master host will result in a 404 error.

The first account that you create becomes the site administrator. You may now use this account to create a new project and start using the workbench to run data science workloads. For a brief example, see Getting Started with the Cloudera Data Science Workbench.

---

**Next Steps**

As a site administrator, you can invite new users, monitor resource utilization, secure the deployment, and upload a license key for the product. For more details on these tasks, see the Administration and Security guides.

You can also start using the product by configuring your personal account and creating a new project. For a quickstart that walks you through creating a simple template project, see Getting Started with Cloudera Data Science Workbench.
Installing Cloudera Data Science Workbench 1.5.x on CDH

on page 98. For more details on collaborating with teams, working on projects, and sharing results, see the Managing Cloudera Data Science Workbench Users on page 102.
Upgrading to the Latest Version of Cloudera Data Science Workbench 1.5.x on CDH

This topic walks you through the upgrade paths available for Cloudera Data Science Workbench 1.5.x. Depending on your existing deployment, choose from one of the upgrade paths listed in the following table.

**Upgrading from CDH 5 > CDH 6** - If you are currently running Cloudera Data Science Workbench on CDH 5 and want to upgrade your cluster to CDH 6, also see Upgrading Cloudera Data Science Workbench from CDH 5 to CDH 6 on page 75.

<table>
<thead>
<tr>
<th>Upgrade Path</th>
<th>Link to Instructions</th>
</tr>
</thead>
<tbody>
<tr>
<td>CSD &gt; CSD</td>
<td>Upgrading Cloudera Data Science Workbench 1.5.x Using Cloudera Manager on page 76</td>
</tr>
<tr>
<td>RPM &gt; CSD</td>
<td>Migrating from an RPM-based Deployment to the Latest 1.5.x CSD on page 79</td>
</tr>
<tr>
<td>RPM &gt; RPM</td>
<td>Upgrading Cloudera Data Science Workbench 1.5.x Using Packages on page 82</td>
</tr>
</tbody>
</table>

Note that you cannot use Cloudera Manager for this upgrade path.

**Upgrading Cloudera Data Science Workbench from CDH 5 to CDH 6**

This section provides a general outline on how to go about upgrading your Cloudera Data Science Workbench cluster from CDH 5 to CDH 6. Refer the relevant linked Cloudera Manager and CDH upgrade documentation for the detailed steps required for this procedure.

Starting with version 1.5, Cloudera Data Science Workbench publishes two separate CSD files: one for CDH 5 and one for CDH 6. Check the CSD file name to ensure that you are using the correct CSD file for your cluster. For example:

- **CDH 6** - CLOUDERA_DATA_SCIENCE_WORKBENCH_CDH6_1.x.y.jar
- **CDH 5** - CLOUDERA_DATA_SCIENCE_WORKBENCH_CDH5_1.x.y.jar

Use the following path to upgrade from running a CSD-based Cloudera Data Science Workbench deployment on CDH 5 to running on CDH 6:

1. Upgrade to Cloudera Manager 6.1 (or higher).
2. Stop Cloudera Data Science Workbench.
3. Download both Cloudera Data Science Workbench CSDs. Log on to the Cloudera Manager Server host, and place the CSD files under /opt/cloudera/csd, which is the default location for CSDs.
4. Restart the Cloudera Manager Server.
5. Upgrade to Cloudera Data Science Workbench 1.5 (or higher). During the upgrade process, as you install, distribute, and activate the new parcel, take care to ensure that both the CDSW CSDs (for CDH 5 and CDH 6) are present on the Cloudera Manager Server host.
6. **Use the Cloudera Manager Upgrade Wizard to upgrade from CDH 5 to CDH 6.1 (or higher)**. As part of the upgrade, the wizard will also remove the Spark 2 parcel from all your cluster hosts. With CDH 6, Spark 2 ships as a part of CDH. The add-on parcel is no longer required.

Cloudera Manager 6 can differentiate between the two active CSDs and will select the right one based on the version of CDH running. Because you already have the CDH 6-compatible CSD installed, no further steps are needed.

7. (Optional) Remove any existing CDH 5 CSDs from the Cloudera Manager Server host.

8. Restart Cloudera Data Science Workbench.

Upgrading RPM Deployments from CDH 5 to CDH 6

Cloudera Data Science Workbench ships a single RPM that can be used to install CDSW on both, CDH 5, and CDH 6 clusters. The upgrade path for RPM deployments is:

1. **Upgrade to Cloudera Manager 6.1 (or higher).**

2. **Use the Cloudera Manager Upgrade Wizard to upgrade from CDH 5 to CDH 6 (or higher).** As part of the upgrade, the wizard will also remove the Spark 2 parcel from all your cluster hosts. This is because with CDH 6, Spark 2 ships as a part of CDH. The add-on parcel is no longer required.

3. **Upgrade to the latest Cloudera Data Science Workbench RPM.**

Upgrading Cloudera Data Science Workbench 1.5.x Using Cloudera Manager

This topic describes how to upgrade a CSD and parcel-based deployment to the latest version of Cloudera Data Science Workbench 1.5.x.

1. Before you begin the upgrade process, make sure you read the [Cloudera Data Science Workbench Release Notes](https://cloudera.com/downloads/releases/documentation.html#data-science-workbench) on page 20 relevant to the version you are upgrading to/from.

2. Depending on the version you are upgrading from, perform one of the following steps to stop Cloudera Data Science Workbench:

   - **(Required for Upgrades from CDSW 1.4.2 or lower)** Safely stop Cloudera Data Science Workbench. To avoid running into the data loss issue described in TSB-346, run the `cdsw.protect_stop_restart.sh` script on the master host and follow the sequence of steps as instructed by the script.
     
     The script will first back up your project files to the specified target folder. It will then temporarily move your project files aside to protect against the data loss condition. At that point, it is safe to **stop the CDSW service in Cloudera Manager**.

     After Cloudera Data Science Workbench has stopped, press enter to continue running the script as instructed. It will then move your project files back into place.

   OR

   - **(Upgrading from CDSW 1.4.3 or higher)** Stop the Cloudera Data Science Workbench service in Cloudera Manager.

3. **(Strongly Recommended)** On the master host, backup all your application data that is stored in the `/var/lib/cdsw` directory.

   To create the backup, run the following command on the master host:

   ```bash
   tar cvzf cdsw.tar.gz /var/lib/cdsw/*
   ```
4. **Required for Upgrades from CDSW 1.4.0 - RedHat only** Cloudera Data Science Workbench 1.4.2 (and higher) includes a fix for a slab leak issue found in RedHat kernels. To have this fix go into effect, RedHat users must reboot all Cloudera Data Science Workbench hosts before proceeding with an upgrade from CDSW 1.4.0.

As a precaution, consult your cluster/IT administrator before you start rebooting hosts.

5. Deactivate the existing Cloudera Data Science Workbench parcel. Go to the Cloudera Manager Admin Console. In the top navigation bar, click **Hosts > Parcels**.

Locate the current active CDSW parcel and click **Deactivate**. On the confirmation pop-up, select **Deactivate Only** and click **OK**.

6. Download and save the latest Cloudera Data Science Workbench CSD to the Cloudera Manager Server host.

   a. Download the latest Cloudera Data Science Workbench CSD.

<table>
<thead>
<tr>
<th>Version</th>
<th>Link to CSD</th>
</tr>
</thead>
<tbody>
<tr>
<td>Cloudera Data Science Workbench 1.5.0</td>
<td><a href="https://archive.cloudera.com/cdw1/1.5.0/parcels/Cloudera_Dat_s_Science_Workbench_CDH6_1.5.0.jar">CDH 6 - Cloudera_Dat_s_Science_Workbench_CDH6_1.5.0.jar</a></td>
</tr>
<tr>
<td>CDH 5 - Cloudera_Dat_s_Science_Workbench_CDH5_1.5.0.jar</td>
<td></td>
</tr>
</tbody>
</table>

   b. Log on to the Cloudera Manager Server host, and place the CSD file under /opt/cloudera/csd, which is the default location for CSD files.
   c. Delete any CSD files belonging to older versions of Cloudera Data Science Workbench from /opt/cloudera/csd.

   **This is required because older versions of the CSD will not work with the latest Cloudera Data Science Workbench 1.4 parcel. Make sure your CSD and parcel are always the same version.**

   ![Note:](image)

   If you have previously configured a custom location for CSD files, place the CSD file there, and delete any CSDs belonging to older versions of Cloudera Data Science Workbench.

   For help, refer the Cloudera Manager documentation at [Configuring the Location of Custom Service Descriptor Files](#).

d. Set the CSD file ownership to **cloudera-scm:cloudera-scm** with permission 644.

e. Restart the Cloudera Manager Server:

   ```
   service cloudera-scm-server restart
   ```

   f. Log into the Cloudera Manager Admin Console and restart the Cloudera Management Service.

      a. Select **Clusters > Cloudera Management Service**.
      b. Select **Actions > Restart**.

   7. Distribute and activate the new parcel on your cluster.

      a. Log into the Cloudera Manager Admin Console.
      b. Click **Hosts > Parcels** in the main navigation bar.
      c. **If the latest CDSW parcel is already available in this list, you can skip this step.**

      Add the Cloudera Data Science Workbench parcel repository URL to Cloudera Manager.

      1. On the **Parcels** page, click **Configuration**.
      2. In the **Remote Parcel Repository URLs** list, click the addition symbol to create a new row.
      3. Enter the path to the repository.

<table>
<thead>
<tr>
<th>Version</th>
<th>Remote Parcel Repository URL</th>
</tr>
</thead>
<tbody>
<tr>
<td>Cloudera Data Science Workbench 1.5.0</td>
<td><a href="https://archive.cloudera.com/cdw1/1.5.0/parcels/">https://archive.cloudera.com/cdw1/1.5.0/parcels/</a></td>
</tr>
</tbody>
</table>
4. Click **Save Changes**.

d. Go back to the Hosts > Parcels page. The latest parcel should now appear in the set of parcels available for download. Click **Download**. Once the download is complete, click **Distribute** to distribute the parcel to all the CDH hosts in your cluster. Then click **Activate**. On the pop-up screen, select **Activate Only** and click **OK**. For more detailed information on each of these tasks, see Managing Parcels.

8. Run the **Prepare Node** command on all Cloudera Data Science Workbench hosts.

1. Before you run **Prepare Node**, you must make sure that the command is allowed to install all the required packages on your cluster. This is controlled by the **Install Required Packages** property.

   1. Navigate to the CDSW service.
   2. Click **Configuration**.
   3. Search for the **Install Required Packages** property. If this property is enabled, you can move on to the next step and run **Prepare Node**.

   However, if the property has been disabled, you must either enable it or manually install the following packages on all Cloudera Data Science Workbench gateway hosts:

   ```
   nfs-utils
   libseccomp
   lvm2
   bridge-utils
   libtool-ltdl
   iptables
   rsync
   policycoreutils-python
   selinux-policy-base
   selinux-policy-targeted
   ntp
   ebtables
   bind-utils
   nmap-ncat
   openssl
   e2fsprogs
   redhat-lsb-core
   socat
   ```

2. Run the **Prepare Node** command.

   1. In Cloudera Manager, navigate to the Cloudera Data Science Workbench service.
   2. Click the **Instances** tab.
   3. Use the checkboxes to select all host instances and click **Actions for Selected** (x).
   4. Click **Prepare Node**. Once again, click **Prepare Node** to confirm the action.

9. Log into the Cloudera Manager Admin Console and restart the Cloudera Data Science Workbench service.

   a. On the **Home > Status** tab, click

   ![Dropdown for Restarting Services](image)

   to the right of the **CDSW** service and select **Restart** from the dropdown.

   b. Confirm your choice on the next screen. Note that a complete restart of the service will take time. Even though the CDSW service status shows **Good** Health, the application itself will take some more time to get ready.

10. **Upgrade Projects to Use the Latest Base Engine Images**

    If the release you have just upgraded to includes a new version of the base engine image (see release notes), you will need to manually configure existing projects to use the new engine. Cloudera recommends you do so to take advantage of any new features and bug fixes included in the newly released engine.
To upgrade a project to the new engine, go to the project’s Settings > Engine page and select the new engine from the dropdown. If any of your projects are using custom extended engines, you will need to modify them to use the new base engine image.

Note that this is a required step if you have upgraded to using Cloudera Data Science Workbench on CDH 6.

The base engine image you use must be compatible with the version of CDH you are running. This is especially important if you are running workloads on Spark. Older base engines (v5 and lower) cannot support the latest versions of CDH 6. That is because these engines were configured to point to the Spark 2 parcel. However, on C6 clusters, Spark is now packaged as part of CDH 6 and the separate add-on Spark 2 parcel is no longer supported. If you want to use Spark on C6, you must upgrade your projects to base engine 7 (or higher).

<table>
<thead>
<tr>
<th>Base Engine Versions</th>
<th>CDH 5</th>
<th>CDH 6</th>
</tr>
</thead>
<tbody>
<tr>
<td>Base engines 6 (and lower)</td>
<td>Yes</td>
<td>No</td>
</tr>
<tr>
<td>Base engines 7 (and higher)</td>
<td>Yes</td>
<td>Yes</td>
</tr>
</tbody>
</table>

Table 3: CDSW Base Engine Compatibility for Spark Workloads on CDH 5 and CDH 6

Migrating from an RPM-based Deployment to the Latest 1.5.x CSD

This topic describes how to migrate from an RPM-based deployment to the latest 1.5.x CSD and parcel-based deployment.

1. Before you begin the migration process, make sure you read the Cloudera Data Science Workbench Release Notes on page 20 relevant to the version you are migrating to/from.

2. Save a backup of the Cloudera Data Science Workbench configuration file located at /etc/cdsw/config/cdsw.conf.

3. **(Required for Upgrades from CDSW 1.4.2 or lower)** Safely stop Cloudera Data Science Workbench. To avoid running into the data loss issue described in TSB-346, run the cdsw_protect_stop_restart.sh script on the master host and follow the sequence of steps as instructed by the script.

   The script will first back up your project files to the specified target folder. It will then temporarily move your project files aside to protect against the data loss condition. At that point, it is safe to stop Cloudera Data Science Workbench. To stop Cloudera Data Science Workbench, run the following command on all Cloudera Data Science Workbench hosts (master and workers):

   ```
cdsw reset
   ```

   After Cloudera Data Science Workbench has stopped, press enter to continue running the script as instructed. It will then move your project files back into place.

   **OR**

   ```
cdsw reset
   ```

4. **(Strongly Recommended)** On the master host, backup all your application data that is stored in the /var/lib/cdsw directory.

   To create the backup, run the following command on the master host:

   ```
tar cvzf cdsw.tar.gz /var/lib/cdsw/*
   ```

5. Save a backup of the Cloudera Data Science Workbench configuration file at /etc/cdsw/config/cdsw.conf.
6. **(Required for Upgrades from CDSW 1.4.0 - RedHat only)** Cloudera Data Science Workbench 1.4.2 (and higher) includes a fix for a slab leak issue found in RedHat kernels. To have this fix go into effect, RedHat users must reboot all Cloudera Data Science Workbench hosts before proceeding with an upgrade from CDSW 1.4.0.

As a precaution, consult your cluster/IT administrator before you start rebooting hosts.

7. Uninstall the previous release of Cloudera Data Science Workbench. Perform this step on the master host, as well as all the worker hosts.

   ```bash
   yum remove cloudera-data-science-workbench
   ```

8. Install the latest version of Cloudera Data Science Workbench using the CSD and parcel. Note that when you are configuring role assignments for the Cloudera Data Science Workbench service, the Master role must be assigned to the same host that was running as master prior to the upgrade.

   For installation instructions, see [Installing Cloudera Data Science Workbench 1.5.x Using Packages](on page 67). You might be able to skip the first few steps assuming you have the wildcard DNS domain and block devices already set up.

9. Use your copy of the backup `cdsw.conf` created in Step 3 to recreate those settings in Cloudera Manager by configuring the corresponding properties under the Cloudera Data Science Workbench service.
   
   a. Log into the Cloudera Manager Admin Console.
   
   b. Go to the Cloudera Data Science Workbench service.
   
   c. Click the Configuration tab.
   
   d. The following table lists all the `cdsw.conf` properties and their corresponding Cloudera Manager properties (in bold). Use the search box to bring up the properties you want to modify.
   
   e. Click Save Changes.

<table>
<thead>
<tr>
<th><code>cdsw.conf</code> Property</th>
<th>Corresponding Cloudera Manager Property and Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>TLS_ENABLE</td>
<td>Enable TLS: Enable and enforce HTTPS (TLS/SSL) access to the web application (optional). Both internal and external termination are supported. To enable internal termination, you must also set the TLS Certificate for Internal Termination and TLS Key for Internal Termination parameters. If these parameters are not set, terminate TLS using an external proxy. For more details on TLS termination, see [Enabling TLS/SSL for Cloudera Data Science Workbench](on page 231).</td>
</tr>
<tr>
<td>TLS_CERT</td>
<td>TLS Certificate for Internal Termination, TLS Key for Internal Termination. Complete path to the certificate and private key (in PEM format) to be used for internal TLS termination. Set these parameters only if you are not terminating TLS externally. You must also set the Enable TLS property to enable and enforce termination. The certificate must include both <code>DOMAIN</code> and <code>*.DOMAIN</code> as hostnames. Self-signed certificates are not supported unless trusted fully by clients. Accepting an invalid certificate manually can cause connection failures for unknown subdomains. Set these only if you are not terminating TLS externally. For details on certificate requirements and enabling TLS termination, see [Enabling TLS/SSL for Cloudera Data Science Workbench](on page 231).</td>
</tr>
<tr>
<td>TLS_KEY</td>
<td>HTTP_PROXY, HTTPS Proxy</td>
</tr>
<tr>
<td>HTTP_PROXY</td>
<td>If your deployment is behind an HTTP or HTTPS proxy, set the respective HTTP Proxy or HTTPS Proxy property to the hostname of the proxy you are using. http://&lt;proxy_host&gt;::&lt;proxy-port&gt;</td>
</tr>
</tbody>
</table>

---

80 | Cloudera Data Science Workbench
### Corresponding Cloudera Manager Property and Description

<table>
<thead>
<tr>
<th>cdsw.conf Property</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>or https://&lt;proxy_host&gt;:&lt;proxy_port&gt;</td>
<td>If you are using an intermediate proxy such as Cntlm to handle NTLM authentication, add the Cntlm proxy address to the HTTP Proxy or HTTPS Proxy fields. That is, either <a href="http://localhost:3128">http://localhost:3128</a> or <a href="https://localhost:3128">https://localhost:3128</a> respectively. If the proxy server uses TLS encryption to handle connection requests, you will need to add the proxy's root CA certificate to your host's store of trusted certificates. This is because proxy servers typically sign their server certificate with their own root certificate. Therefore, any connection attempts will fail until the Cloudera Data Science Workbench host trusts the proxy's root CA certificate. If you do not have access to your proxy's root certificate, contact your Network / IT administrator. To enable trust, copy the proxy's root certificate to the trusted CA certificate store (ca-trust) on the Cloudera Data Science Workbench host. Use the following command to rebuild the trusted certificate store. <code>update-ca-trust extract</code></td>
</tr>
<tr>
<td>ALL_PROXY</td>
<td>SOCKS Proxy: If a SOCKS proxy is in use, set this parameter to <code>socks5://&lt;host&gt;:&lt;port&gt;/</code>.</td>
</tr>
<tr>
<td>NO_PROXY</td>
<td>No Proxy: Comma-separated list of hostnames that should be skipped from the proxy. Starting with version 1.4, if you have defined a proxy in the HTTP_PROXY(S) or ALL_PROXY properties, Cloudera Data Science Workbench automatically appends the following list of IP addresses to the NO_PROXY configuration. Note that this is the minimum required configuration for this field. This list includes 127.0.0.1, localhost, and any private Docker registries and HTTP services inside the firewall that Cloudera Data Science Workbench users might want to access from the engines. <code>&quot;127.0.0.1,localhost,100.66.0.1,100.66.0.2,100.66.0.3,100.66.0.4,100.66.0.5,100.66.0.6,100.66.0.7,100.66.0.8,100.66.0.9,100.66.0.10,100.66.0.11,100.66.0.12,100.66.0.13,100.66.0.14,100.66.0.15,100.66.0.16,100.66.0.17,100.66.0.18,100.66.0.19,100.66.0.20,100.66.0.21,100.66.0.22,100.66.0.23,100.66.0.24,100.66.0.25,100.66.0.26,100.66.0.27,100.66.0.28,100.66.0.29,100.66.0.30,100.66.0.31,100.66.0.32,100.66.0.33,100.66.0.34,100.66.0.35,100.66.0.36,100.66.0.37,100.66.0.38,100.66.0.39,100.66.0.40,100.66.0.41,100.66.0.42,100.66.0.43,100.66.0.44,100.66.0.45,100.66.0.46,100.66.0.47,100.66.0.48,100.66.0.49,100.66.0.50,100.77.0.10,100.77.0.128,100.77.0.129,100.77.0.130,100.77.0.131,100.77.0.132,100.77.0.133,100.77.0.134,100.77.0.135,100.77.0.136,100.77.0.137,100.77.0.138,100.77.0.139&quot;</code></td>
</tr>
<tr>
<td>NVIDIA_GPU_ENABLE</td>
<td>Enable GPU Support: When this property is enabled, and the NVIDIA GPU Driver Library Path parameter is set, the GPUs installed on Cloudera Data Science Workbench hosts will be available for use in its workloads. By default, this parameter is disabled. For instructions on how to enable GPU-based workloads on Cloudera Data Science Workbench, see Using NVIDIA GPUs for Cloudera Data Science Workbench Projects on page 117.</td>
</tr>
<tr>
<td>cdsw.conf Property</td>
<td>Corresponding Cloudera Manager Property and Description</td>
</tr>
<tr>
<td>--------------------</td>
<td>--------------------------------------------------------</td>
</tr>
<tr>
<td>NVIDIA_LIBRARY_PATH</td>
<td>NVIDIA GPU Driver Library Path: Complete path to the NVIDIA driver libraries. For instructions on how to create this directory, see Enable Docker NVIDIA Volumes on GPU Hosts on page 119.</td>
</tr>
</tbody>
</table>

10. Cloudera Manager will prompt you to restart the service if needed.

11. Upgrade Projects to Use the Latest Base Engine Images

If the release you have just upgraded to includes a new version of the base engine image (see release notes), you will need to manually configure existing projects to use the new engine. Cloudera recommends you do so to take advantage of any new features and bug fixes included in the newly released engine.

To upgrade a project to the new engine, go to the project’s Settings > Engine page and select the new engine from the dropdown. If any of your projects are using custom extended engines, you will need to modify them to use the new base engine image.

Note that this is a required step if you have upgraded to using Cloudera Data Science Workbench on CDH 6.

The base engine image you use must be compatible with the version of CDH you are running. This is especially important if you are running workloads on Spark. Older base engines (v5 and lower) cannot support the latest versions of CDH 6. That is because these engines were configured to point to the Spark 2 parcel. However, on C6 clusters, Spark is now packaged as part of CDH 6 and the separate add-on Spark 2 parcel is no longer supported. If you want to use Spark on C6, you must upgrade your projects to base engine 7 (or higher).

Table 4: CDSW Base Engine Compatibility for Spark Workloads on CDH 5 and CDH 6

<table>
<thead>
<tr>
<th>Base Engine Versions</th>
<th>CDH 5</th>
<th>CDH 6</th>
</tr>
</thead>
<tbody>
<tr>
<td>Base engines 6 (and lower)</td>
<td>Yes</td>
<td>No</td>
</tr>
<tr>
<td>Base engines 7 (and higher)</td>
<td>Yes</td>
<td>Yes</td>
</tr>
</tbody>
</table>

Upgrading Cloudera Data Science Workbench 1.5.x Using Packages

This topic describes how to upgrade an RPM-based deployment to the latest version of Cloudera Data Science Workbench 1.5.x.

Before you start upgrading Cloudera Data Science Workbench, read the Cloudera Data Science Workbench Release Notes on page 20 relevant to the version you are upgrading to.

1. Depending on the version you are upgrading from, perform one of the following steps to stop Cloudera Data Science Workbench:

   a. **(Required for Upgrades from CDSW 1.4.2 or lower)** Safely stop Cloudera Data Science Workbench. To avoid running into the data loss issue described in TSB-346, run the `cdsw_protect_stop_restart.sh` script on the master host and follow the sequence of steps as instructed by the script.

   The script will first back up your project files to the specified target folder. It will then temporarily move your project files aside to protect against the data loss condition. At that point, it is safe to stop Cloudera Data Science Workbench. To stop Cloudera Data Science Workbench, run the following command on all Cloudera Data Science Workbench hosts (master and workers):

   ```bash
cdsw reset
   ```

   After Cloudera Data Science Workbench has stopped, press enter to continue running the script as instructed. It will then move your project files back into place.

   OR
• **(Upgrading from CDSW 1.4.3 or higher)** Run the following command on all Cloudera Data Science Workbench hosts (master and workers) to stop Cloudera Data Science Workbench.

```bash
cdsw reset
```

2. **(Strongly Recommended)** On the master host, backup all your application data that is stored in the `/var/lib/cdsw` directory.

   To create the backup, run the following command on the master host:

```bash
tar cvzf cdsw.tar.gz /var/lib/cdsw/*
```

3. Save a backup of the Cloudera Data Science Workbench configuration file at `/etc/cdsw/config/cdsw.conf`.

4. **(Required for Upgrades from CDSW 1.4.0 - RedHat only)** Cloudera Data Science Workbench 1.4.2 (and higher) includes a fix for a slab leak issue found in RedHat kernels. To have this fix go into effect, RedHat users must reboot all Cloudera Data Science Workbench hosts before proceeding with an upgrade from CDSW 1.4.0.

   As a precaution, consult your cluster/IT administrator before you start rebooting hosts.

5. Uninstall the previous release of Cloudera Data Science Workbench. Perform this step on the master host, as well as all the worker hosts.

```bash
yum remove cloudera-data-science-workbench
```

6. Install the latest version of Cloudera Data Science Workbench on the master host and on all the worker hosts.

   During the installation process, you might need to resolve certain incompatibilities in `cdsw.conf`. Even though you will be installing the latest RPM, your previous configuration settings in `cdsw.conf` will remain unchanged. Depending on the release you are upgrading from, you will need to modify `cdsw.conf` to ensure it passes the validation checks run by the 1.5.x release.

**Important:**

### Additional Upgrade Notes for version 1.5.x

- **Proxy Configuration** - If you are using a proxy, the IP addresses for the web and Livelog services (100.77.0.129, 100.77.0.130) must be appended to the `NO_PROXY` parameter. They have been included in the installation instructions.

- **cdsw.conf Parameters**
  - `JAVA_HOME` is a required parameter. Make sure you add `JAVA_HOME` to `cdsw.conf` before you start Cloudera Data Science Workbench.
  - Previous versions allowed `MASTER_IP` to be set to a DNS hostname. If you are still using a DNS hostname, switch to an IP address.

To install the latest version of Cloudera Data Science Workbench, follow the same process to install the package as you would for a fresh installation.

- **Install Cloudera Data Science Workbench on the Master Host** on page 68
- **(Optional) Install Cloudera Data Science Workbench on Worker Hosts** on page 72.

7. **Upgrade Projects to Use the Latest Base Engine Images**

   If the release you have just upgraded to includes a new version of the base engine image (see release notes), you will need to manually configure existing projects to use the new engine. Cloudera recommends you do so to take advantage of any new features and bug fixes included in the newly released engine.

   To upgrade a project to the new engine, go to the project’s **Settings > Engine** page and select the new engine from the dropdown. If any of your projects are using custom **extended engines**, you will need to modify them to use the new base engine image.
Note that this is a required step if you have upgraded to using Cloudera Data Science Workbench on CDH 6.

The base engine image you use must be compatible with the version of CDH you are running. This is especially important if you are running workloads on Spark. Older base engines (v5 and lower) cannot support the latest versions of CDH 6. That is because these engines were configured to point to the Spark 2 parcel. However, on C6 clusters, Spark is now packaged as part of CDH 6 and the separate add-on Spark 2 parcel is no longer supported. If you want to use Spark on C6, you must upgrade your projects to base engine 7 (or higher).

Table 5: CDSW Base Engine Compatibility for Spark Workloads on CDH 5 and CDH 6

<table>
<thead>
<tr>
<th>Base Engine Versions</th>
<th>CDH 5</th>
<th>CDH 6</th>
</tr>
</thead>
<tbody>
<tr>
<td>Base engines 6 (and lower)</td>
<td>Yes</td>
<td>No</td>
</tr>
<tr>
<td>Base engines 7 (and higher)</td>
<td>Yes</td>
<td>Yes</td>
</tr>
</tbody>
</table>
Deploying Cloudera Data Science Workbench 1.5.x on Hortonworks Data Platform

Cloudera Data Science Workbench is a secure, self-service enterprise data science platform that lets data scientists manage their own analytics pipelines, thus accelerating machine learning projects from exploration to production. It allows data scientists to bring their existing skills and tools, such as R, Python, and Scala, to securely run computations on data in Hadoop clusters. It enables data science teams to use their preferred data science packages to run experiments with on-demand access to compute resources. Models can be trained, deployed, and managed centrally for increased agility and compliance.

With version 1.5, Cloudera Data Science Workbench now includes direct integration with the Hortonworks Data Platform for a complete machine learning workflow that supports collaborative development and can run both in the public cloud and on-premises.

A detailed overview of the core capabilities of Cloudera Data Science Workbench is available here: Core Capabilities of Cloudera Data Science Workbench.

On this page:

The rest of this topic describes how Cloudera Data Science Workbench can be deployed on HDP. It walks you through a brief architecture overview, the installation requirements, and the limitations associated with such deployments. It also includes instructions on how to install the Cloudera Data Science Workbench package on HDP.

CDSW-on-HDP Architecture Overview

Cloudera Data Science Workbench runs on one or more dedicated gateway / edge hosts on HDP clusters. A gateway host is one that does not have any cluster services running on them. They only run the clients for cluster services (such as the HDFS Client, YARN Client, Spark2 Client and so on). These clients ensure that Cloudera Data Science Workbench has all the libraries and configuration files necessary to securely access the HDP cluster and their respective services.
Cloudera Data Science Workbench does not support running any other services on these gateway hosts. Each gateway host must be dedicated solely to Cloudera Data Science Workbench. This is because user workloads require dedicated CPU and memory, which might conflict with other services running on these hosts.

From the gateway hosts assigned to Cloudera Data Science Workbench, one will serve as the master host, which also runs the CDSW web application, while others will serve as worker hosts. You should note that worker hosts are not required for a fully-functional Cloudera Data Science Workbench deployment. For proof-of-concept deployments you can deploy a 1-host cluster with just a Master host. The Master host can run user workloads just as a worker can.

Refer the following topics for more details on master and worker hosts, and the other components (such as engines for workload execution) that make up Cloudera Data Science Workbench.

**Related Topics:**
- CDSW Master and Worker Hosts
- CDSW Engines: For Python, R, and Scala Workloads

**Supported Platforms and Requirements**

This topic lists the software and hardware configuration required to successfully install and run Cloudera Data Science Workbench on the Hortonworks Data Platform. Cloudera Data Science Workbench and HDP do not support hosts or clusters that do not conform to the requirements listed on this page.

**Platform Requirements**

- Cloudera Data Science Workbench 1.5.x
- Hortonworks Data Platform 2.6.5, 3.1.0
  - Apache Spark 2 - Use the version of Spark that ships with the version of HDP you are running. Refer the HDP component version lists for details: [HDP 2.6.5](#), [HDP 3.1.0](#).
  - Spark 1.x is not supported.

**Operating System Requirements**

Cloudera Data Science Workbench 1.5.x is supported on HDP on the following operating systems:

<table>
<thead>
<tr>
<th>Operating System</th>
<th>Versions</th>
<th>Notes</th>
</tr>
</thead>
<tbody>
<tr>
<td>RHEL / CentOS / Oracle Linux RHCK</td>
<td>7.2, 7.3, 7.4, 7.5</td>
<td>- When IPv6 is disabled, Cloudera Data Science Workbench installations on RHEL/CentOS 7.3 fail due to an issue in kernel versions 3.10.0-514 - 3.10.0-693. For details, see <a href="https://access.redhat.com/solutions/3039771">https://access.redhat.com/solutions/3039771</a>. Cloudera Data Science Workbench installations on RHEL/CentOS 7.2 might fail due to an issue with certain versions of the nfs-utils package. To fix the issue, either downgrade the nfs-utils package or upgrade to a version with the fix. View the complete Red Hat bug report <a href="#">here</a>.</td>
</tr>
<tr>
<td>Oracle Linux (UEK - default)</td>
<td>7.3</td>
<td>-</td>
</tr>
</tbody>
</table>

**Additional OS-level Settings**

- Enable memory cgroups on your operating system.
Deploying Cloudera Data Science Workbench 1.5.x on Hortonworks Data Platform

- Disable swap for optimum stability and performance. For instructions, see Setting the vm.swappiness Linux Kernel Parameter.
- Cloudera Data Science Workbench uses uid 8536 for an internal service account. Make sure that this user ID is not assigned to any other service or user account.
- Cloudera recommends that all users have the max-user-processes ulimit set to at least 65536.
- Cloudera recommends that all users have the max-open-files ulimit set to 1048576.

Java Requirements

The entire cluster, including Cloudera Data Science Workbench gateway hosts, must use either Oracle JDK 8 or OpenJDK 8.

For Red Hat/CentOS deployments, Java Cryptography Extension (JCE) Unlimited Strength Jurisdiction must be enabled on the Cloudera Data Science Workbench gateway hosts. This is to ensure that JDK uses the same default encryption type (aes256-cts) as that used by Red Hat/CentOS operating systems, Kerberos, and the rest of the cluster. For instructions, see Installing the JCE on Ambari.

The JAVA_HOME configuration property must be configured as part of the CDSW installation process and must match the JAVA_HOME environmental variable configured for your HDP cluster. If you need to modify JAVA_HOME after the fact, restart the master and worker hosts to have the changes go into effect.

Related topics:
- Changing the JDK version on an Existing HDP Cluster

Network and Security Requirements

Important: Make sure that your networking/security settings on Cloudera Data Science Workbench gateway hosts are not being overwritten behind-the-scenes, either by any automated scripts, or by other high-priority configuration such as /etc/sysctl.conf, /etc/krb5.conf, or /etc/hosts.deny.

- All Cloudera Data Science Workbench gateway hosts must be part of the same datacenter and use the same network. Hosts from different data-centers or networks can result in unreliable performance.
- A wildcard subdomain such as *.cdsw.company.com must be configured. Wildcard subdomains are used to provide isolation for user-generated content.
- Disable all pre-existing iptables rules. While Kubernetes makes extensive use of iptables, it’s difficult to predict how pre-existing iptables rules will interact with the rules inserted by Kubernetes. Therefore, Cloudera recommends you use the following commands to disable all pre-existing rules before you proceed with the installation.

```
sudo iptables -P INPUT ACCEPT
sudo iptables -P FORWARD ACCEPT
sudo iptables -P OUTPUT ACCEPT
sudo iptables -t nat -F
sudo iptables -t mangle -F
sudo iptables -F
sudo iptables -X
```

- Cloudera Data Science Workbench sets the following sysctl options in /etc/sysctl.d/k8s.conf:

```
- net.bridge.bridge-nf-call-iptables=1
- net.bridge.bridge-nf-call-ip6tables=1
- net.ipv4.ip_forward=1
```

Underlying components of Cloudera Data Science Workbench (Docker, Kubernetes, and NFS) require these options to work correctly. Make sure they are not overridden by high-priority configuration such as /etc/sysctl.conf.

- SELinux must either be disabled or run in permissive mode.
- Multi-homed networks are supported with Cloudera Data Science Workbench.
Firewall restrictions must be disabled across Cloudera Data Science Workbench or HDP hosts. Internally, the Cloudera Data Science Workbench master and worker hosts require full connectivity with no firewalls. Externally, end users connect to Cloudera Data Science Workbench exclusively through a web server running on the master host, and, therefore, do not need direct access to any other internal Cloudera Data Science Workbench or HDP services.

Review the complete list of ports required by Cloudera Data Science Workbench at [Ports Used By Cloudera Data Science Workbench](#) on page 216.

- Non-root SSH access is not allowed on Cloudera Data Science Workbench hosts.
- `localhost` must resolve to `127.0.0.1`.
- Cloudera Data Science Workbench does not support DNS servers running on `127.0.0.1:53`. This IP address resolves to the container localhost within Cloudera Data Science Workbench containers. As a workaround, use either a non-loopback address or a remote DNS server.

Cloudera Data Science Workbench does not support hosts or clusters that do not conform to these restrictions.

### Hardware Requirements

Cloudera Data Science Workbench hosts are added to your cluster as gateway hosts. The table below lists the recommended minimum hardware configuration for Cloudera Data Science Workbench gateway hosts.

<table>
<thead>
<tr>
<th>Resource Type</th>
<th>Master</th>
<th>Workers</th>
<th>Notes</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>CPU</strong></td>
<td>16+ CPU (vCPU) cores</td>
<td>16+ CPU (vCPU) cores</td>
<td></td>
</tr>
<tr>
<td><strong>RAM</strong></td>
<td>32+ GB</td>
<td>32+ GB</td>
<td></td>
</tr>
<tr>
<td><strong>Disk Space</strong></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td><strong>Root Volume</strong></td>
<td>100+ GB</td>
<td>100+ GB</td>
<td>If you are going to partition the root volume, make sure you allocate at least 20 GB to <code>/</code> so that the installer can proceed without running out of space.</td>
</tr>
<tr>
<td><strong>Application Block Device</strong></td>
<td>1 TB</td>
<td>-</td>
<td>The Application Block Device is only required on the Master where it is mounted to <code>/var/lib/cdsw</code>. You will be asked to create a <code>/var/lib/cdsw</code> directory on all the Worker hosts during the installation process. However, they do not need to be mounted to a block device. This directory is only used to store client configuration for HDP cluster services on Workers.</td>
</tr>
<tr>
<td><strong>Docker Block Device</strong></td>
<td>1 TB</td>
<td>1 TB</td>
<td>The Docker Block Device is required on all Master and Worker hosts.</td>
</tr>
</tbody>
</table>
Python Supported Versions

The default Cloudera Data Science Workbench engine includes **Python 2.7.11** and **Python 3.6.1**. To use PySpark within the HDP cluster, the Spark executors must have access to a matching version of Python. For many common operating systems, the default system Python will not match the minor release of Python included in Cloudera Data Science Workbench.

To ensure that the Python versions match, Python can either be installed on every HDP host or made available per job run using Spark’s ability to distribute dependencies. Given the size of a typical isolated Python environment and the desire to avoid repeated uploads from gateway hosts, Cloudera recommends installing Python 2.7 and 3.6 on the cluster if you are using PySpark with lambda functions.

You can install Python 2.7 and 3.6 on the cluster using any method and set the corresponding `PYSPARK_PYTHON` environment variable in your project. Cloudera Data Science Workbench includes a separate environment variable for Python 3 sessions called `PYSPARK3_PYTHON`. Python 2 sessions continue to use the default `PYSPARK_PYTHON` variable. This will allow you to run Python 2 and Python 3 sessions in parallel without either variable being overridden by the other.

**Anaconda**

Anaconda is a package manager that makes it easier to install, distribute, and manage popular Python libraries and their dependencies. You can use Anaconda for package management with Cloudera Data Science Workbench, but it is not required.

**Note:**
The latest Anaconda package ships with Python 3.7. However, Cloudera Data Science Workbench only supports Python 2.7.11 and Python 3.6.1. For instructions on how to install Anaconda with Python 3.6, refer this section in the Anaconda FAQs: [How do I get Anaconda with Python 3.6?](https://anaconda.org/)

You can install Anaconda before you install Cloudera Data Science Workbench or after. Once Anaconda is installed, perform the following steps to configure Cloudera Data Science Workbench to work with Anaconda:

1. Install the Anaconda package on *all* cluster hosts. For installation instructions, refer to the [Anaconda installation documentation](https://anaconda.org/).
2. Set the `ANACONDA_DIR` property in the Cloudera Data Science Workbench configuration file: `cdsw.conf`. This can be done when you first configure `cdsw.conf` during the installation or later.
3. **Restart Cloudera Data Science Workbench** to have this change go into effect.

**Known Issues and Limitations**

- Cloudera Data Science Workbench cannot be managed by Apache Ambari.
- Apache Phoenix requires additional configuration to run commands successfully from within Cloudera Data Science Workbench engines (sessions, jobs, experiments, models).

**Workaround:** Explicitly set `HBASE_CONF_PATH` to a valid path before running Phoenix commands from engines.

```bash
export HBASE_CONF_PATH=/usr/hdp/hbase/<hdp_version>/0/
```
Installing Cloudera Data Science Workbench 1.5.x on HDP

Use the following steps to install the Cloudera Data Science Workbench RPM package on an HDP cluster.

Prerequisites

Before you begin installing Cloudera Data Science Workbench, perform the following steps to set up a wildcard DNS subdomain for CDSW, disable untrusted SSH access to the hosts, and configure the Application and Docker block devices.

- Set Up a Wildcard DNS Subdomain on page 60
- Disable Untrusted SSH Access on page 60
- Configure the Application and Docker Block Devices

Add Gateway Hosts for Cloudera Data Science Workbench to Your HDP Cluster

To add new hosts to act as Gateway hosts for your cluster:

1. Log in to the Ambari Server.
2. Go to the Hosts page and select Actions > + Add New Hosts.
3. On the Install Options page, enter the fully-qualified domain names for your new hosts.

   The wizard also needs the private key file you created when you set up password-less SSH. Using the host names and key file information, the wizard can locate, access, and interact securely with all the hosts in the cluster. Alternatively, you can manually install and start the Ambari agents on all the new hosts.

   Click Register and Confirm.

   For more detailed instructions, refer to Install Options.

4. The Confirm Hosts page prompts you to confirm that Ambari has located the correct hosts for your cluster and to check those hosts to make sure they have the correct directories, packages, and processes required to continue the install. When you are satisfied with the list of hosts, click Next.

   For detailed instructions, refer to Confirm Hosts.

5. On the Assign Slaves and Clients page, select the Clients that should be installed on the new hosts. To install clients on all hosts, select the Client checkbox for every host. You can use the all option for each available client to expedite this.

   Make sure no other services are running on these hosts. To make this easier, select the none option for all other services.

6. On the Configurations page, select the configuration groups for the new hosts.
7. The Review page displays the host assignments you have made. Check to make sure everything is correct. If you need to make changes, use the left navigation bar to return to the appropriate screen.

   When you are satisfied with your choices, click Deploy.

8. The Install, Start and Test page displays progress as the clients are installed and deployed on each host. When the process is complete, click Next.

9. The Summary page provides you a list of the accomplished tasks. Click Complete and you will be directed back to the Hosts page.

Create HDFS User Directories

To run workloads that leverage HDP cluster services, make sure that HDFS directories (/user/<username>) are created for each user so that they can seamlessly connect to HDP from Cloudera Data Science Workbench.

Perform the following steps for each user directory that must be created.
1. SSH to a host in the cluster that includes the HDFS client.

2. Switch to the `hdfs` system account user:

   ```
   su - hdfs
   ```

3. Create an HDFS directory for the user. For example, you would create the following directory for the default user `admin`:

   ```
   hdfs dfs -mkdir /user/admin
   ```

4. Assign ownership of the new directory to the user. For example, for the new `/user/admin` directory, make the `admin` user the owner of the directory:

   ```
   hdfs dfs -chown admin:hadoop /user/admin
   ```

---

**Install Cloudera Data Science Workbench on the Master Host**

Use the following steps to install Cloudera Data Science Workbench on the master host. *Note that airgapped clusters and non-airgapped clusters use different files for installation.*

1. Download Cloudera Data Science Workbench:

<table>
<thead>
<tr>
<th>Red Hat 7 Repository File</th>
<th>RPM</th>
</tr>
</thead>
</table>
   | For non-airgapped installations, download this file and save it to `/etc/yum.repos.d/:
   | cloudera-cdsw.repo | For airgapped installations, download the Cloudera Data Science Workbench RPM file:
   | Clouderata Data Science Workbench 1.5.0 |

   **Important:** Make sure all Cloudera Data Science Workbench hosts (master and worker) are running the same version of Cloudera Data Science Workbench.

2. Skip this step for airgapped installations. The Cloudera Public GPG repository key verifies that you are downloading genuine packages. Add the repository key:

   ```
   sudo rpm --import
   https://archive.cloudera.com/cdsw1/1.5.0/redhat7/yum/RPM-GPG-KEY-cloudera
   ```

3. **Non-airgapped Installation** - Install the latest RPM with the following command:

   ```
   sudo yum install cloudera-data-science-workbench
   ```

   **Air-gapped Installation** - Copy the RPM to the appropriate gateway host and use the complete filename to install the package. For example:

   ```
   sudo yum install cloudera-data-science-workbench-1.5.0.12345.rpm
   ```

   For guidance on any warnings displayed during the installation process, see *Understanding Installation Warnings* on page 247.

4. Edit the configuration file at `/etc/cdsw/config/cdsw.conf`. The following table lists the configuration properties that can be configured in `cdsw.conf`.
### Table 6: cdsw.conf Properties

<table>
<thead>
<tr>
<th>Properties</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Required Configuration</strong></td>
<td></td>
</tr>
<tr>
<td>DOMAIN</td>
<td>Wildcard DNS domain configured to point to the master host. If the wildcard DNS entries are configured as <code>cdsw.&lt;your_domain&gt;.com</code> and <code>*.cdsw.&lt;your_domain&gt;.com</code>, then <code>DOMAIN</code> should be set to <code>cdsw.&lt;your_domain&gt;.com</code>. Users' browsers should then contact the Cloudera Data Science Workbench web application at <code>http://cdsw.&lt;your_domain&gt;.com</code>. This domain for DNS and is unrelated to Kerberos or LDAP domains.</td>
</tr>
<tr>
<td>MASTER_IP</td>
<td>IPv4 address for the master host that is reachable from the worker hosts. Within an AWS VPC, <code>MASTER_IP</code> should be set to the internal IP address of the master host; for instance, if your hostname is <code>ip-10-251-50-12.ec2.internal</code>, set <code>MASTER_IP</code> to the corresponding IP address, <code>10.251.50.12</code>.</td>
</tr>
<tr>
<td>DISTRO</td>
<td>The Hadoop distribution installed on the cluster. Set this property to HDP.</td>
</tr>
<tr>
<td>DOCKER_BLOCK_DEVICES</td>
<td>Block device(s) for Docker images (space separated if there are multiple). Use the full path to specify the image(s), for instance, <code>/dev/xvde</code>.</td>
</tr>
<tr>
<td>JAVA_HOME</td>
<td>Path where Java is installed on the Cloudera Data Science Workbench hosts. This path must match the <code>JAVA_HOME</code> environment variable that is configured for your HDP cluster. You can find the value in <code>hadoop-env.sh</code> on any node in the HDP cluster. Note that Spark 2.3 requires JDK 1.8. For more details on the specific versions of Oracle JDK recommended for HDP clusters, see the Hortonworks Support Matrix - <a href="https://supportmatrix.hortonworks.com/">https://supportmatrix.hortonworks.com/</a>.</td>
</tr>
<tr>
<td><strong>Optional Configuration</strong></td>
<td></td>
</tr>
<tr>
<td>APPLICATION_BLOCK_DEVICE</td>
<td>(Master Host Only) Configure a block device for application state. If this property is left blank, the filesystem mounted at <code>/var/lib/cdsw</code> on the master host will be used to store all user data. For production deployments, Cloudera strongly recommends you use this option with a dedicated SSD block device for the <code>/var/lib/cdsw</code> mount. (Not recommended) If set, Cloudera Data Science Workbench will format the provided block device as <code>ext4</code>, mount it to <code>/var/lib/cdsw</code>, and store all user data on it. This option has only been provided for proof-of-concept setups, and Cloudera is not responsible for any data loss. Use the full path to specify the mount point, for instance, <code>/dev/xvdf</code>.</td>
</tr>
<tr>
<td>RESERVE_MASTER</td>
<td>Set this property to <code>true</code> to reserve the master host for Cloudera Data Science Workbench’s internal components and services, such as Livelog, the PostgreSQL database, and so on. User workloads will now run exclusively on worker hosts, while the master is reserved for internal application services. <strong>Important:</strong> This feature only applies to deployments with more than one Cloudera Data Science Workbench host. Enabling this feature on single-host deployments will leave Cloudera Data Science Workbench incapable of scheduling any workloads.</td>
</tr>
<tr>
<td>Properties</td>
<td>Description</td>
</tr>
<tr>
<td>---------------</td>
<td>-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------</td>
</tr>
<tr>
<td>DISTRO_DIR</td>
<td>Path where the Hadoop distribution is installed on the Cloudera Data Science Workbench hosts. For HDP clusters, the default location of the packages is /usr/hdp. Specify this property only if you are using a non-default location.</td>
</tr>
<tr>
<td>ANACONDA_DIR</td>
<td>Path where Anaconda is installed. Set this property only if you are using Anaconda for package management.</td>
</tr>
<tr>
<td></td>
<td>By default, the Anaconda package is installed at: /home/&lt;your-username&gt;/anaconda&lt;2 or 3&gt;. Refer to the Anaconda FAQs for more details.</td>
</tr>
<tr>
<td></td>
<td>If you choose to start using Anaconda anytime post-installation, you must set this property and then restart Cloudera Data Science Workbench to have this change take effect.</td>
</tr>
<tr>
<td>TLS_ENABLE</td>
<td>Enable and enforce HTTPS (TLS/SSL) for web access.ian</td>
</tr>
<tr>
<td></td>
<td>You can also set this property to true to enable external TLS termination. For more details on TLS termination, see Enabling TLS/SSL for Cloudera Data Science Workbench on page 231.</td>
</tr>
<tr>
<td>TLS_CERT</td>
<td>Certificate and private key for internal TLS termination.</td>
</tr>
<tr>
<td>TLS_KEY</td>
<td>Setting TLS_CERT and TLS_KEY will enable internal TLS termination. You must also set TLS_ENABLE to true above to enable and enforce internal termination. Set these only if you are not terminating TLS externally.</td>
</tr>
<tr>
<td></td>
<td>Make sure you specify the full path to the certificate and key files, which must be in PEM format.</td>
</tr>
<tr>
<td></td>
<td>For details on certificate requirements and enabling TLS termination, see Enabling TLS/SSL for Cloudera Data Science Workbench on page 231.</td>
</tr>
<tr>
<td>HTTP_PROXY</td>
<td>If your deployment is behind an HTTP or HTTPS proxy, set the respective HTTP_PROXY or HTTPS_PROXY property in /etc/cdsw/config/cdsw.conf to the hostname of the proxy you are using.</td>
</tr>
<tr>
<td>HTTPS_PROXY</td>
<td>If you are using an intermediate proxy, such as Cntlm, to handle NTLM authentication, add the Cntlm proxy address to the HTTP_PROXY or HTTPS_PROXY fields in cdsw.conf.</td>
</tr>
<tr>
<td></td>
<td>If the proxy server uses TLS encryption to handle connection requests, you will need to add the proxy’s root CA certificate to your host’s store of trusted certificates. This is because proxy servers typically sign their server certificate with their own root certificate. Therefore, any connection attempts will fail until the Cloudera Data Science Workbench host trusts the proxy’s root CA certificate. If you do not have access to your proxy’s root certificate, contact your Network / IT administrator.</td>
</tr>
</tbody>
</table>
### Properties

<table>
<thead>
<tr>
<th>Property</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>To enable trust, copy the proxy’s root certificate to the trusted CA certificate store (ca-trust) on the Cloudera Data Science Workbench host.</td>
</tr>
<tr>
<td></td>
<td><strong>cp /tmp/&lt;proxy-root-certificate&gt;.crt /etc/pki/ca-trust/source/anchors/</strong></td>
</tr>
<tr>
<td></td>
<td>Use the following command to rebuild the trusted certificate store.</td>
</tr>
<tr>
<td></td>
<td><strong>update-ca-trust extract</strong></td>
</tr>
<tr>
<td>ALL_PROXY</td>
<td>If a SOCKS proxy is in use, set to socks5://&lt;host&gt;:&lt;port&gt;/.</td>
</tr>
<tr>
<td>NO_PROXY</td>
<td>Comma-separated list of hostnames that should be skipped from the proxy.</td>
</tr>
<tr>
<td></td>
<td>Starting with version 1.4, if you have defined a proxy in the HTTP_PROXY(S) or ALL_PROXY properties, Cloudera Data Science Workbench automatically appends the following list of IP addresses to the NO_PROXY configuration. Note that this is the minimum required configuration for this field.</td>
</tr>
<tr>
<td></td>
<td>This list includes 127.0.0.1, localhost, and any private Docker registries and HTTP services inside the firewall that Cloudera Data Science Workbench users might want to access from the engines.</td>
</tr>
<tr>
<td></td>
<td>&quot;127.0.0.1,localhost,100.66.0.1,100.66.0.2,100.66.0.3,100.66.0.4,100.66.0.5,100.66.0.6,100.66.0.7,100.66.0.8,100.66.0.9,100.66.0.10,100.66.0.11,100.66.0.12,100.66.0.13,100.66.0.14,100.66.0.15,100.66.0.16,100.66.0.17,100.66.0.18,100.66.0.19,100.66.0.20,100.66.0.21,100.66.0.22,100.66.0.23,100.66.0.24,100.66.0.25,100.66.0.26,100.66.0.27,100.66.0.28,100.66.0.29,100.66.0.30,100.66.0.31,100.66.0.32,100.66.0.33,100.66.0.34,100.66.0.35,100.66.0.36,100.66.0.37,100.66.0.38,100.66.0.39,100.66.0.40,100.66.0.41,100.66.0.42,100.66.0.43,100.66.0.44,100.66.0.45,100.66.0.46,100.66.0.47,100.66.0.48,100.66.0.49,100.66.0.50,100.77.0.10,100.77.0.128,100.77.0.129,100.77.0.130,100.77.0.131,100.77.0.132,100.77.0.133,100.77.0.134,100.77.0.135,100.77.0.136,100.77.0.137,100.77.0.138,100.77.0.139&quot;</td>
</tr>
<tr>
<td>NVIDIA_GPU_ENABLE</td>
<td>Set this property to true to enable GPU support for Cloudera Data Science Workbench workloads. When this property is enabled on a host is equipped with GPU hardware, the GPU(s) will be available for use by Cloudera Data Science Workbench hosts.</td>
</tr>
<tr>
<td></td>
<td>If this property is set to true on a host that does not have GPU support, there will be no effect. By default, this property is set to false.</td>
</tr>
<tr>
<td></td>
<td>For detailed instructions on how to enable GPU-based workloads on Cloudera Data Science Workbench, see Using NVIDIA GPUs for Cloudera Data Science Workbench Projects on page 117.</td>
</tr>
<tr>
<td>NVIDIA_LIBRARY_PATH</td>
<td>Complete path to the NVIDIA driver libraries.</td>
</tr>
</tbody>
</table>

5. Initialize and start Cloudera Data Science Workbench:

```
cdsw init
```

The application will take a few minutes to bootstrap. You can watch the status of application installation and startup with `watch cdsw status`. 
(Optional) Install Cloudera Data Science Workbench on Worker Hosts

**Note:** Worker hosts are not required for a fully-functional Cloudera Data Science Workbench deployment. For proof-of-concept deployments, you can deploy a 1-host cluster with just a Master host. The Master host can run user workloads just as a worker host can.

Cloudera Data Science Workbench supports adding and removing additional worker hosts at any time. Worker hosts allow you to transparently scale the number of concurrent workloads users can run.

Use the following steps to add worker hosts to Cloudera Data Science Workbench. **Note that airgapped clusters and non-airgapped clusters use different files for installation.**

1. **Download Cloudera Data Science Workbench:**

<table>
<thead>
<tr>
<th>Red Hat Repository File</th>
<th>RPM</th>
</tr>
</thead>
<tbody>
<tr>
<td>For non-airgapped installations, download this file and save it to /etc/yum.repos.d/:</td>
<td>For airgapped installations, download the Cloudera Data Science Workbench RPM file:</td>
</tr>
<tr>
<td>cloudera-cdsw.repo</td>
<td>Cloudera Data Science Workbench 1.5.0</td>
</tr>
</tbody>
</table>

**Important:** Make sure all Cloudera Data Science Workbench hosts (master and worker) are running the same version of Cloudera Data Science Workbench.

2. **Skip this step for airgapped installations. The Cloudera Public GPG repository key verifies that you are downloading genuine packages. Add the repository key:**

```bash
sudo rpm --import https://archive.cloudera.com/cdsw1/1.5.0/redhat7/yum/RPM-GPG-KEY-cloudera
```

3. **Non-airgapped Installation** - Install the latest RPM with the following command:

```bash
sudo yum install cloudera-data-science-workbench
```

**Air-gapped Installation** - Copy the RPM to the appropriate gateway host and use the complete filename to install the package. For example:

```bash
sudo yum install cloudera-data-science-workbench-1.5.0.12345.rpm
```

For guidance on any warnings displayed during the installation process, see [Understanding Installation Warnings](#) on page 247.

4. **Copy cdsw.conf file from the master host:**

```bash
scp root@<cdsw-master-hostname.your_domain.com>:/etc/cdsw/config/cdsw.conf /etc/cdsw/config/cdsw.conf
```

After initialization, the cdsw.conf file includes a generated bootstrap token that allows worker hosts to securely join the cluster. You can get this token by copying the configuration file from master and ensuring it has 600 permissions.

If your hosts have heterogeneous block device configurations, modify the Docker block device settings in the worker host configuration file after you copy it. Worker hosts do not need application block devices, which store the project files and database state, and this configuration option is ignored.

5. **Create /var/lib/cdsw on the worker host. This directory must exist on all worker hosts. Without it, the next step that registers the worker host with the master will fail.**
Unlike the master host, the /var/lib/cdsw directory on worker hosts does **not** need to be mounted to an Application Block Device. It is only used to store client configuration for HDP services on workers.

6. On the **worker** host, run the following command to add the host to the cluster:

   ```
cdsw join
   ```

   This causes the worker hosts to register themselves with the Cloudera Data Science Workbench master host and increase the available pool of resources for workloads.

7. Return to the master host and verify the host is registered with this command:

   ```
   cdsw status
   ```

Create the Site Administrator Account

Installation typically takes 30 minutes although it might take an additional 60 minutes for the R, Python, and Scala engine to be available on all hosts.

After installation is complete, go to the Cloudera Data Science Workbench web application at http://cdsw.<your_domain>.com.

**Note:** You must access Cloudera Data Science Workbench from the **DOMAIN** previously configured in cdsw.conf, and not the hostname of the master host. Visiting the hostname of the master host will result in a 404 error.

Sign up for a new account. The first account that you create becomes the site administrator. As a site administrator, you can invite new users, secure the deployment, and **upload a license key** for the product. For more details on these tasks, see the **Administration** and **Security** guides.

Getting Started with a New Project on Cloudera Data Science Workbench

To sign up, open the Cloudera Data Science Workbench web application in a browser. The application is typically hosted on the master host at http://cdsw.<your_domain>.com. The first time you log in, you will be prompted to create a username and password. Note that if your site administrator has configured your deployment to require invitations, you will need an invitation link to sign up.

You can use this account to create a new project and start using the workbench to run data science workloads. Watch the following video for a quick demo (demo starts at 00:30): **CDSW Quickstart Demo**

Related Documentation:

The Cloudera Data Science Workbench user guide includes detailed instructions that should help you get started with running workloads on Cloudera Data Science Workbench.

- **Creating and Managing Projects**
- **Collaborating on Projects**
- **Accessing Data from HDFS, Hive, HBase, etc.**
- **Model Training and Deployment:** **Experiments**, **Models**

Upgrading a CDSW 1.5.x Deployment from HDP 2 to HDP 3

Cloudera Data Science Workbench ships a single RPM that can be used to install CDSW on both, HDP 2, and HDP 3 clusters. The upgrade path for RPM deployments is:

1. **Upgrade from HDP 2.6.5 (or higher) to HDP 3.1.x (or higher).**
2. Restart Cloudera Data Science Workbench. Check Ambari to ensure that there are no more stale configs on the cluster hosts.

Frequently Asked Questions (FAQs)

Does CDSW-on-HDP require a license key?

Cloudera Data Science Workbench is fully functional during a 60-day, non-renewable trial period. The trial period starts when you create your first user. When the 60-day period ends, functionality will be limited. You will not be able to create any new projects or schedule any more workloads.

At this point, you must obtain a Cloudera Enterprise license and upload it to Cloudera Data Science Workbench. Cloudera Data Science Workbench will then go back to being fully functional.

For details, see Managing License Keys for Cloudera Data Science Workbench on page 214.

How do I file a support case for CDSW-on-HDP?

If you have encountered an issue, you can create a support ticket in the Cloudera Support portal.

Before you log a support ticket, run the following command on the master host to create a tarball with diagnostic information for your Cloudera Data Science Workbench installation. Attach the resulting bundle to the support case you create.

```
cdsw logs
```

You can also use SmartSense to collect diagnostic data and cluster metrics from Ambari.
Getting Started with Cloudera Data Science Workbench

This topic provides a suggested method for quickly getting started with data science projects on Cloudera Data Science Workbench.

Watch the following video for a quick demo of the steps described in this topic: [CDSW Quickstart Demo](#)

### Sign up

To sign up, open the Cloudera Data Science Workbench web application in a browser. The application is typically hosted on the master host at `http://cdsw.<your_domain>.com`. The first time you log in, you will be prompted to create a username and password. Note that the first account created will receive site administrator privileges.

If your site administrator has configured your deployment to require invitations, you will need an invitation link to sign up.

### Create a Project from a Built-in Template

Cloudera Data Science Workbench is organized around projects. Projects hold all the code, configuration, and libraries needed to reproducibly run analyses.

To help you get started, Cloudera Data Science Workbench includes sample template projects in R, Python, PySpark, and Scala. Using a template project gives you the impetus to start using the Cloudera Data Science Workbench right away.

**Create a Template Project**

![Create a New Project]

To create a template project:
1. Sign in to Cloudera Data Science Workbench.
2. Click **New Project**.
3. Enter the account and project name.
4. Under the **Template** tab, you can choose one of the programming languages to create a project from one of the built-in templates. Alternatively, if your site administrator has added any custom template projects, those will also be available in this dropdown list.
5. Click **Create Project**.

After creating your project, you see your project files and the list of jobs defined in your project. These project files are stored on an internal NFS server, and are available to all your project sessions and jobs, regardless of the gateway hosts they run on. Any changes you make to the code or libraries you install into your project will be immediately available when running an engine.

**Launch a Session to Run the Project**

Cloudera Data Science Workbench provides an interactive environment tailored for data science called the **workbench**. It supports R, Python, and Scala engines, one of which we will use to run the template project.

**Workbench**

Perform the following steps to run the project:

**Open the Workbench to Launch a Session**

To run the project code, open the workbench and launch a new session.

1. Navigate to the new project’s **Overview** page.
2. Click **Open Workbench**.
3. **Launch a New Session**
1. Use **Select Engine Kernel** to choose the programming language that your project uses.
2. Use **Select Engine Profile** to select the number of CPU cores and memory to be used.
3. Click **Launch Session**.

The command prompt at the bottom right of your browser window will turn green when the engine is ready. Sessions typically take between 10 and 20 seconds to start.

**Execute Project Code**

You can enter and execute code using either the editor or the command prompt. The editor is best used for code you want to keep, while the command prompt is best for quick interactive exploration.

**Editor** - To run code in the editor:

1. Select a script from the project files on the left sidebar.
2. To run the whole script click on the top navigation bar, or, highlight the code you want to run and press Ctrl+Enter (Windows/Linux) or cmd+Enter (macOS).

**Command Prompt** - The command prompt functions largely like any other. Enter a command and press Enter to execute it. If you want to enter more than one line of code, use Shift+Enter to move to the next line. The output of your code, including plots, appears in the console.

**Code Autocomplete** - The Python and R kernels include support for automatic code completion, both in the editor and the command prompt. Use single tab to display suggestions and double tab for autocomplete.
Test Terminal Access

Cloudera Data Science Workbench provides terminal access to the running engines from the web console. You can use the terminal to move files around, run Git commands, and understand what resources are already available to you in the project environment.

To access the Terminal from a running session, click Terminal Access above the console pane. The terminal's default working directory is /home/cdsw, which is a temporary directory where all your project files are stored for this session.

Note that the terminal does not provide root or sudo access to the container.

Stop the Session

When you are done with the session, click Stop in the menu bar above the console

Next Steps

Now that you have successfully run a sample workload with the Cloudera Data Science Workbench, further acquaint yourself with Cloudera Data Science Workbench by reading the User, Administration, and Security guides to learn more about the types of users, how to collaborate on projects, how to use Spark 2 for advanced analytics, and how to secure your deployment.

- Managing Cloudera Data Science Workbench Users on page 102
- Cloudera Data Science Workbench Administration Guide on page 207
- Cloudera Data Science Workbench Security Guide on page 224
- Using CDS 2.x Powered by Apache Spark on page 197
Managing Cloudera Data Science Workbench Users

User Contexts

The Cloudera Data Science Workbench UI uses the notion of contexts to separate your personal account from any team accounts you belong to. Depending on the context you are in, you will be able to modify settings for either your personal account, or a team account, and see the projects created in each account. Shared personal projects will show up in your personal account context.

Context changes in the UI are subtle, so if you’re wondering where a project or setting lives, first make sure you are in the right context. The application header will tell you which context you are currently in. You can switch to a different context by going to the drop-down menu in the upper right-hand corner of the page.

Personal

Your personal context can be used to work on private projects, experiments, and so on. For related management tasks, see Managing your Personal Account on page 102.

Team

Teams allow streamlined administration of projects. If you are going to collaborate with teammates on more than one project, you should create a team account and work on shared projects under the team context. For related tasks, see Managing Team Accounts on page 103.

User Roles

Currently, Cloudera Data Science Workbench has only one deployment-wide specialised user role: site administrator. All other roles are created within the scope of a particular team or project and do not affect the rest of the deployment.

Site Administrators

Site administrators are superusers who have complete access to all activity on the deployment. By default, the first user account that signs up for Cloudera Data Science Workbench becomes a site administrator. They can add/invite more users, disable existing users, assign site administrator privileges to others, and so on. For instructions, see Managing Users as a Site Administrator on page 104.

Roles within Teams and Projects

Within the context of each team and project, users can be assigned Viewer, Contributor, or Admin roles. These roles are only applicable within the limited scope of the team/project. More details on the privileges associated with each of these roles are included in the Creating Teams and Project Collaborators topics respectively.

Managing your Personal Account

To manage your personal account settings:

1. Sign in to Cloudera Data Science Workbench.
2. From the upper right drop-down menu, switch context to your personal account.
3. Click Settings.
**Profile**
You can modify your name, email, and bio on this page.

**Teams**
This page lists the teams you are a part of and the role assigned to you for each team.

**SSH Keys**
Your public SSH key resides here. SSH keys provide a useful way to access to external resources such as databases or remote Git repositories. For instructions, see SSH Keys on page 245.

**Hadoop Authentication**
Enter your Hadoop credentials here to authenticate yourself against the cluster KDC. For more information, see Hadoop Authentication with Kerberos for Cloudera Data Science Workbench on page 236.

---

**Important:** You can also access your personal account settings by clicking Account settings in the upper right-hand corner drop-down menu. This option will always take you to your personal settings page, irrespective of the context you are currently in.

---

**Managing Team Accounts**

Users who work together on more than one project and want to facilitate collaboration can create a Team. Teams allow streamlined administration of projects. Team projects are owned by the team, rather than an individual user. Team administrators can add or remove members at any time, assigning each member different permissions.

---

### Creating a Team

To create a team:

1. Click the plus sign (+) in the title bar, to the right of the Search field.
2. Select Create Team.
3. Enter a Team Name.
4. Click Create Team.
5. Add or invite team members. Team members can have one of the following privilege levels:
   - **Viewer** - Read-only access to team projects. Cannot create new projects within the team but can be added to existing ones.
Contributor - Write-level access to all team projects with Team or Public visibility. Can create new projects within the team. They can also be added to existing team projects.

Admin - Has complete access to all team projects, can add new team members, and modify team account information.

6. Click Done.

Note: Team administrators and team members with Admin or Contributor privileges on projects have access to all your sessions and can execute commands within your active sessions. Cloudera Data Science Workbench 1.4.3 introduces a new feature that allows site administrators to restrict this ability by allowing only session creators to execute commands within their own active sessions. For details, see Restricting Access to Active Sessions.

Modifying Team Account Settings

Team administrators can modify account information, add or invite new team members, and view/edit privileges of existing members. To make these changes:

1. From the upper right drop-down menu, switch context to the team account.
2. Click Settings to open up the Account Settings dashboard.
   - Profile
     Modify the team description on this page.
   - Members
     You can add new team members on this page, and modify privilege levels for existing members.
   - SSH Keys
     The team’s public SSH key resides here. Team SSH keys provide a useful way to give an entire team access to external resources such as databases. For instructions, see SSH Keys on page 245. Generally, team SSH keys should not be used to authenticate against Git repositories. Use your personal key instead.

Managing Users as a Site Administrator

Required Role: Site Administrator

This topic describes how to manage Cloudera Data Science Workbench users as a site administrator. By default, the first user account that signs up for the Cloudera Data Science Workbench becomes a site administrator. Site administrators can manage other users, monitor resources, secure access to the deployment, and upload license keys for the product.

Important: Site administrators have complete access to all activity on the deployment. This includes access to all teams and projects on the deployment, even if they have not been explicitly added as team members or collaborators.

To access the site administrator dashboard:

1. Go to the Cloudera Data Science Workbench web application (http://cdsw.company.com) and log in as a site administrator.
2. On the left sidebar, click Admin. You will see an array of tabs for all the tasks you can perform as a site administrator.
As a site administrator you can add new users, assign or modify privileges for existing users, and monitor user activity on the Cloudera Data Science Workbench deployment.

Adding New Users

To invite new users, navigate to the Admin > Users tab. Under Invitations, enter the name or email ID of the person you want to invite and click Invite. This tab will show you a list of all outstanding invitations. Once an invitation has been accepted, the record will no longer show up on this page. The Users tab also displays a list of users of the application. Click on a username to see more details about the user.

If you want new users to join by invitation only, go to the Admin > Settings tab and check the Require invitation to sign up checkbox to require invitation tokens for account creation. By default, invitations are sent from noreply@your-cdsw-domain. To modify this default, see Cloudera Data Science Workbench Email Notifications on page 214.

Assigning the Site Administrator role to an Existing User

To make a regular user a site administrator:

1. Sign in to Cloudera Data Science Workbench as a site administrator.
2. Click Admin.
3. Click the Users tab.
4. Click on the username of the user who you want to make a site administrator.
5. Select the Site Administrator checkbox.
Managing Cloudera Data Science Workbench Users

Disabling User Accounts

Use the following instructions to disable user accounts. Note that disabled users cannot login and do not count towards named users for licensing.

1. Sign in to Cloudera Data Science Workbench as a site administrator.
2. Click Admin.
3. Click the Users tab.
4. Click on the username of the user who you want to disable.
5. Select the Disabled checkbox.
6. Click Update.

Monitoring Users

The Users tab on the admin dashboard displays the complete list of users. You can see which users are currently active, and when a user last logged in to the Cloudera Data Science Workbench. To modify a user’s username, email or permissions, click the Edit button under the Action column.
Managing Projects in Cloudera Data Science Workbench

Projects form the heart of Cloudera Data Science Workbench. They hold all the code, configuration, and libraries needed to reproducibly run analyses. Each project is independent, ensuring users can work freely without interfering with one another or breaking existing workloads.

This topic describes how to create and manage projects in Cloudera Data Science Workbench.

Creating a Project

To create a Cloudera Data Science Workbench project:

1. Go to Cloudera Data Science Workbench and on the left sidebar, click Projects.
2. Click New Project.
3. If you are a member of a team, from the drop-down menu, select the Account under which you want to create this project. If there is only one account on the deployment, you will not see this option.
4. Enter a Project Name.
5. Select Project Visibility from one of the following options.
   - **Private** - Only project collaborators can view or edit the project.
   - **Team** - If the project is created under a team account, all members of the team can view the project. Only explicitly-added collaborators can edit the project.
   - **Public** - All authenticated users of Cloudera Data Science Workbench will be able to view the project. Collaborators will be able to edit the project.
6. Under Initial Setup, you can either create a blank project, or select one of the following sources for your project files.
   - **Built-in Templates** - Template projects contain example code that can help you get started with the Cloudera Data Science Workbench. They are available in R, Python, PySpark, and Scala. Using a template project is not required, but it helps you start using the Cloudera Data Science Workbench right away.
   - **Custom Templates** - Starting with version 1.3, site administrators can add template projects that are customized for their organization’s use-cases. For details, see Custom Template Projects on page 110.
   - **Local** - If you have an existing project on your local disk, use this option to upload compressed files or folders to Cloudera Data Science Workbench.
   - **Git** - If you already use Git for version control and collaboration, you can continue to do so with the Cloudera Data Science Workbench. Specifying a Git URL will clone the project into Cloudera Data Science Workbench. If you use a Git SSH URL, your personal private SSH key will be used to clone the repository. This is the recommended approach. However, you must add the public SSH key from your personal Cloudera Data Science Workbench account to the remote Git hosting service before you can clone the project.
7. Click Create Project. After the project is created, you can see your project files and the list of jobs defined in your project.

   Note that as part of the project filesystem, Cloudera Data Science Workbench also creates the following .gitignore file.

```plaintext
R
node_modules
*.pyc
.*
!.gitignore
```

8. **(Optional)** To work with team members on a project, use the instructions in the following section to add them as collaborators to the project.
Adding Collaborators

If you want to work closely with colleagues on a particular project, use the following steps to add them to the project.

1. Navigate to the project overview page.
2. Click Team to open the Collaborators page.
3. Search for collaborators by either name or email address and click Add.

For a project created under your personal account, anyone who belongs to your organization can be added as a collaborator. For a project created under a team account, you can only add collaborators that already belong to the team. If you want to work on a project that requires collaborators from different teams, create a new team with the required members, and then create a project under that account. If your project was created from a Git repository, each collaborator must create the project from the same central Git repository.

You can grant project collaborators one of three levels of access:

- **Viewer** - Read-only access to code, data, and results.
- **Contributor** - Can view, edit, create, and delete files and environmental variables, run sessions/experiments/jobs/models and execute code in running jobs. Additionally, Contributors can set the default engine for the project.
- **Admin** - Has complete access to all aspects of the project. This includes the ability to add new collaborators, and delete the entire project.

**Warning:**

**Collaborating Securely on Projects**

Before adding project collaborators, you must remember that assigning the Contributor or Admin role to a project collaborator is the same as giving them write access to your data in CDH. This is because project contributors and project administrators have write access to all your project code (including any library code that you might not be actively inspecting). For example, a contributor/admin could modify project file(s) to insert code that deletes some data on the CDH cluster. The next time you launch a session and run the same code, it will appear as though you deleted the data yourself.

Additionally, project collaborators also have access to all actively running sessions and jobs. This means that a malicious user can easily impersonate you by accessing one of your active sessions. Therefore, it is extremely important to restrict project access to trusted collaborators only. Note that Cloudera Data Science Workbench 1.4.3 introduces a new feature that allows site administrators to restrict this ability by allowing only session creators to execute commands within their own active sessions. For details, see Restricting Access to Active Sessions.

For these reasons, Cloudera recommends using Git to collaborate securely on shared projects. This will also help avoid file modification conflicts when your team is working on more elaborate projects.

For more information on collaborating effectively, see Collaborating on Projects with Cloudera Data Science Workbench on page 128.

Modifying Project Settings

Project contributors and administrators can modify aspects of the project environment such as the engine being used to launch sessions, the environment variables, and create SSH tunnels to access external resources. To make these changes:

1. Switch context to the account where the project was created.
2. Click Projects.
3. From the list of projects, select the one you want to modify.
4. Click **Settings** to open up the Project Settings dashboard.

**Options**
Modify the project name and its privacy settings on this page.

**Engine**
Cloudera Data Science Workbench ensures that your code is always run with the specific engine version you selected. You can select the version here. For advanced use cases, Cloudera Data Science Workbench projects can use custom Docker images for their projects. Site administrators can whitelist images for use in projects, and project administrators can use this page to select which of these whitelisted images is installed for their projects. For an example, see [Customized Engine Images](#) on page 184.

**Environment** - If there are any environmental variables that should be injected into all the engines running this project, you can add them to this page. For more details, see [Engine Environment Variables](#) on page 179.

**Tunnels**
In some environments, external databases and data sources reside behind restrictive firewalls. Cloudera Data Science Workbench provides a convenient way to connect to such resources using your SSH key. For instructions, see [SSH Tunnels](#) on page 246.

**Delete Project**
This page can only be accessed by project administrators. Remember that deleting a project is irreversible. All files, data, sessions, and jobs will be lost.

**Managing Files**

⚠️ **Important:** For use cases beyond simple projects, Cloudera strongly recommends using [Git](#) to manage your projects using version control.

Cloudera Data Science Workbench allows you to move, rename, copy, and delete files within the scope of the project where they live. You can also upload new files to a project, or download project files. Files can only be uploaded within the scope of a single project. Therefore, to access a script or data file from multiple projects, you will need to manually upload it to all the relevant projects.

1. Switch context to the account where the project was created.
2. Click **Projects**.
3. From the list of projects, click on the project you want to modify. This will take you to the project overview.
4. Click **Files**.

**Upload Files to a Project**

Click **Upload**. Select **Files** or **Folder** from the dropdown, and choose the files or folder you want to upload from your local filesystem.

In addition to uploading files or a folder, you can upload a `.tar` file of multiple files and folders. After you select and upload the `.tar` file, you can use a terminal session to extract the contents:

1. On the project overview page, click **Open Workbench** and select a running session or create a new one.
2. Click **Terminal access**.
3. In the terminal window, extract the contents of the `.tar` file:

   ```bash
   tar -xvf <file_name>.tar.gz
   ```

   The extracted files are now available for the project.
Download Project Files

Click Download to download the entire project in a .zip file. To download only a specific file, select the checkbox next to the file(s) to be download and click Download.

You can also use the checkboxes to Move, Rename, or Delete files within the scope of this project.

Disabling Project File Uploads and Downloads

Required Role: Site Administrator

By default, all Cloudera Data Science Workbench users are allowed to upload and download files to/from a project. Version 1.5 introduces a new feature flag that allows site administrators to hide the UI features that let users upload and download project files.

Note that this feature flag only removes the relevant features from the Cloudera Data Science Workbench UI. It does not disable the ability to upload and download files through the backend web API.

To disable project file uploads and downloads:

1. Go to Admin > Security.
2. Under the File Upload/Download section, disable the Allow file upload/download through UI checkbox.

Custom Template Projects

Required Role: Site Administrator

Site administrators can add template projects that have been customized for their organization's use-cases. These custom project templates can be added in the form of a Git repository.

To add a new template project, go to Admin > Settings. Under the Project Templates section, provide a template name, the URL to the project's Git repository, and click Add.

The added templates will become available in the Template tab on the Create Project page. Site administrators can add, edit, or delete custom templates, but not the built-in ones. However, individual built-in templates can be disabled using a checkbox in the Project Templates table at Admin > Settings.

Deleting a Project

Important: Deleting a project is an irreversible action. All files, data, and history related to the project will be lost. This includes any jobs, sessions or models you created within the project.

To delete a project:

1. Go to the project Overview page.
2. On the left sidebar, click Settings.
3. Go to the Delete Project.
4. Click Delete Project and click OK to confirm.

Using the Workbench

The workbench console provides an interactive environment tailored for data science, supporting R, Python and Scala. It currently supports R, Python, and Scala engines. You can use these engines in isolation, as you would on your laptop, or connect to your CDH cluster using Cloudera Distribution of Apache Spark 2 and other libraries.

The workbench UI includes four primary components:
• An editor where you can edit your scripts.
• A console where you can track the results of your analysis.
• A command prompt where you can enter commands interactively.
• A terminal where you can use a Bash shell.

Typically, you would use the following steps to run a project in the workbench:

Launch a Session

1. Navigate to your project's Overview page.
2. Click Open Workbench.
3. Launch a New Session
1. Use **Select Engine Kernel** to choose the programming language that your project uses.
2. Use **Select Engine Profile** to select the number of **CPU cores** and memory to be used.
3. Click **Launch Session**.

The command prompt at the bottom right of your browser window will turn green when the engine is ready. Sessions typically take between 10 and 20 seconds to start.

### Execute Code

You can enter and execute code at the command prompt or the editor. The editor is best for code you want to keep, while the command prompt is best for quick interactive exploration.

**Command Prompt** - The command prompt functions largely like any other. Enter a command and press **Enter** to execute it. If you want to enter more than one line of code, use **Shift+Enter** to move to the next line. The output of your code, including plots, appears in the console.

If you created your project from a template, you should see project files in the editor. You can open a file in the editor by clicking the file name in the file navigation bar on the left.

**Editor** - To run code from the editor:

1. Select a script from the project files on the left sidebar.
2. To run the whole script click ➤ on the top navigation bar, or, highlight the code you want to run and press **Ctrl+Enter** (Windows/Linux) or **cmd+Enter** (macOS).

When doing real analysis, writing and executing your code from the editor rather than the command prompt makes it easy to iteratively develop your code and save it along the way.
If you require more space for your editor, you can collapse the file list by double-clicking between the file list pane and the editor pane. You can hide the editor using editor’s View menu.

**Code Autocomplete**

The Python and R kernels include support for automatic code completion, both in the editor and the command prompt. Use single tab to display suggestions and double tab for autocomplete.

**Project Code Files**

All project files are stored to persistent storage within the respective project directory at /var/lib/cdsw/current/projects. They can be accessed within the project just as you would in a typical directory structure. For example, you can import functions from one file to another within the same project.

**Access the Terminal**

Cloudera Data Science Workbench provides full terminal access to running engines from the web console. You can use the terminal to move files around, run Git commands, access the YARN and Hadoop CLIs, or install libraries that cannot be installed directly from the engine. To access the Terminal from a running session, click Terminal Access above the session log pane.

The terminal’s default working directory is /home/cdsw, which is where all your project files are stored. Any modifications you make to this folder will persist across runs, while modifications to other folders are discarded.

If you are using Kerberos authentication, you can run klist to see your Kerberos principal. If you run hdfs dfs -ls you will see the files stored in your HDFS home directory.

Note that the terminal does not provide root or sudo access to the container. To install packages that require root access, see Customizing Engine Images.

**Stop a Session**

When you are done with the session, click Stop in the menu bar above the console, or use code to exit by typing the following command:

R
```
R
> quit()
```

Python
```
Python
> exit
```

Scala
```
Scala
> quit()
```

Sessions automatically stop after an hour of inactivity.
Data Visualization

Each language on Cloudera Data Science Workbench has a visualization system that you can use to create plots, including rich HTML visualizations.

Simple Plots

To create a simple plot, run a console in your favorite language and paste in the following code sample:

R

```r
# A standard R plot
plot(rnorm(1000))

# A ggplot2 plot
library("ggplot2")
qplot(hp, mpg, data=mtcars, color=am,
     facets=gear~cyl, size=I(3),
     xlab="Horsepower", ylab="Miles per Gallon")
```

Python 2

```python
import matplotlib.pyplot as plt
import random
plt.plot([random.normalvariate(0,1) for i in xrange(1,1000)])
```

Cloudera Data Science Workbench processes each line of code individually (unlike notebooks that process code per-cell). This means if your plot requires multiple commands, you will see incomplete plots in the workbench as each line is processed.

To get around this behavior, wrap all your plotting commands in one Python function. Cloudera Data Science Workbench will then process the function as a whole, and not as individual lines. You should then see your plots as expected.

Saved Images

You can also display images, using a command in the following format:

R

```r
library("cdsw")
download.file("https://upload.wikimedia.org/wikipedia/commons/2/29/Minard.png", 
"/cdn/Minard.png")
image("Minard.png")
```

Python 2

```python
import urllib
from IPython.display import Image
urllib.urlretrieve("http://upload.wikimedia.org/wikipedia/commons/2/29/Minard.png", 
"Minard.png")
Image(filename="Minard.png")
```

HTML Visualizations

Your code can generate and display HTML. To create an HTML widget, paste in the following:

R

```r
library("cdsw")
html('<svg><circle cx="50" cy="50" r="50" fill="red" /></svg>')
```
Python 2

```python
from IPython.display import HTML
HTML('<svg><circle cx="50" cy="50" r="50" fill="red" /></svg>
```

Scala

Cloudera Data Science Workbench allows you to build visualization libraries for Scala using `jvm-repr`. The following example demonstrates how to register a custom HTML representation with the "text/html" mimetype in Cloudera Data Science Workbench. This output will render as HTML in your workbench session.

```scala
//HTML representation
case class HTML(html: String)
//Register a displayer to render html
Displayers.register(classOf[HTML],
  new Displayer[HTML] {
    override def display(html: HTML): java.util.Map[String, String] = {
      Map("text/html" -> html.html
    ).asJava
  })
val helloHTML = HTML("<h1> <em>Hello World</em> </h1>")
display(helloHTML)
```

IFrame Visualizations

Note:

Cloudera Data Science Workbench versions 1.4.2 (and higher) added a new feature that allowed users to enable HTTP security headers for responses to Cloudera Data Science Workbench. This setting is enabled by default. However, the X-Frame-Options header added as part of this feature blocks rendering of iFrames injected by third-party data visualization libraries.

To work around this issue, a site administrator can go to the **Admin > Security** page and disable the **Enable HTTP security headers** property. Restart Cloudera Data Science Workbench for this change to take effect.

Most visualizations require more than basic HTML. Embedding HTML directly in your console also risks conflicts between different parts of your code. The most flexible way to embed a web resource is using an IFrame:

R

```r
library("cdsw")
iframe(src="https://www.youtube.com/embed/8pHzROP1D-w", width="854px", height="510px")
```

Python 2

```python
from IPython.display import HTML
HTML('<iframe width="854" height="510"
src="https://www.youtube.com/embed/8pHzROP1D-w"></iframe>
```

You can generate HTML files within your console and display them in iFrames using the `/cdn` folder. The `cdn` folder persists and services static assets generated by your engine runs. For instance, you can embed a full HTML file with iFrames.

Managing Projects in Cloudera Data Science Workbench
R

```r
library("cdsw")
f <- file("/cdn/index.html")
html.content <- paste("<p>Here is a normal random variate: ", rnorm(1), "</p>")
writeLines(c(html.content), f)
close(f)
iframe("index.html")
```

Python 2

```python
from IPython.display import HTML
import random

html_content = "<p>Here is a normal random variate: %f </p>" % random.normalvariate(0,1)
file("/cdn/index.html", "w").write(html_content)
HTML("<iframe src=index.html>")
```

Cloudera Data Science Workbench uses this feature to support many rich plotting libraries such as htmlwidgets, Bokeh, and Plotly.

### Grid Displays

Cloudera Data Science Workbench supports native grid displays of DataFrames across several languages.

**Python 3**

Using DataFrames with the pandas package requires per-session activation:

```python
import pandas as pd
pd.options.display.html.table_schema = True
pd.DataFrame(data=range(1,100))
```

For PySpark DataFrames, use pandas and run `df.toPandas()` on a PySpark DataFrame. This will bring the DataFrame into local memory as a pandas DataFrame.

**Note:**

A Python project originally created with engine 1 will be running pandas version 0.19, and will not auto-upgrade to version 0.20 by simply selecting engine 2 in the project's Settings > Engine page.

The pandas data grid setting only exists starting in version 0.20.1. To upgrade, manually install version 0.20.1 at the session prompt.

```bash
!pip install pandas==0.20.1
```

**R**

In R, DataFrames will display as grids by default. For example, to view the Iris data set, you would just use:

```r
iris
```

Similar to PySpark, bringing Sparklyr data into local memory with `as.data.frame` will output a grid display.

```r
sparklyr_df %>% as.data.frame
```

**Scala**

Calling the `display()` function on an existing dataframe will trigger a collect, much like `df.show()`.

```scala
val df = sc.parallelize(1 to 100).toDF()
display(df)
```
Documenting Your Analysis

Cloudera Data Science Workbench supports Markdown documentation of your code written in comments. This allows you to generate reports directly from valid Python and R code that runs anywhere, even outside Cloudera Data Science Workbench. To add documentation to your analysis, create comments in Markdown format:

R

```r
# Heading
# -------
# This documentation is **important.**
# Inline math: $e^x$
# Display math: $$y = \Sigma x + \epsilon$$
print("Now the code!")
```

Python

```python
# Heading
# -------
# This documentation is **important.**
# Inline math: $e^x$
# Display math: $$y = \Sigma x + \epsilon$$
print("Now the code!")
```

Using NVIDIA GPUs for Cloudera Data Science Workbench Projects

**Minimum Required Roles:** Cloudera Manager Cluster Administrator, CDSW Site Administrator

A GPU is a specialized processor that can be used to accelerate highly parallelized computationally-intensive workloads. Because of their computational power, GPUs have been found to be particularly well-suited to deep learning workloads. Ideally, CPUs and GPUs should be used in tandem for data engineering and data science workloads. A typical machine learning workflow involves data preparation, model training, model scoring, and model fitting. You can use existing general-purpose CPUs for each stage of the workflow, and optionally accelerate the math-intensive steps with the selective application of special-purpose GPUs. For example, GPUs allow you to accelerate model fitting using frameworks such as Tensorflow, PyTorch, Keras, MXNet, and Microsoft Cognitive Toolkit (CNTK).

By enabling GPU support, data scientists can share GPU resources available on Cloudera Data Science Workbench hosts. Users can requests a specific number of GPU instances, up to the total number available on a host, which are then allocated to the running session or job for the duration of the run. Projects can use isolated versions of libraries, and even different CUDA and cuDNN versions via Cloudera Data Science Workbench’s extensible engine feature.

**Prerequisite**

GPU support for workloads was added in Cloudera Data Science Workbench 1.1.0. The rest of this topic assumes you have already installed or upgraded to the latest version of Cloudera Data Science Workbench.

**Key Points to Note**

- Cloudera Data Science Workbench only supports CUDA-enabled NVIDIA GPU cards.
- Cloudera Data Science Workbench does not support heterogeneous GPU hardware in a single deployment.
- Cloudera Data Science Workbench does not include an engine image that supports NVIDIA libraries. Create your own custom CUDA-capable engine image using the instructions described in this topic.
Enabling Cloudera Data Science Workbench to use GPUs

To enable GPU usage on Cloudera Data Science Workbench, perform the following steps to provision the Cloudera Data Science Workbench hosts. As noted in the following instructions, certain steps must be repeated on all gateway hosts that have GPU hardware installed on them.

The steps described in this document have been tested and validated on the following setup:

<table>
<thead>
<tr>
<th>CDSW</th>
<th>OS &amp; Kernel</th>
<th>NVIDIA Driver</th>
<th>nvidia-docker</th>
<th>CUDA</th>
<th>Tensorflow</th>
</tr>
</thead>
<tbody>
<tr>
<td>1.8.0</td>
<td>RHEL 7.4 3.10.0-862.el7.x86_64</td>
<td>396.26</td>
<td>nvidia-docker-1.0.1-1</td>
<td>CUDA 9.2</td>
<td>1.8.0</td>
</tr>
</tbody>
</table>

Set Up the Operating System and Kernel

**Perform this step on all hosts with GPU hardware installed on them.**

1. Install the `kernel-devel` package.

   ```
sudo yum install -y kernel-devel-`uname -r`
   
   If the previous command fails to find a matching version of the `kernel-devel` package, list all the `kernel/kernel-devel` versions that are available from the RHEL/CentOS package repositories, and pick the desired version to install.

   You can use a bash script as demonstrated here to do this:
   ```
   if ! yum install kernel-devel-`uname -r`; then
     yum install -y kernel kernel-devel; retValue=$?
     if [ $retValue -eq 0 ]; then echo "Reboot is required since new version of kernel was installed"; fi
   fi
   ```

2. If you upgraded to a new kernel version in the previous step, run the following command to reboot.

   ```
sudo reboot
   ```

3. Install the Development tools package.

   ```
sudo yum groupinstall -y "Development tools"
   ```

Install the NVIDIA Driver on GPU Hosts

**Perform this step on all hosts with GPU hardware installed on them.**

Cloudera Data Science Workbench does not ship with any of the NVIDIA drivers needed to enable GPUs for general purpose processing. System administrators are expected to install the version of the drivers that are compatible with the CUDA libraries that will be consumed on each host.

Use the [NVIDIA UNIX Driver archive](#) to find out which driver is compatible with your GPU card and operating system. Make sure the driver you select is also compatible with the `nvidia-docker` plugin we will be installing in the next step. See [nvidia-docker installation prerequisites](#).
To download and install the NVIDIA driver, make sure you follow the instructions on the respective driver’s download page. It is crucial that you download the correct version.

For example, if you use the .run file method (Linux 64 bit), you would download and install the driver as follows:

```bash
wget http://us.download.nvidia.com/.../NVIDIA-Linux-x86_64--<driver_version>.run
export NVIDIA_DRIVER_VERSION=<driver_version>
chmod 755 ./NVIDIA-Linux-x86_64--$NVIDIA_DRIVER_VERSION.run
./NVIDIA-Linux-x86_64--$NVIDIA_DRIVER_VERSION.run -asq
```

Once the installation is complete, run the following command to verify that the driver was installed correctly:

```
/usr/bin/nvidia-smi
```

Enable Docker NVIDIA Volumes on GPU Hosts

**Perform this step on all hosts with GPU hardware installed on them.**

To enable Docker containers to use the GPUs, the previously installed NVIDIA driver libraries must be consolidated in a single directory named after the `<driver_version>`, and mounted into the containers. This is done using the `nvidia-docker` package, which is a thin wrapper around the Docker CLI and a Docker plugin.

The following sample steps demonstrate how to use `nvidia-docker` to set up the directory structure for the drivers so that they can be easily consumed by the Docker containers that will leverage the GPU. Perform these steps on all hosts with GPU hardware installed on them.

1. Download and install `nvidia-docker`. Use a version that is suitable for your deployment.

```bash
wget https://github.com/NVIDIA/nvidia-docker/releases/download/v1.0.1/nvidia-docker-1.0.1-1.x86_64.rpm
sudo yum install -y nvidia-docker-1.0.1-1.x86_64.rpm
```

2. Start the necessary services and plugins:

```bash
systemctl start nvidia-docker
systemctl enable nvidia-docker
```

3. Run a small container to create the Docker volume structure:

```bash
sudo nvidia-docker run --rm nvidia/cuda:9.2-devel-ubuntu16.04 nvidia-smi
```

4. Verify that the `/var/lib/nvidia-docker/volumes/nvidia_driver/$NVIDIA_DRIVER_VERSION/` directory was created.

5. Use the following Docker command to verify that Cloudera Data Science Workbench can access the GPU.

```bash
sudo docker run --net host \
--device=/dev/nvidiactl \
--device=/dev/nvidia-uvm \
--device=/dev/nvidia0 \
-v /var/lib/nvidia-docker/volumes/nvidia_driver/$NVIDIA_DRIVER_VERSION:/usr/local/nvidia/ \
-it nvidia/cuda:9.2-devel-ubuntu16.04 \
/usr/local/nvidia/bin/nvidia-smi
```

On a multi-GPU machine the output of this command will show exactly one GPU. This is because we have run this sample Docker container with only one device (/dev/nvidia0).

**Important:** Cloudera Data Science Workbench does not detect GPUs after a machine reboot. This is because certain NVIDIA modules do not load automatically after a reboot. Review the associated Known Issue and workaround [here](#).
Enable GPU Support in Cloudera Data Science Workbench

Minimum Required Cloudera Manager Role: Cluster Administrator

Depending on your deployment, use one of the following sets of steps to enable Cloudera Data Science Workbench to identify the GPUs installed:

CSD Deployments

1. Go to the CDSW service in Cloudera Manager. Click Configuration. Set the following parameters as directed in the following table.

<table>
<thead>
<tr>
<th>Enable GPU Support</th>
<th>Use the checkbox to enable GPU support for Cloudera Data Science Workbench workloads. When this property is enabled on a host that is equipped with GPU hardware, the GPU(s) will be available for use by Cloudera Data Science Workbench.</th>
</tr>
</thead>
<tbody>
<tr>
<td>NVIDIA Driver Library Path</td>
<td>Complete path to the NVIDIA driver libraries. In this example, the path would be, /var/lib/nvidia-docker/volumes/nvidia_driver/$NVIDIA_DRIVER_VERSION/</td>
</tr>
</tbody>
</table>

2. Restart the CDSW service in Cloudera Manager.
3. Test whether Cloudera Data Science Workbench is detecting GPUs.

RPM Deployments

1. Set the following parameters in /etc/cdsw/config/cdsw.conf on all Cloudera Data Science Workbench hosts. You must make sure that cdsw.conf is consistent across all hosts, irrespective of whether they have GPU hardware installed on them.

<table>
<thead>
<tr>
<th>NVIDIA_GPU_ENABLE</th>
<th>Set this property to true to enable GPU support for Cloudera Data Science Workbench workloads. When this property is enabled on a host that is equipped with GPU hardware, the GPU(s) will be available for use by Cloudera Data Science Workbench.</th>
</tr>
</thead>
<tbody>
<tr>
<td>NVIDIA_LIBRARY_PATH</td>
<td>Complete path to the NVIDIA driver libraries. In this example, the path would be, &quot;/var/lib/nvidia-docker/volumes/nvidia_driver/$NVIDIA_DRIVER_VERSION/&quot;</td>
</tr>
</tbody>
</table>

2. On the master host, run the following command to restart Cloudera Data Science Workbench.

```bash
cdsw restart
```

If you modified cdsw.conf on a worker host, run the following commands to make sure the changes go into effect:

```bash
cdsw reset
cdsw join
```

3. Use the following section to test whether Cloudera Data Science Workbench can now detect GPUs.

Test whether Cloudera Data Science Workbench can detect GPUs

Once Cloudera Data Science Workbench has successfully restarted, if NVIDIA drivers have been installed on the Cloudera Data Science Workbench hosts, Cloudera Data Science Workbench will now be able to detect the GPUs available on its hosts.
Additionally, the output of this command will also indicate that there are hosts with GPUs present.

```
cdsw status
```

**Create a Custom CUDA-capable Engine Image**

**Note:** Before you proceed, review the list of known issues and limitations associated with custom engines [here](#).

The base engine image (`docker.repository.cloudera.com/cdsw/engine:<version>`) that ships with Cloudera Data Science Workbench will need to be extended with CUDA libraries to make it possible to use GPUs in jobs and sessions.

The following sample Dockerfile illustrates an engine on top of which machine learning frameworks such as Tensorflow and PyTorch can be used. This Dockerfile uses a deep learning library from NVIDIA called NVIDIA CUDA Deep Neural Network (cuDNN). For detailed information about compatibility between NVIDIA driver versions and CUDA, refer the cuDNN installation guide (prerequisites).

Make sure you also check with the machine learning framework that you intend to use in order to know which version of cuDNN is needed. As an example, Tensorflow’s NVIDIA hardware and software requirements for GPU support are listed in the Tensorflow documentation [here](#). Additionally, the Tensorflow version compatibility matrix for CUDA and cuDNN is documented [here](#).

The following sample Dockerfile uses NVIDIA’s official Dockerfiles for CUDA and cuDNN images.

**cuda.Dockerfile**

```bash
FROM docker.repository.cloudera.com/cdsw/engine:5

RUN NVIDIA_GPGKEY_SUM=d1be581509378368edeeb8c0eb2eb958702feef3bc3d17011adbf24efacce4ab5 &
    NVIDIA_GPGKEY_FPR=ae09fe4b22da8422ccf36f4b3d7fa2af80 &
    apt-key adv --fetch-keys http://developer.download.nvidia.com/compute/cuda/repos/ubuntu1604/x86_64/7fa2af80.pub &
    apt-key adv --export --no-emit-version -a $NVIDIA_GPGKEY_FPR | tail -n +5 >
    cudasign.pub &
    echo "$NVIDIA_GPGKEY_SUM  cudasign.pub"  | sha256sum -c --strict - &
    rm cudasign.pub &
    echo "deb http://developer.download.nvidia.com/compute/cuda/repos/ubuntu1604/x86_64/" > /etc/apt/sources.list.d/cuda.list

ENV CUDA_VERSION 9.2.148
LABEL com.nvidia.cuda.version="$CUDA_VERSION"

ENV CUDA_PKG_VERSION 9-2-0
RUN apt-get update &
    apt-get install -y --no-install-recommends \
    cuda-cudart-"$CUDA_PKG_VERSION" &
    ln -s cuda-9.2 /usr/local/cuda &
    rm -rf /var/lib/apt/lists/*
```
You can now build a custom engine image out of cuda.Dockerfile using the following sample command:

```
docker build --network host -t <company-registry>/cdsw-cuda:5 . -f cuda.Dockerfile
```

Push this new engine image to a public Docker registry so that it can be made available for Cloudera Data Science Workbench workloads. For example:

```
docker push <company-registry>/cdsw-cuda:5
```

### Allocate GPUs for Sessions and Jobs

**Required CDSW Role:** [Site Administrator](#)

Once Cloudera Data Science Workbench has been enabled to use GPUs, a site administrator must whitelist the CUDA-capable engine image created in the previous step. Site administrators can also set a limit on the maximum number of GPUs that can be allocated per session or job.

1. Sign in to Cloudera Data Science Workbench.
2. Click **Admin**.
3. Go to the **Engines** tab.
4. From the **Maximum GPUs per Session/Job** dropdown, select the maximum number of GPUs that can be used by an engine.
5. Under **Engine Images**, add the custom CUDA-capable engine image created in the previous step. This whitelists the image and allows project administrators to use the engine in their jobs and sessions.
6. Click **Update**.

Project administrators can now whitelist the CUDA engine image to make it available for sessions and jobs within a particular project.

1. Navigate to the project’s **Overview** page.
2. Click **Settings**.
3. Go to the **Engines** tab.
4. Under **Engine Image**, select the CUDA-capable engine image from the dropdown.

### Example: Tensorflow

This is a simple example that walks you through a simple Tensorflow workload that uses GPUs.

1. Open the workbench console and start a Python session. Make sure you select at least 1 GPU from the **Select Engine Profile** dropdown before you launch the session.
2. Install Tensorflow.
3. Restart the session. This requirement of a session restart after installation is a known issue specific to Tensorflow.

4. Create a new file with the following sample code. The code first performs a multiplication operation and prints the session output, which should mention the GPU that was used for the computation. The latter half of the example prints a list of all available GPUs for this engine.

```python
import tensorflow as tf
a = tf.constant([1.0, 2.0, 3.0, 4.0, 5.0, 6.0], shape=[2, 3], name='a')
b = tf.constant([1.0, 2.0, 3.0, 4.0, 5.0, 6.0], shape=[3, 2], name='b')
c = tf.matmul(a, b)

# Creates a session with log_device_placement set to True.
sess = tf.Session(config=tf.ConfigProto(log_device_placement=True))
# Runs the operation.
print(sess.run(c))
# Prints a list of GPUs available
from tensorflow.python.client import device_lib
def get_available_gpus():
    local_device_protos = device_lib.list_local_devices()
    return [x.name for x in local_device_protos if x.device_type == 'GPU']
print(get_available_gpus())
```

### Accessing Web User Interfaces from Cloudera Data Science Workbench

This topic describes the different ways in which Cloudera Data Science Workbench allows you to access user interfaces for applications such as Cloudera Manager, Hue, and even the transient per-session UIs for frameworks such as Spark 2, TensorFlow, Shiny, and so on.

**Cloudera Manager, Hue, and the Spark History Server**

Cloudera Data Science Workbench also gives you a way to access your CDH cluster's Cloudera Manager and Hue UIs from within the Cloudera Data Science Workbench application. Spark 2 provides a UI that displays information and logs for completed Spark applications, which is useful for debugging and performance monitoring. This UI, called the History Server, runs on the CDH cluster, on a configurable host and port.

To access these applications, click the grid icon in the upper right hand corner of the Cloudera Data Science Workbench web application, and select the UI you want to visit from the dropdown.
Web UIs Embedded in Jobs and Sessions

Many data science libraries and processing frameworks include user interfaces to help track progress of your jobs and break down workflows. These are instrumental in debugging and using the platforms themselves. For example, Spark provides a Spark Web UI to monitor running applications and TensorFlow visualizations can be run on TensorBoard. Other web application frameworks such as Shiny and Flask are popular ways for data scientists to display additional interactive analysis in the languages they already know.

Cloudera Data Science Workbench allows you to access these web UIs directly from sessions and jobs. This feature is particularly helpful when you want to monitor and track progress for batch jobs. Even though jobs don’t give you access to the interactive workbench console, you can still track long running jobs through the UI. However, note that the UI is only active so long as the job/session is active. If your session times out after 60 minutes (default timeout value), so will the UI.

Cloudera Data Science Workbench exposes web applications in one of the following ways:

**Spark 2 Web UIs (CDSW_SPARK_PORT)**

Spark 2 exposes one web UI for each Spark application driver running in Cloudera Data Science Workbench. The UI will be running within the container, on the port specified by the environmental variable CDSW_SPARK_PORT. By default, CDSW_SPARK_PORT is set to **20049**. The web UI will exist only as long as a SparkContext is active within a session. The port is freed up when the SparkContext is shutdown.

Spark 2 web UIs are available in browsers at: https://spark-<$CDSW_ENGINE_ID>.<$CDSW_DOMAIN>. To access the UI while you are in an active session, click the grid icon in the upper right hand corner of the Cloudera Data Science Workbench web application, and select **Spark UI** from the dropdown. For a job, navigate to the job overview page and click the **History** tab. Click on a job run to open the session output for the job. You can now click the grid icon in the upper right hand corner of the Cloudera Data Science Workbench web application to access the Spark UI for this session.

**TensorBoard, Shiny, and others (CDSW_PUBLIC_PORT)**

CDSW_PUBLIC_PORT is an environment variable that points to a general purpose public port. By default, CDSW_PUBLIC_PORT is set to port **8080**. Any HTTP services running in containers that bind to CDSW_PUBLIC_PORT are available in browsers at: http://<$CDSW_ENGINE_ID>.<$CDSW_DOMAIN>. Therefore, TensorBoard, Shiny, Flask or any other web framework accompanying a project can be accessed directly from within a session or job, as long as it is run on CDSW_PUBLIC_PORT.

Note: In Cloudera Data Science Workbench 1.5.x, setting CDSW_PUBLIC_PORT to a non-default port number is not supported.

To access the UI while you are in an active session, click the grid icon in the upper right hand corner of the Cloudera Data Science Workbench web application, and select the UI from the dropdown. For a job, navigate to the job overview page and click the **History** tab. Click on a job run to open the session output for the job. You can now click the grid icon in the upper right hand corner of the Cloudera Data Science Workbench web application to access the UI for this session.

**Example: A Shiny Application**

This example demonstrates how to create and run a Shiny application and view the associated UI while in an active session.

Create a new, blank project and run an R console. Create the files, **ui.R** and **server.R**, in the project, and copy the contents of the following example files provided by [Shiny by RStudio](https://shiny.rstudio.com):

```R
# ui.R
library(shiny)
```
# Define UI for application that draws a histogram
shinyUI(fluidPage(
    # Application title
titlePanel("Hello Shiny!"),
    # Sidebar with a slider input for the number of bins
    sidebarLayout(
        sidebarPanel(
            sliderInput("bins", "Number of bins:",
                min = 1,
                max = 50,
                value = 30)
        ),
        # Show a plot of the generated distribution
        mainPanel(
            plotOutput("distPlot")
        )
    )
))

# server.R
library(shiny)

# Define server logic required to draw a histogram
shinyServer(function(input, output) {
    # Expression that generates a histogram. The expression is
    # wrapped in a call to renderPlot to indicate that:
    # 1) It is "reactive" and therefore should re-execute automatically
    #    when inputs change
    # 2) Its output type is a plot
    output$distPlot <- renderPlot({
        x <- faithful[, 2]  # Old Faithful Geyser data
        bins <- seq(min(x), max(x), length.out = input$bins + 1)

        # draw the histogram with the specified number of bins
        hist(x, breaks = bins, col = 'darkgray', border = 'white')
    })
})

Run the following code in the interactive workbench prompt to install the Shiny package, load the library into the engine, and run the Shiny application.

```
R
install.packages('shiny')
library('shiny')
runApp(port=as.numeric(Sys.getenv("CDSW_PUBLIC_PORT")),
      host=Sys.getenv("CDSW_IP_ADDRESS"),
      launch.browser="FALSE")
```

Finally, click the grid icon in the upper right hand corner of the Cloudera Data Science Workbench web application, and select the Shiny UI, **Hello Shiny!**, from the dropdown. The UI will be active as long as the session is still running.

## Distributed Computing with Workers

For distributed computing, such as cross-validating a model or tuning some hyper-parameters, Cloudera Data Science Workbench provides basic support for leveraging multiple engine instances from a single run. Any R or Python engine
can spawn other engines, known as workers, and give them code to execute when they start up. Worker output is displayed in the main console to allow you to debug your code. These workers are terminated when the session exits.

For more significant distributed computing needs, using CDS 2.x Powered by Apache Spark from within Cloudera Data Science Workbench is strongly recommended.

Spawning Workers

Select a language from the code samples below to launch workers:

**R**

```r
library("cdsw")
workers <- launch.workers(n=2, cpu=0.2, memory=0.5, env="", code="print('Hello from a
cDSW Worker')")
```

**Note:** The `env` parameter has been defined due to a bug that appears when parsing the `launch.workers` function. When not defined, the `env` parameter is serialized internally into a format that is incompatible with Cloudera Data Science Workbench. This bug does not affect the Python engine.

**Python**

```python
import cdsw
workers = cdsw.launch_workers(n=2, cpu=0.2, memory=0.5, code="print 'Hello from a CDSW
Worker'")
```

Worker Network Communication

Workers are a low-level feature to help use higher level libraries that can operate across multiple hosts. As such, you will generally want to use workers only to launch the backends for these libraries.

To help you get your workers or distributed computing framework components talking to one another, every worker engine run includes an environmental variable `CDSW_MASTER_IP` with the fully addressable IP of the master engine. Every engine has a dedicated IP access with no possibility of port conflicts.

For instance, the following are trivial examples of two worker engines talking to the master engine.

**R**

From the master engine, the following `master.r` script will launch two workers and accept incoming connections from them.

```r
# master.r
library("cdsw")

# Launch two CDSW workers. These are engines that will run in
# the same project, execute a given code or script, and exit.
workers <- launch.workers(n=2, cpu=0.2, memory=0.5, env="", script="worker.r")

# Accept two connections, one from each worker. Workers will
# execute worker.r.
for(i in c(1,2)) {
  # Receive a message from each worker and return a response.
  con <- socketConnection(host="0.0.0.0", port = 6000, blocking=TRUE, server=TRUE, open="r+")
  data <- readLines(con, 1)
  print(paste("Server received:", data))
  writeLines("Hello from master!", con)
  close(con)
}
```
The workers will execute the following `worker.r` script and respond to the master.

```r
# worker.r
print(Sys.getenv("CDSW_MASTER_IP"))
con <- socketConnection(host=Sys.getenv("CDSW_MASTER_IP"), port = 6000, blocking=TRUE, server=FALSE, open="r+")
write_resp <- writeLines("Hello from Worker", con)
server_resp <- readLines(con, 1)
print(paste("Worker received: ", server_resp))
close(con)
```

Python

From the master engine, the following `master.py` script will launch two workers and accept incoming connections from them.

```python
# master.py
import cdsw, socket
# Launch two CDSW workers. These are engines that will run in
# the same project, execute a given code or script, and exit.
workers = cdsw.launch_workers(n=2, cpu=0.2, memory=0.5, script="worker.py")

# Listen on TCP port 6000
s = socket.socket(socket.AF_INET, socket.SOCK_STREAM)
s.bind(("0.0.0.0", 6000))
s.listen(1)

# Accept two connections, one from each worker. Workers will
# execute worker.py.
for i in range(2):
    # Receive a message from each worker and return a response.
    data = s.recv(20)
    if not data: break
    print "Master received:", data
    s.send("Hello From Server!")
s.close()
```

The workers will execute the following `worker.py` script and respond to the master.

```python
# worker.py
import os, socket
# Open a TCP connection to the master.
s = socket.socket(socket.AF_INET, socket.SOCK_STREAM)
s.connect((os.environ["CDSW_MASTER_IP"], 6000))

# Send some data and receive a response.
s.send("Hello From Worker!")
data = s.recv(1024)
s.close()
print "Worker received:", data
```
Collaborating on Projects with Cloudera Data Science Workbench

Cloudera Data Science Workbench supports several collaboration models.

Project Collaborators

If you want to work closely with trusted colleagues on a particular project, you can add them to the project as collaborators. For instructions, see Adding Collaborators on page 108.

Warning:

Collaborating Securely on Projects

Before adding project collaborators, you must remember that assigning the Contributor or Admin role to a project collaborator is the same as giving them write access to your data in CDH. This is because project contributors and project administrators have write access to all your project code (including any library code that you might not be actively inspecting). For example, a contributor/admin could modify project file(s) to insert code that deletes some data on the CDH cluster. The next time you launch a session and run the same code, it will appear as though you deleted the data yourself.

Additionally, project collaborators also have access to all actively running sessions and jobs. This means that a malicious user can easily impersonate you by accessing one of your active sessions. Therefore, it is extremely important to restrict project access to trusted collaborators only. Note that Cloudera Data Science Workbench 1.4.3 introduces a new feature that allows site administrators to restrict this ability by allowing only session creators to execute commands within their own active sessions. For details, see Restricting Access to Active Sessions.

For these reasons, Cloudera recommends using Git to collaborate securely on shared projects. This will also help avoid file modification conflicts when your team is working on more elaborate projects.

Restricting Collaborator and Administrator Access to Active Sessions

Required Role: Site Administrator

By default, the following Cloudera Data Science Workbench users have the ability to execute commands within any active sessions you have created:

- All Site Administrators
- Users who have been assigned Admin or Contributor privileges for the project where the session is created.
- For team projects, Team Admins have complete access to all team projects and any active sessions running within these projects. Additionally, any team members who have been assigned the Admin or Contributor roles for your projects will also have the ability to execute commands within your active sessions.

Starting with Cloudera Data Science Workbench 1.4.3, site administrators can now restrict this ability by allowing only the user who launched the session to execute commands within their own active sessions. To enable this restriction:

1. Log into Cloudera Data Science Workbench with site administrator privileges.
2. Click Admin > Security.
3. Under the General section, select the checkbox to enable the Only session creators can execute commands on active sessions property.

When this property is enabled, only the user that creates a session will be able to execute commands in that session. No other users, regardless of their permissions in the team or as project collaborators, will be able to execute commands on active sessions that are not created by them. Even site administrators will not be able to execute commands in other users’ active sessions. However, keep in mind that all site administrators still have access to the Site Administrator dashboard and can reverse this change at any time.
Teams

Users who work together on more than one project and want to facilitate collaboration can create a Team. Teams allow streamlined administration of projects. Team projects are owned by the team, rather than an individual user. Team administrators can add or remove members at any time, assigning each member different permissions.

For more details, see:

- Creating a Team on page 103
- Modifying Team Account Settings on page 104

Sharing Personal Projects

When you create a project in your personal context, Cloudera Data Science Workbench asks you to assign one of the following visibility levels to the project - Private or Public. Public projects on Cloudera Data Science Workbench grant read-level access to everyone with access to the Cloudera Data Science Workbench application. That means everyone can view the project’s files and results, but only those whom you have explicitly added as a collaborator can edit files, run engines, or view the project’s environment variables.

You can include a markdown-formatted README.md file in public projects to document your project’s purpose and usage.

If you are a project admin, you can set a project’s visibility to Public from the Project > Settings > Options page. For instructions, see Modifying Project Settings on page 108.

Forking Projects

You can fork another user’s project by clicking Fork on the Project page. Forking creates a new project under your account that contains all the files, libraries, configuration, and jobs from the original project.

Creating sample projects that other users can fork helps to bootstrap new projects and encourage common conventions.

Note: An issue exists where a timeout might occur when forking large projects.

Collaborating with Git

Cloudera Data Science Workbench provides seamless access to Git projects. Whether you are working independently, or as part of a team, you can leverage all of benefits of version control and collaboration with Git from within Cloudera Data Science Workbench. Teams that already use Git for collaboration can continue to do so. Each team member will need to create a separate Cloudera Data Science Workbench project from the central Git repository.

For anything but simple projects, Cloudera recommends using Git for version control. You should work on Cloudera Data Science Workbench the same way you would work locally, and for most data scientists and developers that means using Git.

For more details, see Using Git to Collaborate on Projects on page 130.

Sharing Job and Session Console Outputs

Cloudera Data Science Workbench lets you easily share the results of your analysis with one click. Using rich visualizations and documentation comments, you can arrange your console log so that it is a readable record of your analysis and results. This log continues to be available even after the session stops. This method of sharing allows you to show colleagues and collaborators your progress without your having to spend time creating a report.
To share results from an interactive session, click **Share** at the top of the console page. From here you can generate a link that includes a secret token that gives access to that particular console output. For jobs results, you can either share a link to the latest job result or a particular job run. To share the latest job result, click the **Latest Run** link for a job on the Overview page. This link will always have the latest job results. To share a particular run, click on a job run in the job's **History** page and share the corresponding link.

You can share console outputs with one of the following sets of users.

- **All anonymous users with the link** - By default, Cloudera Data Science Workbench allows anonymous access to shared consoles. However, site administrators can disable anonymous sharing at any time by going to **Admin > Security**, disabling the **Allow anonymous access to shared console outputs** checkbox, and clicking **Disable anonymous access** to confirm.

  Once anonymous sharing has been disabled, all existing publicly shared console outputs will be updated to be viewable only by authenticated users.

- **All authenticated users with the link** - This means any user with a Cloudera Data Science Workbench account will have access to the shared console.

- **Specific users and teams** - Click **Change** to search for users and teams to give access to the shared console. You can also come back to the session and revoke access from a user or team the same way.

### Sharing Data Visualizations

If you want to share a single data visualization rather than an entire console, you can embed it in another web page. Click the small circular ‘link’ button located to the left of most rich visualizations to view the HTML snippet that you can use to embed the visualization.

### Using Git to Collaborate on Projects

Cloudera Data Science Workbench provides seamless access to Git projects. Whether you are working independently, or as part of a team, you can leverage all of benefits of version control and collaboration with Git from within Cloudera Data Science Workbench. Teams that already use Git for collaboration can continue to do so. Each team member will need to create a separate Cloudera Data Science Workbench project from the central Git repository.

For anything but simple projects, Cloudera recommends using Git for version control. You should work on Cloudera Data Science Workbench the same way you would work locally, and for most data scientists and developers that means using Git.

Cloudera Data Science Workbench does not include significant UI support for Git, but instead allows you to use the full power of the command line. If you run an engine and open a terminal, you can run any Git command, including `init`, `add`, `commit`, `branch`, `merge` and `rebase`. Everything should work exactly as it does locally, except that you are running on a distributed edge host directly connected to your Apache Hadoop cluster.

### Importing a Project From Git

When you create a project, you can optionally supply an HTTPS or SSH Git URL that points to a remote repository. The new project is a clone of that remote repository. You can commit, push and pull your code by running a console and opening a terminal.

**Using SSH** - If you want to use SSH to clone the repo, you will need to first add your personal Cloudera Data Science Workbench SSH key to your GitHub account. For instructions, see **Adding SSH Key to GitHub** on page 245.

If you see Git commands hanging indefinitely, check with your cluster administrators to make sure that the SSH ports on the Cloudera Data Science Workbench hosts are not blocked.

### Linking an Existing Project to a Git Remote

If you did not create your project from a Git repository, you can link an existing project to a Git remote (for example, `git@github.com:username/repo.git`) so that you can push and pull your code.
To link to a Git remote:

1. Launch a new session.
2. Open a terminal.
3. Enter the following commands:

   ```shell
   git init
   git add *
   git commit -a -m 'Initial commit'
   git remote add origin git@github.com:username/repo.git
   ```

You can run `git status` after `git init` to make sure your `.gitignore` includes a folder for libraries and other non-code artifacts.
Importing Data into Cloudera Data Science Workbench

Cloudera Data Science Workbench allows you to run analytics workloads on data imported from local files, Apache HBase, Apache Kudu, Apache Impala, Apache Hive or other external data stores such as Amazon S3.

Accessing Local Data from Your Computer

If you want to perform analytics operations on existing data files (.csv, .txt, etc.) from your computer, you can upload these files directly to your Cloudera Data Science Workbench project. Go to the project's Overview page. Under the Files section, click Upload and select the relevant data files to be uploaded.

The following sections use the tips.csv dataset to demonstrate how to work with local data stored within your project. Upload this dataset to the data folder in your project before you run these examples.

Pandas (Python)

```python
import pandas as pd

tips = pd.read_csv('data/tips.csv')

tips
  .query('sex == "Female"')
  .groupby('day')
  .agg({"tip": 'mean'})
  .rename(columns={"tip": 'avg_tip_dinner'})
  .sort_values('avg_tip_dinner', ascending=False)
```

dplyr (R)

```r
library(readr)
library(dplyr)

# load data from .csv file in project

tips <- read_csv("data/tips.csv")

# query using dplyr

tips %>%
  filter(sex == "Female") %>%
  group_by(day) %>%
  summarise(
    avg_tip = mean(tip, na.rm = TRUE)
  ) %>%
  arrange(desc(avg_tip))
```

Accessing Data from HDFS

There are many ways to access HDFS data from R, Python, and Scala libraries. The following code samples demonstrate how to count the number of occurrences of each word in a simple text file in HDFS.

Navigate to your project and click Open Workbench. Create a file called sample_text_file.txt and save it to your project in the data folder. Now write this file to HDFS. You can do this in one of the following ways:

- Click Terminal above the Cloudera Data Science Workbench console and enter the following command to write the file to HDFS:

```bash
hdfs dfs -put data/sample_text_file.txt /tmp
```

OR
• Use the workbench command prompt:

**Python Session**
```bash
!hdfs dfs -put data/sample_text_file.txt /tmp
```

**R Session**
```r
system("hdfs dfs -put data/tips.csv /user/hive/warehouse/tips/")
```

The following examples use Python and Scala to read `sample_text_file.txt` from HDFS (written above) and perform the count operation on it.

**Python**
```python
from __future__ import print_function
import sys, re
from operator import add
from pyspark.sql import SparkSession

spark = SparkSession.builder.
    .appName("PythonWordCount")
    .getOrCreate()

# Access the file
lines = spark.read.text("/tmp/sample_text_file.txt").rdd.map(lambda r: r[0])
counts = lines.flatMap(lambda x: x.split(' ')) 
    .map(lambda x: (x, 1)) 
    .reduceByKey(add) 
    .sortBy(lambda x: x[1], False)
output = counts.collect()
for (word, count) in output:
    print("%s: %i" % (word, count))
spark.stop()
```

**Scala**
```scala
//count lower bound
val threshold = 2

// read the file added to hdfs
val tokenized = sc.textFile("/tmp/sample_text_file.txt").flatMap(_.split(" "))

// count the occurrence of each word
val wordCounts = tokenized.map((_ , 1)).reduceByKey(_ + _)

// filter out words with fewer than threshold occurrences
val filtered = wordCounts.filter(_._2 >= threshold)
System.out.println(filtered.collect().mkString(",")
```

**Accessing Data from Apache HBase**

This section demonstrates how to use the HappyBase Python library to access data from HBase.

**Load Data into HBase Table**

For this example, we’re going to import data from a CSV file into HBase using the `importTsv` package.

1. Log into Cloudera Data Science Workbench and launch a Python 3 session within a new/existing project.
2. For this example, we will be using the following sample CSV file. Create the following `employees.csv` file in your project.
3. In the workbench, click **Terminal access**. Perform the following steps in the Terminal:

   a. Start the HBase shell and create a new blank table called **employees**.

   ```sh
hbase shell
create 'employees', 'name', 'department'
exit
```

   b. Load **employees.csv** into HDFS.

   ```sh
hdfs dfs -put employees.csv /tmp
```

   c. Use **ImportTsv** to load data from HDFS (/tmp/employees.csv) into the HBase table created in the previous step.

   ```sh
hbase org.apache.hadoop.hbase.mapreduce.ImportTsv -Dimporttsv.separator=',' -Dimporttsv.columns=HBASE_ROW_KEY,name,department employees /tmp/employees.csv
```

   d. Go back to the HBase shell and run the following command to make sure data was loaded into the HBase table.

   ```sh
hbase shell
scan 'employees'
```

### Query Data Using HappyBase

1. Launch a Python 3 session and use the workbench command prompt to install the **happybase** package.

   ```sh
!pip3 install happybase
```

2. Use **happybase** to connect to the **employees** table created in the previous step.

   ```py
import happybase
connection = happybase.Connection(host='<hbase_thrift_server_hostname>', port=9090, autoconnect=True)
table = connection.table('employees')
rows = table.rows(["1","2","3"])
for key, data in rows:
    print(key, data)
```

### Accessing Data from Apache Hive

The following code sample demonstrates how to establish a connection with the Hive metastore and access data from tables in Hive.

```py
import os
import pandas
from impala.dbapi import connect
from impala.util import import as_pandas
```
# Specify HIVE_HMS_HOST as an environment variable in your project settings
HIVE_HMS_HOST = os.getenv('HIVE_HS2_HOST', '<hiveserver2_hostname>

# This connection string depends on your cluster setup and authentication mechanism
conn = connect(host=HIVE_HS2_HOST,
                port='10000',
                auth_mechanism='GSSAPI',
                kerberos_service_name='hive')
cursor = conn.cursor()
cursor.execute('SHOW TABLES')
tables = as_pandas(cursor)
tables

## Accessing Data from Apache Impala

In this section, we take some sample data in the form of a CSV file, save the contents of this file to a table in Impala, and then use some common Python and R libraries to run simple queries on this data.

### Loading CSV Data into an Impala Table

For this demonstration, we will be using the `tips.csv` dataset. Use the following steps to save this file to a project in Cloudera Data Science Workbench, and then load it into a table in Apache Impala.

1. Create a new Cloudera Data Science Workbench project.
2. Create a folder called `data` and upload `tips.csv` to this folder. For detailed instructions, see Managing Project Files.
3. The next steps require access to services on the CDH cluster. If Kerberos has been enabled on the cluster, enter your credentials (username, password/keytab) in Cloudera Data Science Workbench to enable access. For instructions, see Hadoop Authentication with Kerberos for Cloudera Data Science Workbench on page 236.
4. Navigate back to the project Overview page and click Open Workbench.
5. Launch a new session (Python or R).
6. Open the Terminal.

   a. Run the following command to create an empty table in Impala called `tips`. Replace `<impala_daemon_hostname>` with the hostname for your Impala daemon.

   ```
   impala-shell -i <impala_daemon_hostname>:21000 -q 'CREATE TABLE default.tips (  `total_bill` FLOAT,  `tip` FLOAT,  `sex` STRING,  `smoker` STRING,  `day` STRING,  `time` STRING,  `size` TINYINT)  ROW FORMAT DELIMITED FIELDS TERMINATED BY ","  LOCATION "hdfs:///user/hive/warehouse/tips/";'
   ```

   b. Run the following command to load data from the `/data/tips.csv` file into the Impala table.

   ```
   hdfs dfs -put data/tips.csv /user/hive/warehouse/tips/
   ```

### Running Queries on Impala Tables

This section demonstrates how to run queries on the `tips` table created in the previous section using some common Python and R libraries such as Pandas, Impyla, Sparklyr and so on. All the examples in this section run the same query, but use different libraries to do so.
PySpark (Python)

```python
from pyspark.sql import SparkSession
spark = SparkSession.builder.master('yarn').getOrCreate()

# load data from .csv file in HDFS
# tips = spark.read.csv("/user/hive/warehouse/tips/", header=True, inferSchema=True)
# OR load data from table in Hive metastore
tips = spark.table('tips')

from pyspark.sql.functions import col, lit, mean

# query using DataFrame API
tips
  .filter(col('sex').like("%Female%"))
  .groupBy('day')
  .agg(mean('tip').alias('avg_tip'))
  .orderBy('avg_tip',ascending=False)
  .show()

# query using SQL
spark.sql('''
SELECT day,AVG(tip) AS avg_tip
FROM tips
WHERE sex LIKE "%Female%"
GROUP BY day
ORDER BY avg_tip DESC''').show()

spark.stop()
```

Impyla (Python)

Due to an incompatibility with the `thrift_sasl` package, Impyla has been known to fail with Python 3.

Python 2

```python
# (Required) Install the impyla package
# !pip install impyla
# !pip install thrift_sasl
import os
import pandas
from impala.dbapi import connect
from impala.util import as_pandas

# Connect to Impala using Impyla
# Secure clusters will require additional parameters to connect to Impala.
# Recommended: Specify IMPALA_HOST as an environment variable in your project settings
IMPALA_HOST = os.getenv('IMPALA_HOST', '<impala_daemon_hostname>')
conn = connect(host=IMPALA_HOST, port=21050)

cursor = conn.cursor()
cursor.execute('''
SELECT day,AVG(tip) AS avg_tip
FROM tips
WHERE sex ILIKE "%Female%"
GROUP BY day
ORDER BY avg_tip DESC''')

# Pretty output using Pandas
tables = as_pandas(cursor)
tables
```
Ibis (Python)

```python
# (Required) Install the ibis-framework[impala] package
# !pip3 install ibis-framework[impala]

import ibis
import os
ibis.options.interactive = True
ibis.options.verbose = True

# Connection to Impala
# Secure clusters will require additional parameters to connect to Impala.
# Recommended: Specify IMPALA_HOST as an environment variable in your project settings

IMPALA_HOST = os.getenv('IMPALA_HOST', '<impala_daemon_hostname>')</con = ibis.impala.connect(host=IMPALA_HOST, port=21050, database='default')
con.list_tables()
tips = con.table('tips')
tips 
  .filter(tips.sex.like(['%Female%'])) 
  .group_by('day') 
  .aggregate(
    avg_tip=tips.tip.mean() 
  ) 
  .sort_by(ibis.desc('avg_tip')) 
  .execute()
```

Sparklyr (R)

```r
# (Required) Install the sparklyr package
# install.packages("sparklyr")

library(stringr)
library(sparklyr)
library(dplyr)

spark <- spark_connect(master = "yarn")

# load data from file in HDFS
tips <- spark_read_csv(
  sc = spark,
  name = "tips",
  path = "/user/hive/warehouse/tips/"
)

# OR load data from table
tips <- tbl(spark, "tips")

# query using dplyr
tips %>%
  filter(sex %like% "%Female") %>%
  group_by(day) %>%
  summarise(
    avg_tip = mean(tip, na.rm = TRUE)
  ) %>%
  arrange(desc(avg_tip))

# query using SQL
tbl(spark, sql("SELECT day,AVG(tip) AS avg_tip 
  FROM tips 
  WHERE sex LIKE '%Female%' 
  GROUP BY day 
  ORDER BY avg_tip DESC"))

spark_disconnect(spark)
```
Accessing Data in Amazon S3 Buckets

Every language in Cloudera Data Science Workbench has libraries available for uploading to and downloading from Amazon S3.

To work with S3:

1. Add your Amazon Web Services access keys to your project's environment variables as AWS_ACCESS_KEY_ID and AWS_SECRET_ACCESS_KEY.
2. Pick your favorite language from the code samples below. Each one downloads the R 'Old Faithful' dataset from S3.

```{r}
library("devtools")
install_github("armstrtw/AWS.tools")
Sys.setenv("AWSACCESSKEY"=Sys.getenv("AWS_ACCESS_KEY_ID"))
Sys.setenv("AWSSECRETKEY"=Sys.getenv("AWS_SECRET_ACCESS_KEY"))
library("AWS.tools")
s3.get("s3://sense-files/faithful.csv")
```

```{python}
# Install Boto to the project
!pip install boto
# Create the Boto S3 connection object.
from boto.s3.connection import S3Connection
aws_connection = S3Connection()
# Download the dataset to file 'faithful.csv'.
bucket = aws_connection.get_bucket('sense-files')
key = bucket.get_key('faithful.csv')
key.get_contents_to_filename('/home/cdsw/faithful.csv')
```

Accessing External SQL Databases

Every language in Cloudera Data Science Workbench has multiple client libraries available for SQL databases.

If your database is behind a firewall or on a secure server, you can connect to it by creating an SSH tunnel to the server, then connecting to the database on localhost.

If the database is password-protected, consider storing the password in an environmental variable to avoid displaying it in your code or in consoles. The examples below show how to retrieve the password from an environment variable and use it to connect.

```{r}
# dplyr lets you program the same way with local data frames and remote SQL databases.
install.packages("dplyr")
library("dplyr")
db <- src_postgres(dbname="test_db", host="localhost", port=5432, user="cdswuser",
password=Sys.getenv("POSTGRESQL_PASSWORD"))
flights_table <- tbl(db, "flights")
select(flights_table, year:day, dep_delay, arr_delay)
```
Python

You can access data using `pyodbc` or `SQLAlchemy`.

```python
import os

db = pyodbc.connect("DRIVER={PostgreSQL Unicode};SERVER=localhost;PORT=5432;DATABASE=test_db;USER=cdswuser;OPTION=3;PASSWORD=%s" % os.environ["POSTGRESQL_PASSWORD"])
cursor = cnxn.cursor()
cursor.execute("select user_id, user_name from users")
```

```python
import sqlalchemy
import os
from sqlalchemy.orm import sessionmaker
from sqlalchemy import create_engine

db = create_engine("postgresql://cdswuser:%s@localhost:5432/test_db" % os.environ["POSTGRESQL_PASSWORD"])
session = sessionmaker(bind=db)
user = session.query(User).filter_by(name='ed').first()
```
Experiments

Starting with version 1.4, Cloudera Data Science Workbench allows data scientists to run batch experiments that track different versions of code, input parameters, and output (both metrics and files).

**Demo:** Watch the following video for a quick demonstration of the steps described in this topic: Experiments with Cloudera Data Science Workbench

**Related:**
- Engines for Experiments and Models on page 171
- Debugging Issues with Experiments on page 145

Purpose

**Challenge**

As data scientists iteratively develop models, they often experiment with datasets, features, libraries, algorithms, and parameters. Even small changes can significantly impact the resulting model. This means data scientists need the ability to iterate and repeat similar experiments in parallel and on demand, as they rely on differences in output and scores to tune parameters until they obtain the best fit for the problem at hand. Such a training workflow requires versioning of the file system, input parameters, and output of each training run.

Without versioned experiments you would need intense process rigor to consistently track training artifacts (data, parameters, code, etc.), and even then it might be impossible to reproduce and explain a given result. This can lead to wasted time/effort during collaboration, not to mention the compliance risks introduced.

**Solution**

Starting with version 1.4, Cloudera Data Science Workbench uses experiments to facilitate ad-hoc batch execution and model training. Experiments are batch executed workloads where the code, input parameters, and output artifacts are versioned. This feature also provides a lightweight ability to track output data, including files, metrics, and metadata for comparison.

**Concepts**

The term experiment refers to a non interactive batch execution script that is versioned across input parameters, project files, and output. Batch experiments are associated with a specific project (much like sessions or jobs) and have no notion of scheduling; they run at creation time. To support versioning of the project files and retain run-level artifacts and metadata, each experiment is executed in an isolated container.

**Lifecycle of an Experiment**

The rest of this section describes the different stages in the lifecycle of an experiment - from launch to completion.
1. **Launch Experiment**

   In this step you will select a script from your project that will be run as part of the experiment, and the resources (memory/GPU) needed to run the experiment. The engine kernel will be selected by default based on your script. For detailed instructions on how to launch an experiment, see Running an Experiment (QuickStart) on page 141.

2. **Build**

   When you launch the experiment, Cloudera Data Science Workbench first builds a new versioned engine image where the experiment will be executed in isolation. This new engine includes:
   
   - the base engine image used by the project (check **Project > Settings**)
   - a snapshot of the project filesystem
   - environmental variables inherited from the project.
   - packages explicitly specified in the project's build script (`cdsw-build.sh`)

   It is your responsibility to provide the complete list of dependencies required for the experiment via the `cdsw-build.sh` file. As part of the engine's build process, Cloudera Data Science Workbench will run the `cdsw-build.sh` script and install the packages or libraries requested there on the new image.

   For details about the build process and examples on how to specify dependencies, see Engines for Experiments and Models on page 171.

3. **Schedule**

   Once the engine is built the experiment is scheduled for execution like any other job or session. Once the requested CPU/GPU and memory have been allocated to the experiment, it will move on to the execution stage.

   Note that if your deployment is running low on memory and CPU, your runs may spend some time in this stage.

4. **Execute**

   This is the stage where the script you have selected will be run in the newly built engine environment. This is the same output you would see if you had executed the script in a session in the Workbench console.

   You can watch the execution in progress in the individual run's **Session** tab.

   You can also go to the project **Overview > Experiments** page to see a table of all the experiments launched within that project and their current status.

   **Run ID:** A numeric ID that tracks all experiments launched on a Cloudera Data Science Workbench deployment. It is not limited to the scope of a single user or project.

---

**Running an Experiment (QuickStart)**

The following steps describe how to launch an experiment from the Workbench console. In this example we are going to run a simple script that adds all the numbers passed as arguments to the experiment.

1. Go to the project **Overview** page.
2. Click **Open Workbench**.
3. Create/modify any project code as needed. You can also launch a session to simultaneously test code changes on the interactive console as you launch new experiments.

   As an example, you can run this Python script that accepts a series of numbers as command-line arguments and prints their sum.

   ```python
   import sys
   import cdsw
   ```
To test the script, launch a Python session and run the following command from the workbench command prompt:

```
!python add.py 1 2 3 4
```

4. Click Run Experiment. If you’re already in an active session, click Run > Run Experiment. Fill out the following fields:

- **Script** - Select the file that will be executed for this experiment.
- **Arguments** - If your script requires any command line arguments, enter them here.

  **Note:** Arguments are not supported with Scala experiments.

- **Engine Kernel and Engine Profile** - Select the kernel and computing resources needed for this experiment.

For this example we will run the `add.py` script and pass some numbers as arguments.

5. Click Start Run.

6. To track progress for the run, go back to the project Overview. On the left navigation bar click Experiments. You should see the experiment you’ve just run at the top of the list. Click on the Run ID to view an overview for each individual run. Then click Build.

On this Build tab you can see realtime progress as Cloudera Data Science Workbench builds the Docker image for this experiment. This allows you to debug any errors that might occur during the build stage.
7. Once the Docker image is ready, the run will begin execution. You can track progress for this stage by going to the Session tab.

For example, the Session pane output from running `add.py` is:

```
> import sys
> import cdsw
> args = len(sys.argv) - 1
> sum = 0
> x = 1
> while (args >= x):
    print ('Parameter %i: %s' % (x, sys.argv[x]))
    sum = sum + int(sys.argv[x])
    x = x + 1
Parameter 1: 18
Parameter 2: 90
Parameter 3: 34
> print ('Sum of the numbers is: %i.' % sum)
Sum of the numbers is: 142.
```

8. (Optional) The `cdsw` library that is bundled with Cloudera Data Science Workbench includes some built-in functions that you can use to compare experiments and save any files from your experiments.

For example, to track the sum for each run, add the following line to the end of the `add.py` script.

```python
cdsw.track_metric("Sum", sum)
```

This will be tracked in the Experiments table:
For more details, see Tracking Metrics on page 144 and Saving Files on page 144.

Tracking Metrics

The cdsw library includes a `track_metric` function that can be used to log up to 50 metrics associated with a run, thus allowing accuracy and scores to be tracked over time.

The function accepts input in the form of key value pairs.

```
    cdsw.track_metric(key, value)
```

**Python**

```
    cdsw.track_metric("R_squared", 0.79)
```

**R**

```
    cdsw::track.metric("R_squared", 0.62)
```

These metrics will be available on the project's Experiments tab where you can view, sort, and filter experiments on the values. The table on the Experiments page will allow you to display only three metrics at a time. You can select which metrics are displayed from the metrics dropdown.

**Note:** This function is not supported with Scala experiments.

Saving Files

Cloudera Data Science Workbench allows you to select which artifacts you'd like to access and evaluate after an experiment is complete. These artifacts could be anything from a text file to an image or a model that you have built through the run.

The cdsw library includes a `track_file` function that can be used to specify which artifacts should be retained after the experiment is complete.

**Python**

```
    cdsw.track_file('model.pkl')
```

**R**

```
    cdsw::track.file('model.pkl')
```

Specified artifacts can be accessed from the run's Overview page. These files can also be saved to the top-level project filesystem and downloaded from there.

**Note:** This function is not supported with Scala experiments.

Disabling the Experiments Feature

**Required Role:** Site Administrator
Important: The feature flag mentioned here only hides the Experiments feature from the UI. It will not stop any experiments that have already been queued for execution.

To disable this feature on your Cloudera Data Science Workbench deployment:

1. Log in to Cloudera Data Science Workbench.
2. Click Admin > Settings.
3. Under the Feature Flags section, disable the Enable users to run experiments checkbox.

Limitations

- Experiments do not store snapshots of project files. You cannot automatically restore code that was run as part of an experiment.
- Experiments will fail if your project filesystem is too large for the Git snapshot process. As a general rule, any project files (code, generated model artifacts, dependencies, etc.) larger than 50 MB must be part of your project's .gitignore file so that they are not included in snapshots for experiment builds.
- Experiments cannot be deleted. As a result, be conscious of how you use the track_metrics and track_file functions:
  - Do not track files larger than 50MB.
  - Do not track more than 100 metrics per experiment. Excessive metric calls from an experiment may cause Cloudera Data Science Workbench to hang.
- The Experiments table will allow you to display only three metrics at a time. You can select which metrics are displayed from the metrics dropdown. If you are tracking a large number of metrics (100 or more), you might notice some performance lag in the UI.
- Arguments are not supported with Scala experiments.
- The track_metrics and track_file functions are not supported with Scala experiments.
- The UI does not display a confirmation when you start an experiment or any alerts when experiments fail.

Debugging Issues with Experiments

This topic lists some common issues to watch out for during an experiment's build and execution process:

Experiment spends too long in Scheduling/Built stage

If your experiments are spending too long in any particular stage, check the resource consumption statistics for the cluster. When the cluster starts to run out of resources, often experiments (and other entities like jobs, models) will spend too long in the queue before they can be executed.

Resource consumption by experiments (and jobs, sessions) can be tracked by site administrators on the Admin > Activity page.

Experiment fails in the Build stage

During the build stage Cloudera Data Science Workbench creates a new Docker image for the experiment. You can track progress for this stage on each experiment's Build page. The build logs on this page should help point you in the right direction.

Common issues that might cause failures at this stage include:

- Lack of execute permissions on the build script itself.
- Inability to reach the Python package index or R mirror when installing packages.
Experiments

- Typo in the name of the build script (cdsw-build.sh). Note that the build process will only execute a script called cdsw-build.sh, not any other bash scripts from your project.
- Using pip3 to install packages in cdsw-build.sh, but selecting a Python 2 kernel when you actually launch the experiment. Or vice versa.

Experiment fails in the Execute stage

Each experiment includes a Session page where you can track the output of the experiment as it executes. This is similar to the output you would see if you test the experiment in the workbench console. Any runtime errors will display on the Session page just as they would in an interactive session.
Models

Starting with version 1.4, Cloudera Data Science Workbench allows data scientists to build, deploy, and manage models as REST APIs to serve predictions.

**Demo:** Watch the following video for a quick demonstration of the steps described in this topic: [Model Deployment with Cloudera Data Science Workbench](#).

**Related:**
- [Engines for Experiments and Models](#) on page 171
- [Model Training and Deployment Example - Iris Dataset](#)
- [Model Monitoring and Administration](#) on page 160
- [Debugging Issues with Models](#) on page 162

**Purpose**

**Challenge**

Data scientists often develop models using a variety of Python/R open source packages. The challenge lies in actually exposing those models to stakeholders who can test the model. In most organizations, the model deployment process will require assistance from a separate DevOps team who likely have their own policies about deploying new code.

For example, a model that has been developed in Python by data scientists might be rebuilt in another language by the devops team before it is actually deployed. This process can be slow and error-prone. It can take months to deploy new models, if at all. This also introduces compliance risks when you take into account the fact that the new re-developed model might not be even be an accurate reproduction of the original model.

Once a model has been deployed, you then need to ensure that the devops team has a way to rollback the model to a previous version if needed. This means the data science team also needs a reliable way to retain history of the models they build and ensure that they can rebuild a specific version if needed. At any time, data scientists (or any other stakeholders) must have a way to accurately identify which version of a model is/was deployed.

**Solution**

Starting with version 1.4, Cloudera Data Science Workbench allows data scientists to build and deploy their own models as REST APIs. Data scientists can now select a Python or R function within a project file, and Cloudera Data Science Workbench will:

- Create a snapshot of model code, model parameters, and dependencies.
- Package a trained model into an immutable artifact and provide basic serving code.
- Add a REST endpoint that automatically accepts input parameters matching the function, and that returns a data structure that matches the function’s return type.
- Save the model along with some metadata.
- Deploy a specified number of model API replicas, automatically load balanced.

**Concepts and Terminology**

**Model**

*Model* is a high level abstract term that is used to describe several possible incarnations of objects created during the model deployment process. For the purpose of this discussion you should note that 'model' does not always refer to a specific artifact. More precise terms (as defined later in this section) should be used whenever possible.

**Stages of the Model Deployment Process**
The rest of this section contains supplemental information that describes the model deployment process in detail.

Create

- **File** - The R or Python file containing the function to be invoked when the model is started.
- **Function** - The function to be invoked inside the file. This function should take a single JSON-encoded object (for example, a python dictionary) as input and return a JSON-encodable object as output to ensure compatibility with any application accessing the model using the API. JSON decoding and encoding for model input/output is built into Cloudera Data Science Workbench.

The function will likely include the following components:

- **Model Implementation**
  The code for implementing the model (e.g. decision trees, k-means). This might originate with the data scientist or might be provided by the engineering team. This code implements the model's predict function, along with any setup and teardown that may be required.

- **Model Parameters**
  A set of parameters obtained as a result of model training/fitting (using experiments). For example, a specific decision tree or the specific centroids of a k-means clustering, to be used to make a prediction.

Build

This stage takes as input the file that calls the function and returns an artifact that implements a single concrete model, referred to as a model **build**.

- **Built Model**
  A built model is a static, immutable artifact that includes the model implementation, its parameters, any runtime dependencies, and its metadata. If any of these components need to be changed, for example, code changes to the implementation or its parameters need to be retrained, a new build must be created for the model. Model builds are versioned using **build numbers**.

  To create the model build, Cloudera Data Science Workbench creates a Docker image based on the engine designated as the project's default engine. This image provides an isolated environment where the model implementation code will run.

  To configure the image environment, you can specify a list of dependencies to be installed in a build script called **cdsw-build.sh**.

  For details about the build process and examples on how to install dependencies, see *Engines for Experiments and Models* on page 171.

- **Build Number**
  Build numbers are used to track different versions of builds within the scope of a single model. They start at 1 and are incremented with each new build created for the model.
Deploy

This stage takes as input the memory/CPU resources required to power the model, the number of replicas needed, and deploys the model build created in the previous stage to a REST API.

- **Deployed Model**
  A deployed model is a model build in execution. A built model is deployed in a model serving environment, likely with multiple replicas.

- **Environmental Variable**
  You can set environmental variables each time you deploy a model. Note that models also inherit any environment variables set at the project and global level. However, in case of any conflicts, variables set per-model will take precedence.

  **Note:** If you are using any model-specific environmental variables, these must be specified every time you re-deploy a model. Models do not inherit environmental variables from previous deployments.

- **Model Replicas**
  The engines that serve incoming requests to the model. Note that each replica can only process one request at a time. Multiple replicas are essential for load-balancing, fault tolerance, and serving concurrent requests. Cloudera Data science Workbench allows you to deploy a maximum of 9 replicas per model.

- **Deployment ID**
  Deployment IDs are numeric IDs used to track models deployed across Cloudera Data Science Workbench. They are not bound to a model or project.

Creating and Deploying a Model (QuickStart)

Using Cloudera Data Science Workbench, you can create any function within a script and deploy it to a REST API. In a machine learning project, this will typically be a predict function that will accept an input and return a prediction based on the model’s parameters.

For the purpose of this quick start demo we are going to create a very simple function that adds two numbers and deploy it as a model that returns the sum of the numbers. This function will accept two numbers in JSON format as input and return the sum.

1. Create a new project. Note that models are always created within the context of a project.
2. Click Open Workbench and launch a new Python 3 session.
3. Create a new file within the project called add_numbers.py. This is the file where we define the function that will be called when the model is run. For example:

```python
def add(args):
    result = args['a'] + args['b']
    return result
```

  **Note:** In practice, do not assume that users calling the model will provide input in the correct format or enter good values. Always perform input validation.
4. Before deploying the model, test it by running the `add_numbers.py` script, and then calling the `add` function directly from the interactive workbench session. For example:

```
add({"a": 3, "b": 5})
```

5. Deploy the `add` function to a REST endpoint.
   a. Go to the project Overview page.
   b. Click Models > New Model.
   c. Give the model a Name and Description.
   d. Enter details about the model that you want to build. In this case:
      - File: `add_numbers.py`
      - Function: `add`
      - Example Input: `{"a": 3, "b": 5}`
      - Example Output: 8
   e. Select the resources needed to run this model, including any replicas for load balancing.
   f. Click Deploy Model.

6. Click on the model to go to its Overview page. Click Builds to track realtime progress as the model is built and deployed. This process essentially creates a Docker container where the model will live and serve requests.
7. Once the model has been deployed, go back to the model Overview page and use the Test Model widget to make sure the model works as expected.

If you entered example input when creating the model, the Input field will be pre-populated with those values. Click Test. The result returned includes the output response from the model, as well as the ID of the replica that served the request.

Model response times depend largely on your model code. That is, how long it takes the model function to perform the computation needed to return a prediction. It is worth noting that model replicas can only process one request at a time. Concurrent requests will be queued until the model can process them.

Calling a Model

This section lists some requirements for model requests and how to test a model using Cloudera Data Science Workbench.

- **(Requirement) JSON for Model Requests/Responses**
- **(Requirement) Model Access Key**
- **Test Calls to a Model**

**(Requirement) JSON for Model Requests/Responses**

Every model function in Cloudera Data Science Workbench takes a single argument in the form of a JSON-encoded object, and returns another JSON-encoded object as output. This format ensures compatibility with any application accessing the model using the API, and gives you the flexibility to define how JSON data types map to your model’s datatypes.

**Model Requests**

When making calls to a model, keep in mind that JSON is not suitable for very large requests and has high overhead for binary objects such as images or video. Consider calling the model with a reference to the image or video such as a URL instead of the object itself. Requests to models should not be more than 5 MB in size. Performance may degrade and memory usage increase for larger requests.

**Note:** In Cloudera Data Science Workbench 1.4.0, model request sizes were limited to 100 KB. With version 1.4.2 (and higher), this limit has been increased to 5 MB. To take advantage of this higher threshold, you will need to upgrade to version 1.4.2 (or higher) and rebuild your existing models.

Ensure that the JSON request represents all objects in the request or response of a model call. For example, JSON does not natively support dates. In such cases consider passing dates as strings, for example in ISO-8601 format, instead.
For a simple example of how to pass JSON arguments to the model function and make calls to deployed model, see [Creating and Deploying a Model (QuickStart)](page 149).

**Model Responses**

Models return responses in the form of a JSON-encoded object. Model response times depend on how long it takes the model function to perform the computation needed to return a prediction. Model replicas can only process one request at a time. Concurrent requests are queued until a replica is available to process them.

When Cloudera Data Science Workbench receives a call request for a model, it attempts to find a free replica that can answer the call. If the first arbitrarily selected replica is busy, Cloudera Data Science Workbench will keep trying to contact a free replica for 30 seconds. If no replica is available, Cloudera Data Science Workbench will return a `model.busy` error with HTTP status code 429 (Too Many Requests). If you see such errors, re-deploy the model build with a higher number of replicas.

**(Requirement) Access Key**

Each model in Cloudera Data Science Workbench has a unique access key associated with it. This access key serves two purposes: 1) it is a unique identifier for the model, and 2) it serves as an authentication token that allows you to make calls to the model.

Models deployed using Cloudera Data Science Workbench are not public. In order to call an active model your request must include the model's access key for authentication (as demonstrated in the sample calls above).

To locate the access key for a model, go to the model Overview page and click Settings.

Important:

Only one access key per model is active at any time. If you regenerate the access key, you will need to re-distribute this access key to users/applications using the model.

Alternatively, you can use this mechanism to revoke access to a model by regenerating the access key. Anyone with an older version of the key will not be able to make calls to the model.

**Testing Calls to a Model**

Cloudera Data Science Workbench provides two ways to test calls to a model:

- **Test Model Widget**

  On each model's Overview page, Cloudera Data Science Workbench provides a widget that makes a sample call to the deployed model to ensure it is receiving input and returning results as expected.
Sample Request Strings

On the model Overview page, Cloudera Data Science Workbench also provides sample curl and POST request strings that you can use to test calls to the model. Copy/paste the curl request directly into a Terminal to test the call.

Note that these sample requests already include the example input values you entered while building the model, and the access key required to query the model.

Add Two Numbers

Overview - Deployments - Builds - Monitoring - Settings

Description

Add two numbers.

Sample Code

Shell  Python  R

curl -H 'Content-Type: application/json' -X POST http://\models\acl\\models\call\model -d '{"accessKey":"mfgflyfjc5esdy8t6jyold9cfzw\","request":{"a":1,"b":2}}'

Updating Active Models

Active Model - A model that is in the Deploying, Deployed, or Stopping stages.

You can make changes to a model even after it has been deployed and is actively serving requests. Depending on business factors and changing resource requirements, such changes will likely range from changes to the model code itself, to simply modifying the number of CPU/GPUs requested for the model. In addition, you can also stop and restart active models.

Depending on your requirement, you can perform one of the following actions:
Re-deploy an Existing Build

Re-deploying a model involves re-publishing a previously-deployed model in a new serving environment - this is, with an updated number of replicas or memory/CPU/GPU allocation. For example, circumstances that require a re-deployment might include:

- An active model that previously requested a large number of CPUs/GPUs that are not being used efficiently.
- An active model that is dropping requests because it is falling short of replicas.
- An active model needs to be rolled back to one of its previous versions.

**Warning:** Currently, Cloudera Data Science Workbench only allows one active deployment per model. This means when you re-deploy a build, the current active deployment will go offline until the re-deployment process is complete and the new deployment is ready to receive requests. Prepare for model downtime accordingly.

To re-deploy an existing model:

1. Go to the model **Overview** page.
2. Click **Deployments**.
3. Select the version you want to deploy and click **Re-deploy this Build**.

**Note:** If you are using any model-specific environmental variables, these must be specified every time you re-deploy a model. Models do not inherit environmental variables from previous deployments.

4. Modify the model serving environment as needed.
5. Click **Deploy Model**.

Deploy a New Build for a Model

Deploying a new build for a model involves both, re-building the Docker image for the model, and deploying this new build. Note that this is not required if you only need to update the resources allocated to the model. As an example, changes that require a new build might include:

- Code changes to the model implementation.
- Renaming the function that is used to invoke the model.

**Warning:** Currently, Cloudera Data Science Workbench does not allow you to create a new build for a model without also deploying it. This combined with the fact that you can only have one active deployment per model means that once the new model is built, the current active deployment will go offline so that the new build can be deployed. Prepare for model downtime accordingly.

To create a new build and deploy it:

1. Go to the model **Overview** page.
2. Click **Deploy New Build**.
3. Complete the form and click **Deploy Model**.

**Stop a Model**

To stop a model (all replicas), go to the model **Overview** page and click **Stop**. Click **OK** to confirm.

**Restart a Model**

To restart a model (all replicas), go to the model **Overview** page and click **Restart**. Click **OK** to confirm.

Restarting a model does not let you make any code changes to the model. It should primarily be used as a way to quickly re-initialize or re-connect to resources.

**Usage Guidelines**

This section calls out some important guidelines you should keep in mind when you start deploying models with Cloudera Data Science Workbench.

**Model Code**

Models in Cloudera Data Science Workbench are designed to execute any code that is wrapped into a function. This means you can potentially deploy a model that returns the result of a `SELECT *` query on a very large table. However, Cloudera strongly recommends against using the models feature for such use cases.

As a best practice, your models should be returning simple JSON responses in near-real time speeds (within a fraction of a second). If you have a long-running operation that requires extensive computing and takes more than 15 seconds to complete, consider using batch jobs instead.

**Model Artifacts**

Once you start building larger models, make sure you are storing these model artifacts in HDFS, S3, or any other external storage. Do not use the project filesystem to store large output artifacts.

In general, any project files larger than 50 MB must be part of your project’s `.gitignore` file so that they are not included in snapshots for future experiments/model builds. Note that in case your models require resources that are stored outside the model itself, it is up to you to ensure that these resources are available and immutable as model replicas may be restarted at any time.

**Resource Consumption and Scaling**

Models should be treated as any other long-running applications that are continuously consuming memory and computing resources. If you are unsure about your resource requirements when you first deploy the model, start with a single replica, monitor its usage, and scale as needed.

If you notice that your models are getting stuck in various stages of the deployment process, check the **monitoring** page to make sure that the cluster has sufficient resources to complete the deployment operation.

**Security Considerations**

As stated previously, models do not impose any limitations on the code they can execute. Additionally, models run with the permissions of the user that creates the model (same as sessions and jobs). Therefore, be conscious of potential data leaks especially when querying underlying data sets to serve predictions.
Cloudera Data Science Workbench models are not public by default. Each model has an access key associated with it. Only users/applications who have this key can make calls to the model. Be careful with who has permission to view this key.

Cloudera Data Science Workbench also prints stderr/stdout logs from models to an output pane in the UI. Make sure you are not writing any sensitive information to these logs.

**Deployment Considerations**

Cloudera Data Science Workbench does not currently support high availability for models. Additionally, there can only be one active deployment per model at any given time. This means you should plan for model downtime if you want to deploy a new build of the model or re-deploy with more/less replicas.

Keep in mind that models that have been developed and trained using Cloudera Data Science Workbench are essentially Python/R code that can easily be persisted and exported to external environments using popular serialization formats such as Pickle, PMML, ONNX, and so on.

**Known Issues and Limitations**

- **Known Issues with Model Builds and Deployed Models**
  - Re-deploying or re-building models results in model downtime (usually brief).
  - Re-starting Cloudera Data Science Workbench does not automatically restart active models. These models must be manually restarted so they can serve requests again.
    Cloudera Bug: DSE-4950
  - Model deployment will fail if your project filesystem is too large for the Git snapshot process. As a general rule, any project files (code, generated model artifacts, dependencies, etc.) larger than 50 MB must be part of your project’s .gitignore file so that they are not included in snapshots for model builds.
  - Model builds will fail if your project filesystem includes a .git directory (likely hidden or nested). Typical build stage errors include:

    ```
    Error: 2 UNKNOWN: Unable to schedule build: [Unable to create a checkpoint of current source: [Unable to push sources to git server: ...
    ```
    To work around this, rename the .git directory (for example, NO.git) and re-build the model.
    Cloudera Bug: DSE-4657
  - JSON requests made to active models should not be more than 5 MB in size. This is because JSON is not suitable for very large requests and has high overhead for binary objects such as images or video. Call the model with a reference to the image or video, such as a URL, instead of the object itself.
  - Any external connections, for example, a database connection or a Spark context, must be managed by the model's code. Models that require such connections are responsible for their own setup, teardown, and refresh.
  - Model logs and statistics are only preserved so long as the individual replica is active. Cloudera Data Science Workbench may restart a replica at any time it is deemed necessary (such as bad input to the model).
  - **(Affects version 1.4.x, 1.5.x)** The model deployment example (predict.py) in the in-built Python template project does not work anymore due to a change in dependencies in the sklearn package. A working replacement for the predict.py file has been provided here: Deploy the Model - Iris Dataset.
    Cloudera Bug: DSE-5314

- **Limitations**
  - Scala models are not supported.
- **Spawning worker threads** is not supported with models.
- Models deployed using Cloudera Data Science Workbench are not highly-available.
- Dynamic scaling and auto-scaling are not currently supported. To change the number of replicas in service, you will have to re-deploy the build.

**Model Training and Deployment - Iris Dataset**

This topic uses Cloudera Data Science Workbench’s built-in Python template project to walk you through an end-to-end example where we use experiments to develop and train a model, and then deploy it using Cloudera Data Science Workbench.

This example uses the canonical Iris dataset from Fisher and Anderson to build a model that predicts the width of a flower’s petal based on the petal’s length.

**Create a Project**

The scripts for this example are available in the Python template project that ships with Cloudera Data Science Workbench. First, create a new project from the Python template:

![Create a New Project](image)

Once you’ve created the project, go to the project’s Files page. The following files are used for the demo:

- **cdsw-build.sh** - A custom build script used for models and experiments. Pip installs our dependencies, primarily the scikit-learn library.
- **fit.py** - A model training example to be run as an experiment. Generates the `model.pkl` file that contains the fitted parameters of our model.
- **predict.py** - A sample function to be deployed as a model. Uses `model.pkl` produced by `fit.py` to make predictions about petal width.

**Note:** Due to a change in dependencies in the sklearn package, the `predict.py` example does not work anymore. A complete working replacement has been provided as part of the steps in this topic.
Train the Model

Run experiments using fit.py to develop a model. The fit.py script tracks metrics, mean squared error (MSE) and $R^2$, to help compare the results of different experiments. It also writes the fitted model to a model.pkl file.

To run an experiment:

1. Navigate to the Iris project's Overview > Experiments page.
2. Click Run Experiment.
3. Fill out the form as follows and click Start Run. Make sure you use the Python 3 kernel.

4. The new experiment should now show up on the Experiments table. Click on the Run ID to go to the experiment's Overview page. The Build and Session tabs display realtime progress as the experiment builds and executes.

5. Once the experiment has completed successfully, go back to its Overview page. The tracked metrics show us that our test set had an MSE of ~0.0078 and an $R^2$ of ~0.0493. For the purpose of this demo, let's consider this an accurate enough model to deploy and use for predictions.
6. Once you have finished training and comparing metrics from different experiments, go to the experiment that generated the best model. From the experiment's Overview page, select the model.pkl file and click Add to Project.

This saves the model to the project filesystem, available on the project's Files page. We will now deploy this model as a REST API that can serve predictions.

**Deploy the Model**

To deploy the model we use the following predict.py script. This script contains the predict function that accepts petal length as input and uses the model built in the previous step to predict petal width.

```
# Read the fitted model from the file model.pkl
# and define a function that uses the model to
# predict petal width from petal length
import pickle
import numpy as np

model = pickle.load(open('model.pkl', 'rb'))

def predict(args):
    iris_x = np.reshape(float(args.get('petal_length')), (-1,1))
    result = model.predict(iris_x)
    return result[0][0]
```

1. Due to a change in dependencies in the sklearn package, the predict.py example included with the Python template project does not work anymore.

   Copy and paste the code sample provided above to replace the template project's pre-existing predict.py file.

2. Navigate to the Iris project's Overview > Models page.

3. Click New Model and fill out the fields. Make sure you use the Python 3 kernel. For example:
4. Deploy the model.
5. Click on the model to go to its **Overview** page. As the model builds you can track progress on the **Build** page. Once deployed, you can see the replicas deployed on the **Monitoring** page.

6. To test the model, use the **Test Model** widget on the model’s **Overview** page.

---

### Test Model

**Input**

```json
{
   "petal_length": 5.4
}
```

---

**Result**

<table>
<thead>
<tr>
<th>Status</th>
<th>success</th>
</tr>
</thead>
<tbody>
<tr>
<td>Response</td>
<td>1.8826221434150965</td>
</tr>
<tr>
<td>Replica ID</td>
<td>predict-petal-width-2-9-7cf557b957-5w</td>
</tr>
</tbody>
</table>
Monitoring Individual Models

When a model is deployed, Cloudera Data Science Workbench allows you to specify a number of replicas that will be deployed to serve requests. For each active model, you can monitor its replicas by going to the model's Monitoring page. On this page you can track the number of requests being served by each replica, success and failure rates, and their associated stderr and stdout logs. Depending on future resource requirements, you can increase or decrease the number of replicas by re-deploying the model.

The most recent logs are at the top of the pane (see image). stderr logs are displayed next to a red bar while stdout logs are by a green bar. Note that model logs and statistics are only preserved so long as the individual replica is active. When a replica restarts (for example, in case of bad input) the logs also start with a clean slate.

Monitoring All Active Models

Required Role: Site Administrator

To see a complete list of all the models that have been deployed on a deployment, and review resource usage across the deployment by models alone, go to Admin > Models. On this page, site administrators can also Stop/Reraest/Rebuild any of the currently deployed models.

Deleting a Model

Important:
- You must stop all active deployments before you delete a model. If not stopped, active models will continue serving requests and consuming resources even though they do not show up in Cloudera Data Science Workbench UI.
- Deleted models are not actually removed from disk. That is, this operation will not free up storage space.

Deleting a model removes all of the model's builds and its deployment history from Cloudera Data Science Workbench. To delete a model, go to the model Overview > Settings and click Delete Model.
You can also delete specific builds from a model's history by going to the model's Overview > Build page.

Disabling the Models Feature

Required Role: Site Administrator

**Important:** The feature flag mentioned here only hides the Models feature from the UI. It will not stop any active models that have already been deployed. Make sure you stop all active models from the Admin > Models page before you disable the feature.

To disable this feature on your Cloudera Data Science Workbench deployment:

1. Log in to Cloudera Data Science Workbench.
2. Click Admin > Settings.
3. Under the Feature Flags section, disable the Enable users to create models checkbox.

Debugging Issues with Models

This topic describes some common issues to watch out for during different stages of the model build and deployment process.

As a general rule, if your model spends too long in any of the afore-mentioned stages, check the resource consumption statistics for the cluster. When the cluster starts to run out of resources, often models will spend some time in a queue before they can be executed.

Resource consumption by active models on a deployment can be tracked by site administrators on the Admin > Models page.

**Building**

Live progress for this stage can be tracked on the model's Build tab. It shows the details of the build process that creates a new Docker image for the model. Potential issues:

- If you specified a custom build script (cdsw-build.sh), ensure that the commands inside the script complete successfully.
- If you are in an environment with restricted network connectivity, you might need to manually upload dependencies to your project and install them from local files.

**Pushing**

Once the model has been built, it is copied to an internal Docker registry to make it available to all the Cloudera Data Science Workbench hosts. Depending on network speeds, your model may spend some time in this stage.

**Deploying**

If you see issues occurring when Cloudera Data Science Workbench is attempting to start the model, use the following guidelines to begin troubleshooting:

- Make sure your model code works in a workbench session. To do this, launch a new session, run your model file, and then interactively call your target function with the input object. For a simple example, see the Model Quickstart.
- Ensure that you do not have any syntax errors. For python, make sure you have the kernel with the appropriate python version (Python 2 or Python 3) selected for the syntax you have used.
- Make sure that your cdsw-build.sh file provides a complete set of dependencies. Dependencies manually installed during a session on the workbench are not carried over to your model. This is to ensure a clean, isolated, build for each model.
• If your model accesses resources such as data on the CDH cluster or an external database make sure that those resources can accept the load your model may exert on them.

Deployed

Once a model is up and running, you can track some basic logs and statistics on the model’s Monitoring page. In case issues arise:

• Check that you are handling bad input from users. If your function throws an exception, Cloudera Data Science Workbench will restart your model to attempt to get back to a known good state. The user will see an unexpected model shutdown error.

For most transient issues, model replicas will respond by restarting on their own before they actually crash. This auto-restart behavior should help keep the model online as you attempt to debug runtime issues.

• Make runtime troubleshooting easier by printing errors and output to stderr and stdout. You can catch these on each model’s Monitoring tab. Be careful not to log sensitive data here.

• The Monitoring tab also displays the status of each replica and will show if the replica cannot be scheduled due to a lack of cluster resources. It will also display how many requests have been served/dropped by each replica.

When Cloudera Data Science Workbench receives a call request for a model, it attempts to find a free replica that can answer the call. If the first arbitrarily selected replica is busy, Cloudera Data Science Workbench will keep trying to contact a free replica for 30 seconds. If no replica is available, Cloudera Data Science Workbench will return a model.busy error with HTTP status code 429 (Too Many Requests). If you see such errors, re-deploy the model build with a higher number of replicas.
Managing Jobs and Pipelines in Cloudera Data Science Workbench

Cloudera Data Science Workbench allows you to automate analytics workloads with a built-in job and pipeline scheduling system that supports real-time monitoring, job history, and email alerts. A job automates the action of launching an engine, running a script, and tracking the results, all in one batch process. Jobs are created within the purview of a single project and can be configured to run on a recurring schedule. You can customize the engine environment for a job, set up email alerts for successful or failed job runs, and email the output of the job to yourself or a colleague.

As data science projects mature beyond ad hoc scripts, you might want to break them up into multiple steps. For example, a project may include one or more data acquisition, data cleansing, and finally, data analytics steps. For such projects, Cloudera Data Science Workbench allows you to schedule multiple jobs to run one after another in what is called a pipeline, where each job is dependent on the output of the one preceding it.

Creating a Job

Jobs are created within the scope of a project. When you create a job, you will be asked to select a script to execute as part of the job, and create a schedule for when the job should run. Optionally, you can configure a job to be dependent on another existing job, thus creating a pipeline of tasks to be accomplished in a sequence. Note that the script files and any other job dependencies must exist within the scope of the same project.

1. Navigate to the project for which you want to create a job.
2. On the left-hand sidebar, click Jobs.
3. Click New Job.
4. Enter a Name for the job.
5. Select a script to execute for this job by clicking on the folder icon. You will be able to select a script from a list of files that are already part of the project. To upload more files to the project, see Managing Files on page 109.
6. Depending on the code you are running, select an Engine Kernel for the job from one of the following options: Python 2, Python 3, R, or Scala.
7. Select a Schedule for the job runs from one of the following options.
   - Manual - Select this option if you plan to run the job manually each time.
   - Recurring - Select this option if you want the job to run in a recurring pattern every X minutes, or on an hourly, daily, weekly or monthly schedule.
   - Dependent - Use this option when you are building a pipeline of jobs to run in a predefined sequence. From a dropdown list of existing jobs in this project, select the job that this one should depend on. Once you have configured a dependency, this job will run only after the preceding job in the pipeline has completed a successful run.
8. Select an Engine Profile to specify the number of cores and memory available for each session.
9. Enter an optional timeout value in minutes.
10. Click Set environment variables if you want to set any values to override the overall project environment variables.
11. Specify a list of Job Report Recipients to whom you can send email notifications with detailed job reports for job success, failure, or timeout. You can send these reports to yourself, your team (if the project was created under a team account), or any other external email addresses.
12. Add any Attachments such as the console log to the job reports that will be emailed.
13. Click Create Job.

Starting with version 1.1.x, you can use the Jobs API to schedule jobs from third party workflow tools. For details, see Cloudera Data Science Workbench Jobs API on page 165.
Creating a Pipeline

The **Jobs** overview presents a list of all existing jobs created for a project along with a dependency graph to display any pipelines you’ve created. Job dependencies do not need to be configured at the time of job creation. Pipelines can be created after the fact by modifying the jobs to establish dependencies between them. From the job overview, you can modify the settings of a job, access the history of all job runs, and view the session output for individual job runs.

Let’s take an example of a project that has two jobs, Read Weblogs and Write Weblogs. Given that you must read the data before you can run analyses and write to it, the Write Weblogs job should only be triggered after the Read Weblogs job completes a successful run. To create such a two-step pipeline:

1. Navigate to the project where the Read Weblogs and Write Weblogs jobs were created.
2. Click **Jobs**.
3. From the list of jobs, select Write Weblogs.
4. Click the **Settings** tab.
5. Click on the Schedule dropdown and select **Dependent**. Select Read Weblogs from the dropdown list of existing jobs in the project.
6. Click **Update Job**.

Viewing Job History

1. Navigate to the project where the job was created.
2. Click **Jobs**.
3. Select the relevant job.
4. Click the **History** tab. You will see a list of all the job runs with some basic information such as who created the job, run duration, and status. Click individual runs to see the session output for each run.

Cloudera Data Science Workbench Jobs API

Cloudera Data Science Workbench exposes a REST API that allows you to schedule jobs from third-party workflow tools. You must authenticate yourself before you can use the API to submit a job run request. The Jobs API supports HTTP Basic Authentication, accepting the same users and credentials as Cloudera Data Science Workbench.

API Key Authentication

Cloudera recommends using your API key for requests instead of your actual username/password so as to avoid storing and sending your credentials in plaintext. The API key is a randomly generated token that is unique to each user. It must be treated as highly sensitive information because it can be used to start jobs via the API. To look up your Cloudera Data Science Workbench API key:

1. Sign in to Cloudera Data Science Workbench.
2. From the upper right drop-down menu, switch context to your personal account.
3. Click **Settings**.
4. Select the **API Key** tab.

The following example demonstrates how to construct an HTTP request using the standard basic authentication technique. Most tools and libraries, such as Curl and Python Requests, support basic authentication and can set the required Authorization header for you. For example, with curl you can pass the API Key to the --user flag and leave the password field blank.

```
curl -v -XPOST http://cdsw.example.com/api/v1/<path_to_job> --user "<API_KEY>:
```

To access the API using a library that does not provide Basic Authentication convenience methods, set the request’s Authorization header to Basic <API_KEY_encoded_in_base64>. For example, if your API key is...
Managing Jobs and Pipelines in Cloudera Data Science Workbench

Start a Job Run Using the API

Once a job has been created and configured through the Cloudera Data Science Workbench web application, you can start a run of the job through the API. This will constitute sending a POST request to a job start URL of the form:

http://cdsw.example.com/api/v1/projects/<$USERNAME>/<$PROJECT_NAME>/jobs/<$JOB_ID>/start

To construct a request, use the following steps to derive the username, project name, and job ID from the job’s URL in the web application.

1. Log in to the Cloudera Data Science Workbench web application.
2. Switch context to the team/personal account where the parent project lives.
3. Select the project from the list.
4. From the project’s Overview, select the job you want to run. This will take you to the job Overview page. The URL for this page is of the form: http://cdsw.example.com/<$USERNAME>/<$PROJECT_NAME>/jobs/<$JOB_ID>.
5. Use the $USERNAME, $PROJECT_NAME, and $JOB_ID parameters from the job Overview URL to create the following job start URL:

http://cdsw.example.com/api/v1/projects/<$USERNAME>/<$PROJECT_NAME>/jobs/<$JOB_ID>/start

For example, if your job Overview page has the URL http://cdsw.example.com/alice/sample-project/jobs/123, then a sample POST request would be of the form:

```
curl -v -XPOST http://cdsw.example.com/api/v1/projects/alice/sample-project/jobs/123/start
  --user "uysgxtj7jzkps96njextnxxmq05usp0b:" --header "Content-type: application/json"
```

Note that the request must have the Content-Type header set to application/json, even if the request body is empty.

Setting Environment Variables

You can set environment variables for a job run by passing parameters in the API request body in a JSON-encoded object with the following format.

```
{
  "environment": {
    "ENV_VARIABLE": "value 1",
    "ANOTHER_ENV_VARIABLE": "value 2"
  }
}
```

The values set here will override the defaults set for the project and the job in the web application. This request body is optional and can be left blank.

Be aware of potential conflicts with existing defaults for environment variables that are crucial to your job, such as PATH and the CDSW_* variables.

Sample Job Run

As an example, let’s assume user Alice has created a project titled Risk Analysis. Under the Risk Analysis project, Alice has created a job with the ID, 208. Using curl, Alice can use her API Key (uysgxtj7jzkps96njextnxxmq05usp0b) to create an API request as follows:

```
curl -v -XPOST http://cdsw.example.com/api/v1/projects/alice/risk-analysis/jobs/208/start
  --user "uysgxtj7jzkps96njextnxxmq05usp0b:" --header "Content-type: application/json"
  --data '{"environment": {"START_DATE": "2017-01-01", "END_DATE": "2017-01-31"}}'
```

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In this example, `START_DATE` and `END_DATE` are environment variables that are passed as parameters to the API request in a JSON object.

In the resulting HTTP request, `curl` automatically encodes the `Authorization` request header in base64 format.

You can confirm that the job was started by going to the Cloudera Data Science Workbench web application.

**Starting a Job Run Using Python**

To start a job run using Python, Cloudera recommends using `Requests`, an HTTP library for Python; it comes with a convenient API that makes it easy to submit job run requests to Cloudera Data Science Workbench. Extending the Risk Analysis example from the previous section, the following sample Python code will create an HTTP request to run the job with the job ID, 208.

**Python 2**

```python
# example.py
import requests
import json

HOST = "http://cdsw.example.com"
USERNAME = "alice"
API_KEY = "uysgxtj7jzkps96njextnxxmq05usp0b"
PROJECT_NAME = "risk-analysis"
JOB_ID = "208"

url = "/".join([HOST, "api/v1/projects" ,USERNAME, PROJECT_NAME, "jobs", JOB_ID, "start"])
job_params = {"START_DATE": "2017-01-01", "END_DATE": "2017-01-31"}
res = requests.post(url, headers = {"Content-Type": "application/json"}, auth = (API_KEY,""),
                     data = json.dumps({"environment": job_params})

print "URL", url
print "HTTP status code", res.status_code
print "Engine ID", res.json().get('engine_id')
```

When you run the code, you should see output of the form:

```
python example.py
URL http://cdsw.example.com/api/v1/projects/alice/risk-analysis/jobs/208/start
HTTP status code 200
Engine ID r11w5q3q589ryg9o
```
Limitations

- Cloudera Data Science Workbench does not support changing your API key, or having multiple API keys.
- Currently, you cannot create a job, stop a job, or get the status of a job using the Jobs API.
Cloudera Data Science Workbench Engines

In the context of Cloudera Data Science Workbench, engines are responsible for running data science workloads and intermediating access to the underlying CDH cluster. This topic gives an overview of engines and walks you through some of the ways you can customize engine environments to meet the requirements of your users and projects.

Basic Concepts and Terminology

Base Engine Image

The base engine image is a Docker image that contains all the building blocks needed to launch a Cloudera Data Science Workbench session and run a workload. It consists of kernels for Python, R, and Scala along with additional libraries that can be used to run common data analytics operations. When you launch a session to run a project, an engine is kicked off from a container of this image. The base image itself is built and shipped along with Cloudera Data Science Workbench.

New versions of the base engine image are released periodically. However, existing projects are not automatically upgraded to use new engine images. Older images are retained to ensure you are able to test code compatibility with the new engine before upgrading to it manually.

For more details on the libraries shipped within the base engine image, see Cloudera Data Science Workbench Engine Versions and Packaging on page 187.

Engine

The term engine refers to a virtual machine-style environment that is created when you run a project (via session or job) in Cloudera Data Science Workbench. You can use an engine to run R, Python, and Scala workloads on data stored in the underlying CDH cluster.

Cloudera Data Science Workbench allows you to run code using either a session or a job. A session is a way to interactively launch an engine and execute code while a job lets you batch process those actions and schedule them to run recursively. Each session and job launches its own engine that lives as long as the workload is running (or until it times out).

A running engine includes the following components:

- **Kernel**
  
  Each engine runs a kernel with an R, Python or Scala process that can be used to execute code within the engine. The kernel launched differs based on the option you select (either Python 2/3, PySpark, R, or Scala) when you launch the session or configure a job.

  The Python kernel is based on the Jupyter IPython kernel; the R kernel is custom-made for CDSW; and the Scala kernel is based on the Apache Toree kernel.

- **Project Filesystem Mount**
Cloudera Data Science Workbench Engines

Cloudera Data Science Workbench uses a persistent filesystem to store project files such as user code, installed libraries, or even small data files. Project files are stored on the master host at /var/lib/cdsw/current/projects.

Every time you launch a new session or run a job for a project, a new engine is created, and the project filesystem is mounted into the engine’s environment at /home/cdsw. Once the session/job ends, the only project artifacts that remain are a log of the workload you ran, and any files that were generated or modified, including libraries you might have installed. All of the installed dependencies persist through the lifetime of the project. The next time you launch a session/job for the same project, those dependencies will be mounted into the engine environment along with the rest of the project filesystem.

- **CDH and Host Mounts**

  To ensure that each engine is able to access the CDH cluster, a number of folders are mounted from the CDSW gateway host into the engine’s environment. For example, on a CSD deployment, this includes the path to the parcel repository (/opt/cloudera), client configurations for HDFS, Spark, YARN, as well as the host’s JAVA_HOME.

  Cloudera Data Science Workbench works out-of-the-box for CDH clusters that use the default file system layouts configured by Cloudera Manager. If you customized your CDH cluster’s filesystem layout (for example, modified the CDH parcel directory) or if there are other files on the hosts that should be mounted into the engines, use the Site Administration panel to include them.

  For detailed instructions, see CDH Parcel Directory and Host Mounts.

**Project Environments**

This section describes how you can configure engine environments to meet the requirements of a project. This can be done by using environmental variables and by installing dependencies.

**Environmental Variables**

Environmental variables help you customize engine environments, both globally and for individual projects/jobs. For example, if you need to configure a particular timezone for a project or increase the length of the session/job timeout windows, you can use environmental variables to do so. Environmental variables can also be used to assign variable names to secrets, such as passwords or authentication tokens, to avoid including these directly in the code.

For a list of the environmental variables you can configure and instructions on how to configure them, see Engine Environment Variables on page 179.

**Dependencies**

You can provide packages, such as Python libraries, in addition to the pre-installed package through the following methods:

- Directly installing packages within projects
- Creating a custom engine with the required packages
- Mounting a path from the host which contains additional packages

One method may be more appropriate for your deployment than another method. For more information about each option, see Managing Engine Dependencies on page 176.

**Configuring Engine Environments for Experiments and Models**

To allow for versioning of experiments and models, Cloudera Data Science Workbench executes each experiment and model in a completely isolated engine. Every time a model or experiment is kicked off, Cloudera Data Science Workbench creates a new isolated Docker image where the model or experiment is executed. These engines are built by extending the project’s designated default engine image to include the code to be executed and any dependencies as specified.
For details on how this process works and how to configure these environments, see Engines for Experiments and Models on page 171.

Engines for Experiments and Models

In Cloudera Data Science Workbench, models, experiments, jobs, and sessions are all created and executed within the context of a project. We've described the different ways in which you can customize a project's engine environment for sessions and jobs here. However, engines for models and experiments are completely isolated from the rest of the project.

Every time a model or experiment is kicked off, Cloudera Data Science Workbench creates a new isolated Docker image where the model or experiment is executed. This isolation in build and execution makes it possible for Cloudera Data Science Workbench to keep track of input and output artifacts for every experiment you run. In case of models, versioned builds give you a way to retain build history for models and a reliable way to rollback to an older version of a model if needed.

The rest of this topic describes the engine build process that occurs when you kick off a model or experiment.

Snapshot Code

When you first launch an experiment or model, Cloudera Data Science Workbench takes a Git snapshot of the project filesystem at that point in time. It is important to note that this Git server functions behind the scenes and is completely separate from any other Git version control system you might be using for the project as a whole.

However, this Git snapshot will recognize the .gitignore file defined in the project. This means if there are any artifacts (files, dependencies, etc.) larger than 50 MB stored directly in your project filesystem, make sure to add those files or folders to .gitignore so that they are not recorded as part of the snapshot. This ensures that the experiment/model environment is truly isolated and does not inherit dependencies that have been previously installed in the project workspace.

By default, each project is created with the following .gitignore file:

```bash
R
  node_modules
  *.pyc
  *
  !.gitignore
```

Augment this file to include any extra dependencies you have installed in your project workspace to ensure a truly isolated workspace for each model/experiment.

Build Image

Once the code snapshot is available, Cloudera Data Science Workbench creates a new Docker image with a copy of the snapshot. This new image is based off the project's designated default engine image (configured at Project Settings > Engine). The image environment can be customized by using environmental variables and a build script that specifies which packages should be included in the new image.

Environmental Variables

Both models and experiments inherit environmental variables from their parent project. Furthermore, in case of models, you can specify environment variables for each model build. In case of conflicts, the variables specified per-build will override any values inherited from the project.

For more information, see Engine Environment Variables on page 179.
Build Script - cdsw-build.sh

As part of the Docker build process, Cloudera Data Science Workbench runs a build script called `cdsw-build.sh` file. You can use this file to customize the image environment by specifying any dependencies to be installed for the code to run successfully. One advantage to this approach is that you now have the flexibility to use different tools and libraries in each consecutive training run. Just modify the build script as per your requirements each time you need to test a new library or even different versions of a library.

Important:
- The `cdsw-build.sh` script does not exist by default — it has to be created by you within each project as needed.
- The name of the file is not customizable. It must be called `cdsw-build.sh`.

The following sections demonstrate how to specify dependencies in Python and R projects so that they are included in the build process for models and experiments.

Python 3

For Python, create a `requirements.txt` file in your project with a list of packages that must be installed. For example:

```
beautifulsoup4==4.6.0
seaborn==0.7.1
```

*Figure 1: requirements.txt*

Then, create a `cdsw-build.sh` file in your project and include the following command to install the dependencies listed in `requirements.txt`.

```
pip3 install -r requirements.txt
```

*Figure 2: cdsw-build.sh*

Now, when `cdsw-build.sh` is run as part of the build process, it will install the `beautifulsoup4` and `seaborn` packages to the new image built for the experiment/model.

R

For R, create a script called `install.R` with the list of packages that must be installed. For example:

```
install.packages(repos="https://cloud.r-project.org", c("tidyr", "stringr"))
```

*Figure 3: install.R*

Then, create a `cdsw-build.sh` file in your project and include the following command to run `install.R`.

```
Rscript install.R
```

*Figure 4: cdsw-build.sh*

Now, when `cdsw-build.sh` is run as part of the build process, it will install the `tidyr` and `stringr` packages to the new image built for the experiment/model.

If you do not specify a build script, the build process will still run to completion, but the Docker image will not have any additional dependencies installed. At the end of the build process, the built image is then pushed to an internal Docker registry so that it can be made available to all the Cloudera Data Science Workbench hosts. This push is largely transparent to the end user.
Run Experiment / Deploy Model

Once the Docker image has been built and pushed to the internal registry, the experiment/model can now be executed within this isolated environment.

In case of experiments, you can track live progress as the experiment executes in the experiment’s Session tab.

Unlike experiments, models do not display live execution progress in a console. Behind the scenes, Cloudera Data Science Workbench will move on to deploying the model in a serving environment based on the computing resources and replicas you requested. Once deployed you can go to the model’s Monitoring page to view statistics on the number of requests served/dropped and stderr/stdout logs for the model replicas.

Configuring Cloudera Data Science Workbench Engines

This topic describes how to configure and manage engines in Cloudera Data Science Workbench. Cloudera Data Science Workbench currently supports R, Python, and Scala engines. You can use these engines to run data science projects either in isolation, as you would on your laptop, or connect to your CDH cluster using Cloudera Distribution of Apache Spark 2 and other libraries.

Concepts and Terminology

Review basic concepts and terminology related to engines at Cloudera Data Science Workbench Engines on page 169.

Managing Engines

**Required Role:**Site Administrator

Site administrators and project administrators are responsible for making sure that all projects on the deployment have access to the engines they need. Site admins can create engine profiles, determine the default engine version to be used across the deployment, and white-list any custom engines that teams require. As a site administrator, you can also customize engine environments by setting global environmental variables and configuring any files/folders that need to be mounted into project environments on run time.

**Managing Engine Profiles**

Engine profiles define how many vCPUs and how much memory Cloudera Data Science Workbench will reserve for a particular session/job. As a site administrator you can create several different vCPU, GPU, and memory configurations which will be available when launching a session/job. When launching a new session, users will be able to select one of the available engine profiles depending on their project’s requirements.
To create engine profiles, go to the Admin > Engines page, under Engines Profiles. Cloudera recommends that all profiles include at least 2 GB of RAM to avoid out of memory errors for common user operations.

You will see the option to add GPUs to the engine profiles only if your Cloudera Data Science Workbench hosts are equipped with GPUs, and you have enabled them for use by setting the relevant properties either in Cloudera Manager (for CSD) or in `cdsw.conf` (for RPM).

Managing Engine Images

By default, Cloudera Data Science Workbench ships a base engine image that includes kernels for Python, R, and Scala, along with some additional libraries that can be used to run common data analytics operations. Occasionally, new engine versions are released and shipped with Cloudera Data Science Workbench releases.

Engine images are available in the Site Administrator panel at Admin > Engines, under the Engine Images section. As a site administrator, you can select which engine version is used by default for new projects. Furthermore, project administrators can explicitly select which engine image should be used as the default image for a project. To do so, go to the project's Overview page and click Settings on the left navigation bar.

If a user publishes a new custom Docker image, site administrators are responsible for white-listing such images for use across the deployment. For more information on creating and managing custom Docker images, see Customized Engine Images on page 184.

Configuring the Engine Environment

This section describes some of the ways you can configure engine environments to meet the requirements of your projects.

Environmental Variables

For information on how environmental variables can be used to configure engine environments in Cloudera Data Science Workbench, see Engine Environment Variables on page 179.

CDH Parcel Directory

Starting with Cloudera Data Science Workbench 1.5, the CDH parcel directory property is no longer available in the Site Administration panel. By default, Cloudera Data Science Workbench looks for the CDH parcel at `/opt/cloudera/parcels`.

If you want to use a custom location for your parcels, use one of the following methods to configure this custom location:

**CSD deployments:** If you are using the default parcel directory, `/opt/cloudera/parcels`, no action is required. If you want to use a custom location for the parcel directory, configure this in Cloudera Manager as documented here.

**OR**
RPM deployments: If you are using the default parcel directory, `/opt/cloudera/parcels`, no action is required. If you want to specify a custom location for the parcel directory, configure the `DISTRO_DIR` property in the `cdsw.conf` file on both master and worker hosts. Run `cdsw restart` after you make this change.

Configuring Host Mounts

By default, Cloudera Data Science Workbench will automatically mount the CDH parcel directory and client configuration for required services such as HDFS, Spark, and YARN into each project’s engine. However, if users want to reference any additional files/folders on the host, site administrators will need to configure them here so that they are loaded into engine containers at runtime. Note that the directories specified here will be available to all projects across the deployment.

To configure additional mounts, go to Admin > Engines and add the paths to be mounted from the host to the Mounts section.

The following table summarizes how mounts are loaded into engine containers in current and previous Cloudera Data Science Workbench releases.

<table>
<thead>
<tr>
<th>CDSW Version</th>
<th>Mount Point Permissions in Engines</th>
</tr>
</thead>
<tbody>
<tr>
<td>1.4.2 (and higher)</td>
<td>By default, mount points are loaded into engine containers with read-only permissions. CDSW 1.4.2 (and higher) also include a Write Access checkbox (see image) that you can use to enable read-write access for individual mounted directories. Note that these permissions will apply to all projects across the deployment.</td>
</tr>
<tr>
<td>1.4.0</td>
<td>Mount points are loaded into engine containers with read-write permissions.</td>
</tr>
<tr>
<td>1.3.x (and lower)</td>
<td>Mount points are loaded into engine containers with read-only permissions.</td>
</tr>
</tbody>
</table>

Points to Remember:

- When adding host mounts, try to be as generic as possible without mounting common system files. For example, if you want to add several files under `/etc/spark2-conf`, you can simplify and mount the `/etc/spark2-conf` directory; but adding the parent `/etc` might prevent the engine from running.

  As a general rule, do not mount full system directories that are already in use; such as `/var`, `/tmp`, or `/etc`. This also serves to avoid accidentally exposing confidential information in running sessions.

- Do not add duplicate mount points. This will lead to sessions crashing in the workbench.

Setting Time Zones for Sessions and Jobs

The default time zone for Cloudera Data Science Workbench sessions is UTC. This is the default regardless of the time zone setting on the Master host.

To change to your preferred time zone, for example, Pacific Standard Time (PST), navigate to Admin > Engines. Under the Environmental Variables section, add a new variable with the name set to `TZ` and value set to `America/Los_Angeles`, and click Add.
Managing Engine Dependencies

This page describes the options available to you for mounting a project's dependencies into its engine environment:

**Important:** Even though experiments and models are created within the scope of a project, the engines they use are completely isolated from those used by sessions or jobs launched within the same project. For details, see Engines for Experiments and Models on page 171.

Installing Packages Directly Within Projects

Cloudera Data Science Workbench engines are preloaded with a few common packages and libraries for R, Python, and Scala. In addition to these, Cloudera Data Science Workbench allows you to install any other packages or libraries required by your projects just as you would on your local computer. Each project's environment is completely isolated from others, which means you can install different versions of libraries pinned to different projects.

Libraries can be installed from the workbench using the inbuilt interactive command prompt or the terminal. Any dependencies installed this way are mounted to the project environment at /home/cdsw. Alternatively, you could choose to use a package manager such as Conda to install and maintain packages and their dependencies.

Note that overriding pre-installed packages by installing packages directly in the workbench can have unwanted side effects. It is not recommended or supported.

For detailed instructions, see Installing Additional Packages on page 181.
Creating a Customized Engine with the Required Package(s)

1. Build Custom Engine A with the dependencies.

2. Push Custom Engine A to a visible registry.

3. CDSW Site Admin whitelists Custom Engine A: Admin > Engines.


Directly installing a package to a project as described above might not always be feasible. For example, packages that require root access to be installed, or that must be installed to a path outside `/home/cdsw` (outside the project mount), cannot be installed directly from the workbench. For such circumstances, Cloudera recommends you extend the base Cloudera Data Science Workbench engine image to build a customized image with all the required packages installed to it.

This approach can also be used to accelerate project setup across the deployment. For example, if you want multiple projects on your deployment to have access to some common dependencies out of the box or if a package just has a complicated setup, it might be easier to simply provide users with an engine environment that has already been customized for their project(s).

For detailed instructions with an example, see Customized Engine Images on page 184
Mounting Additional Dependencies from the Host

1. CDSW Site Admin selects a sub-tree of the filesystem to act as the mount path for dependencies.

2. Site Admin (as a root user) installs Package X to that sub-tree on ALL CDSW Gateway hosts.

3. CDSW Site Admin adds the filesystem mount path for the dependencies: Admin > Engines.

Limitation

If you use this option, you must self-manage the dependencies installed on the gateway hosts to ensure consistency across them all. Cloudera Data Science Workbench cannot manage or control the packages with this method. For example, you must manually ensure that version mismatches do not occur. Additionally, this method can lead to issues with experiment repeatability since CDSW does not control or keep track of the packages on the host. Contrast this with Installing Packages Directly Within Projects on page 176 and Creating a Customized Engine with the Required Package(s) on page 177, where Cloudera Data Science Workbench is aware of the packages and manages them.

For instructions on how to configure host mounts, see Configuring Host Mounts on page 175. Note that the directories specified here will be available to all projects across the deployment.

Managing Dependencies for Spark 2 Projects

With Spark projects, you can add external packages to Spark executors on startup. To add external dependencies to Spark jobs, specify the libraries you want added by using the appropriate configuration parameters in a spark-defaults.conf file.
Engine Environment Variables

Environmental variables allow you to customize engine environments for projects. For example, if you need to configure a particular timezone for a project, or increase the length of the session/job timeout windows, you can use environmental variables to do so. Environmental variables can also be used to assign variable names to secrets such as passwords or authentication tokens to avoid including these directly in the code.

In general, Cloudera recommends that you do not include passwords, tokens, or any other secrets directly in your code because anyone with read access to your project will be able to view this information. A better place to store secrets is in your project's environment variables, where only project collaborators and admins have view access. They can therefore be used to securely store confidential information such as your AWS keys or database credentials.

Cloudera Data Science Workbench allows you to define environmental variables for the following scopes:

**Global**

A site administrator for your Cloudera Data Science Workbench deployment can set environmental variables on a global level. These values will apply to every project on the deployment.

To set global environmental variables, go to Admin > Engines.

**Project**

Project administrators can set project-specific environmental variables to customize the engines launched for a project. Variables set here will override the global values set in the site administration panel.

To set environmental variables for a project, go to the project's Overview page and click Settings > Engine.

**Job**

Environments for individual jobs within a project can be customized while creating the job. Variables set per-job will override the project-level and global settings.

To set environmental variables for a job, go to the job's Overview page and click Settings > Set Environmental Variables.

**Experiments**

Engines created for execution of experiments are completely isolated from the project. However, these engines inherit values from environmental variables set at the project-level and/or global level. Variables set at the project-level will override the global values set in the site administration panel.

**Models**

Model environments are completely isolated from the project. Environmental variables for these engines can be configured during the build stage of the model deployment process. Models will also inherit any environment variables set at the project and global level. However, variables set per-model build will override other settings.

Environment Variables from Cloudera Manager

In addition to the environment variables that you can specify with different scopes, Cloudera Data Science Workbench inherits a set of environment variables from Cloudera Manager:

- HTTP_PROXY
- HTTPS_PROXY
- ALL_PROXY
- NO_PROXY

For information about what these variables are used for, see Configuring Cloudera Data Science Workbench Deployments Behind a Proxy on page 235.
Site and project administrators can change these values by manually modifying them at the project or global level. The values set within Cloudera Data Science Workbench take precedence over the ones inherited from Cloudera Manager.

**Accessing Environmental Variables from Projects**

Environmental variables are injected into every engine launched for a project, contingent on the scope at which the variable was set (global, project, etc.). The following code samples show how to access a sample environment variable called `DATABASE_PASSWORD` from your project code.

**R**

```r
database.password <- Sys.getenv("DATABASE_PASSWORD")
```

**Python**

```python
import os
database_password = os.environ["DATABASE_PASSWORD"]
```

**Scala**

```scala
System.getenv("DATABASE_PASSWORD")
```

**Engine Environment Variables**

The following table lists Cloudera Data Science Workbench environment variables that you can use to customize your experience within the Workbench console. These can be set either as a site administrator, or within the scope of a project or a job.

<table>
<thead>
<tr>
<th>Environment Variable</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>MAX_TEXT_LENGTH</td>
<td>Maximum number of characters that can be displayed in a single text cell. By default, this value is set to 800,000 and any more characters will be truncated. Default: 800,000</td>
</tr>
<tr>
<td>SESSION_MAXIMUM_MINUTES</td>
<td>Maximum number of minutes a session can run before it times out. Default: 60<em>24</em>7 minutes (7 days) Maximum Value: 35,000 minutes</td>
</tr>
<tr>
<td>JOB_MAXIMUM_MINUTES</td>
<td>Maximum number of minutes a job can run before it times out. Default: 60<em>24</em>7 minutes (7 days) Maximum Value: 35,000 minutes</td>
</tr>
<tr>
<td>IDLE_MAXIMUM_MINUTES</td>
<td>Maximum number of minutes a session can remain idle before it exits. Default: 60 minutes Maximum Value: 35,000 minutes</td>
</tr>
<tr>
<td>CONDA_DEFAULT_ENV</td>
<td>Points to the default Conda environment so you can use Conda to install/manage packages in the Workbench. For more details on when to use this variable, see Using Conda with Cloudera Data Science Workbench on page 182.</td>
</tr>
</tbody>
</table>

**Per-Engine Environmental Variables**: In addition to the previous table, there are some more built-in environmental variables that are set by the Cloudera Data Science Workbench application itself and do not need to be modified by users. These variables are set `per-engine` launched by Cloudera Data Science Workbench and only apply within the scope of each engine.
<table>
<thead>
<tr>
<th>Environment Variable</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>CDSW_PROJECT</td>
<td>The project to which this engine belongs.</td>
</tr>
<tr>
<td>CDSW_ENGINE_ID</td>
<td>The ID of this engine. For sessions, this appears in your browser’s URL bar.</td>
</tr>
<tr>
<td>CDSW_MASTER_ID</td>
<td>If this engine is a worker, this is the CDSW_ENGINE_ID of its master.</td>
</tr>
<tr>
<td>CDSW_MASTER_IP</td>
<td>If this engine is a worker, this is the IP address of its master.</td>
</tr>
<tr>
<td>CDSW_PUBLIC_PORT</td>
<td>A port on which you can expose HTTP services in the engine to browsers. HTTP services that bind CDSW_PUBLIC_PORT will be available in browsers at: http(s): //&lt;CDSW_ENGINE_ID&gt;.-&lt;CDSW_DOMAIN&gt;. By default, CDSW_PUBLIC_PORT is set to 8080. A direct link to these web services will be available from the grid icon in the upper right corner of the Cloudera Data Science Workbench web application, as long as the job or session is still running. For more details, see Accessing Web User Interfaces from Cloudera Data Science Workbench on page 123.</td>
</tr>
<tr>
<td>CDSW_DOMAIN</td>
<td>The domain on which Cloudera Data Science Workbench is being served. This can be useful for iframing services, as demonstrated in the Shiny example.</td>
</tr>
<tr>
<td>CDSW_CPU_MILLCORES</td>
<td>The number of CPU cores allocated to this engine, expressed in thousandths of a core.</td>
</tr>
<tr>
<td>CDSW_MEMORY_MB</td>
<td>The number of megabytes of memory allocated to this engine.</td>
</tr>
<tr>
<td>CDSW_IP_ADDRESS</td>
<td>Other engines in the Cloudera Data Science Workbench cluster can contact this engine on this IP address.</td>
</tr>
</tbody>
</table>

### Installing Additional Packages

Cloudera Data Science Workbench engines are preloaded with a few common packages and libraries for R, Python, and Scala. However, a key feature of Cloudera Data Science Workbench is the ability of different projects to install and use libraries pinned to specific versions, just as you would on your local computer.

Generally, Cloudera recommends you install all required packages locally into your project. This will ensure you have the exact versions you want and that these libraries will not be upgraded when Cloudera upgrades the base engine image. You only need to install libraries and packages once per project. From then on, they are available to any new engine you spawn throughout the lifetime of the project.

You can install additional libraries and packages from the workbench, using either the command prompt or the terminal. Alternatively, you might choose to use a package manager such as Conda to install and maintain packages and their dependencies. For some basic usage guidelines for Conda, see Using Conda with Cloudera Data Science Workbench on page 182.

**Note:**

Cloudera Data Science Workbench does not currently support installation of packages that require root access to the hosts. For such use-cases, you will need to create a new custom engine that extends the base engine image to include the required packages. For instructions, see Creating Custom Engine Images.
To install a package from the command prompt:

1. Navigate to your project’s Overview page. Click Open Workbench and launch a session.
2. At the command prompt in the bottom right, enter the command to install the package. Some examples using Python and R have been provided.

**R**

```r
# Install from CRAN
install.packages("ggplot2")

# Install using devtools
install.packages('devtools')
library(devtools)
install_github("hadley/ggplot2")
```

**Python 2**

```python
# Installing from console using ! shell operator and pip:
!pip install beautifulsoup

# Installing from terminal
pip install beautifulsoup
```

**Python 3**

```python
# Installing from console using ! shell operator and pip3:
!pip3 install beautifulsoup4

# Installing from terminal
pip3 install beautifulsoup4
```

**(Python Only) Using a Requirements File**

For a Python project, you can specify a list of the packages you want in a requirements.txt file that lives in your project. The packages can be installed all at once using pip/pip3.

1. Create a new file called requirements.txt file within your project:

   ```
   beautifulsoup4==4.6.0
   seaborn==0.7.1
   ```

2. To install the packages in a Python 3 engine, run the following command in the workbench command prompt.

   ```
   !pip3 install -r requirements.txt
   ```

   For Python 2 engines, use pip.

   ```
   !pip install -r requirements.txt
   ```

**Using Conda with Cloudera Data Science Workbench**

Cloudera Data Science Workbench recommends using pip for package management along with a requirements.txt file (as described in the previous section). However, for users that prefer Conda, the default engine in Cloudera Data Science Workbench includes two environments called python2.7, and python3.6. These environments are added to sys.path, depending on the version of Python selected when you launch a new session.

In Python 2 and Python 3 sessions and attached terminals, Cloudera Data Science Workbench automatically sets the CONDA_DEFAULT_ENV and CONDA_PREFIX environment variables to point to Conda environments under /home/cdsw/.conda.

However, Cloudera Data Science Workbench does not automatically configure Conda to pin the actual Python version. Therefore if you are using Conda to install a package, you must specify the version of Python. For example, to use
Conda to install the `feather-format` package into the python3.6 environment, run the following command in the Workbench command prompt:

```
!conda install -y -c conda-forge python=3.6.1 feather-format
```

To install a package into the python2.7 environment, run:

```
!conda install -y -c conda-forge python=2.7.11 feather-format
```

Note that on `sys.path`, pip packages have precedence over conda packages.

Note:
- If your project is using an older base engine image (version 3 and lower), you will need to specify both the Python version as well as the Conda environment. For example:

```
!conda install -y -c conda-forge --name python3.6 python=3.6.1 feather-format
```

The Conda environment is also required when you create an extensible engine using Conda (as described in the following section).
- Cloudera Data Science Workbench does not automatically configure a Conda environment for R and Scala sessions and attached terminals. If you want to use Conda to install packages from an R or Scala session or terminal, you must manually configure Conda to install packages into the desired environment.

Creating an Extensible Engine With Conda

Cloudera Data Science Workbench also allows you to [extend its base engine image](#) to include packages of your choice using Conda. To create an extended engine:

1. Add the following lines to a Dockerfile to extend the base engine, push the engine image to your Docker registry, and whitelist the new engine for your project. For more details on this step, see [Extensible Engines](#).

   **Python 2**

   ```
   RUN mkdir -p /opt/conda/envs/python2.7
   RUN conda install -y nbconvert python=2.7.11 -n python2.7
   ```

   **Python 3**

   ```
   RUN mkdir -p /opt/conda/envs/python3.6
   RUN conda install -y nbconvert python=3.6.1 -n python3.6
   ```

2. Set the `PYTHONPATH` environmental variable as shown below. You can set this either globally in the site administrator dashboard, or for a specific project by going to the project’s Settings > Engine page.

   **Python 2**

   ```
   PYTHONPATH=$PYTHONPATH:/opt/conda/envs/python2.7/lib/python2.7/site-packages
   ```

   **Python 3**

   ```
   PYTHONPATH=$PYTHONPATH:/opt/conda/envs/python3.6/lib/python3.6/site-packages
   ```
Customized Engine Images

By default, Cloudera Data Science Workbench engines are preloaded with a few common packages and libraries for R, Python, and Scala. In addition to these, Cloudera Data Science Workbench also allows you to install any other packages or libraries that are required by your projects. However, directly installing a package to a project as described above might not always be feasible. For example, packages that require root access to be installed, or that must be installed to a path outside /home/cdsw (outside the project mount), cannot be installed directly from the workbench.

For such circumstances, Cloudera Data Science Workbench allows you to extend the base Docker image and create a new Docker image with all the libraries and packages you require. Site administrators can then whitelist this new image for use in projects, and project administrators set the new white-listed image to be used as the default engine image for their projects. For an end-to-end example of this process, see End-to-End Example: MeCab on page 185.

Note that this approach can also be used to accelerate project setup across the deployment. For example, if you want multiple projects on your deployment to have access to some common dependencies (package or software or driver) out of the box, or even if a package just has a complicated setup, it might be easier to simply provide users with an engine that has already been customized for their project(s).

Creating a Customized Engine Image

This section walks you through the steps required to create your own custom engine based on the Cloudera Data Science Workbench base image. For a complete example, see End-to-End Example: MeCab on page 185.

Create a Dockerfile for the New Custom Image

The first step when building a customized image is to create a Dockerfile that specifies which packages you would like to install in addition to the base image.

For example, the following Dockerfile installs the `beautifulsoup4` package on top of the base Ubuntu image that ships with Cloudera Data Science Workbench.

```bash
# Dockerfile
FROM docker.repository.cloudera.com/cdsw/engine:5
# Update packages on the base image and install beautifulsoup4
RUN apt-get update
RUN pip install beautifulsoup4 && pip3 install beautifulsoup4
```

Build the New Image

A new custom Docker image can be built on any host where Docker binaries are installed. To install these binaries, run the following command on the host where you want to build the new image:

```
docker build -t <image-name>:<tag> . -f Dockerfile
```

If you want to build your image on a Cloudera Data Science Workbench gateway host, you must add the `--network=host` option to the build command:

```
docker build --network=host -t <image-name>:<tag> . -f Dockerfile
```

Distribute the Image

Once you have built a new custom engine, use one of the following ways to distribute the new image to all your Cloudera Data Science Workbench hosts:

**Push the image to a public registry such as DockerHub**

For instructions, refer the Docker documentation: docker push, Push Images to Docker Cloud.

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**Push the image to your company’s Docker registry**

When using this method, make sure to tag your image with the following schema:

```bash
docker tag <image-name> <company-registry>/<user-name>/<image-name>:<tag>
```

Once the image has been tagged properly, use the following command to push the image:

```bash
docker push <company-registry>/<user-name>/<image-name>:<tag>
```

The MeCab example at the end of this topic uses this method.

**Distribute the image manually**

Use the following steps to manually distribute the image on the cluster:

1. **Save the docker image as a tarball on the host where it was built**

   ```bash
docker image save -o ./<new_customized_engine>.tar <image-name>
   `` `

2. **Distribute the image to all the Cloudera Data Science Workbench gateway hosts.**

   ```bash
   scp ./<new_customized_engine>.tar root@<cdsw.your_company.com>:/tmp/
   ```

3. **Load the image on all the Cloudera Data Science Workbench gateway hosts.**

   ```bash
docker load --input /tmp/.<new_customized_engine>.tar
   `` `

4. **To verify that the image was successfully distributed and loaded, run:**

   ```bash
docker images
   ``

**Whitelist the Image in Cloudera Data Science Workbench**

White-listing a customized image in Cloudera Data Science Workbench is a two-step process.

1. **Whitelist Image for the Deployment**

   First, a site administrator will need to clear the new image for use on the deployment.

   1. Log in as a site administrator.
   2. Click `Admin > Engines`
   3. Add `<company-registry>/<user-name>/<image-name>:<tag>` to the list of whitelisted engine images.

2. **Whitelist Image for Per-Project**

   If you want to start using the image in a project, the project administrator will need to set this image as the default image for the project.

   1. Go to the project `Settings` page.
   2. Click `Engines`
   3. Select the new customized engine from the dropdown list of available Docker images. Sessions and jobs you run in your project will now have access to this engine.

**End-to-End Example: MeCab**

This section demonstrates how to customize the Cloudera Data Science Workbench base engine image to include the MeCab (a Japanese text tokenizer) library.
This is a sample Dockerfile that adds MeCab to the Cloudera Data Science Workbench base image.

```docker
# Dockerfile
FROM docker.repository.cloudera.com/cdsw/engine:5
RUN apt-get update && \
    apt-get install -y -q mecab \ 
    libmecab-dev \ 
    mecab-ipadic-utf8 && \
    apt-get clean && \
    rm -rf /var/lib/apt/lists/*
RUN cd /tmp && \
    git clone --depth 1 https://github.com/neologd/mecab-ipadic-neologd.git && \
    /tmp/mecab-ipadic-neologd/bin/install-mecab-ipadic-neologd -y -n -p \
    /var/lib/mecab/dic/neologd && \
    rm -rf /tmp/mecab-ipadic-neologd
RUN pip install --upgrade pip
RUN pip install mecab-python==0.996
```

To use this image on your Cloudera Data Science Workbench project, perform the following steps.

1. Build a new image with the Dockerfile.
   ```bash
docker build --network=host -t <company-registry>/user/cdsw-mecab:latest . -f Dockerfile
   ```

2. Push the image to your company's Docker registry.
   ```bash
docker push <your-company-registry>/user/cdsw-mecab:latest
   ```

3. Whitelist the image, `<your-company-registry>/user/cdsw-mecab:latest`. Only a site administrator can do this.
   Go to Admin > Engines and add `<company-registry>/user/cdsw-mecab:latest` to the list of whitelisted engine images.

4. Ask a project administrator to set the new image as the default for your project. Go to the project Settings, click Engines, and select `company-registry/user/cdsw-mecab:latest` from the dropdown.

You should now be able to run this project on the customized MeCab engine.
Limitations

- Cloudera Data Science Workbench only supports customized engines that are based on the Cloudera Data Science Workbench base image.
- Cloudera Data Science Workbench does not support creation of custom engines larger than 10 GB.
  **Cloudera Bug:** DSE-4420
- Cloudera Data Science Workbench does not support pulling images from registries that require Docker credentials.
  **Cloudera Bug:** DSE-1521
- The contents of certain pre-existing standard directories such as /home/cdsw, /tmp, /opt/cloudera, and so on, cannot be modified while creating customized engines. This means any files saved in these directories will not be accessible from sessions that are running on customized engines.
  **Workaround:** Create a new custom directory in the Dockerfile used to create the customized engine, and save your files to that directory. Or, create a new custom directory on all the Cloudera Data Science Workbench gateway hosts and save your files to those directories. Then, mount this directory to the custom engine.

Related Resources

- This Cloudera Engineering Blog post on Customizing Docker Images in Cloudera Data Science Workbench describes an end-to-end example on how to build and publish a customized Docker image and use it as an engine in Cloudera Data Science Workbench.
- For an example of how to extend the base engine image to include Conda, see Creating an Extensible Engine With Conda on page 183.

Cloudera Data Science Workbench Engine Versions and Packaging

This topic lists the packages included in the Python and R kernels of the base engine images that ship with each release of Cloudera Data Science Workbench.

Base Engine 7

This section lists the Python, R, and Scala libraries that ship with engine 7.

**Python Libraries in Base Engine 7**

Engine 7 ships **Python 2.7.11 and 3.6.1**. This section lists the libraries that ship with the Python kernels in engine 7. Items in bold indicate a new version since the last release.

<table>
<thead>
<tr>
<th>Library</th>
<th>Version</th>
</tr>
</thead>
<tbody>
<tr>
<td>ipython</td>
<td>5.1.0</td>
</tr>
<tr>
<td>requests</td>
<td>2.13.0</td>
</tr>
<tr>
<td>Flask</td>
<td>0.12.0</td>
</tr>
<tr>
<td>simplejson</td>
<td>3.10.0</td>
</tr>
<tr>
<td>numpy</td>
<td>1.12.1</td>
</tr>
<tr>
<td>pandas</td>
<td>0.20.1</td>
</tr>
<tr>
<td>pandas-datareader</td>
<td>0.2.1</td>
</tr>
<tr>
<td>py4j</td>
<td>0.10.7</td>
</tr>
<tr>
<td>futures</td>
<td>2.1.4</td>
</tr>
</tbody>
</table>
### R Libraries in Base Engine 7

Engine 7 ships **R version 3.5.1**. This section lists the libraries that ship with the R kernel in engine 7.

Items in **bold** indicate a new version since the last release.

<table>
<thead>
<tr>
<th>Package</th>
<th>Version</th>
</tr>
</thead>
<tbody>
<tr>
<td>RCurl</td>
<td>1.95.4.11</td>
</tr>
<tr>
<td>caTools</td>
<td>1.17.1.1</td>
</tr>
<tr>
<td>svTools</td>
<td>0.9.5</td>
</tr>
<tr>
<td>png</td>
<td>0.1.7</td>
</tr>
<tr>
<td>RJSONIO</td>
<td>1.3.1.1</td>
</tr>
<tr>
<td>ggplot2</td>
<td>3.1.0</td>
</tr>
<tr>
<td>cluster</td>
<td>2.0.7.1</td>
</tr>
<tr>
<td>codetools</td>
<td>0.2.16</td>
</tr>
<tr>
<td>foreign</td>
<td>0.8.71</td>
</tr>
<tr>
<td>dplyr</td>
<td>0.7.8</td>
</tr>
<tr>
<td>httr</td>
<td>1.4.0</td>
</tr>
<tr>
<td>httpuv</td>
<td>1.4.5.1</td>
</tr>
<tr>
<td>jsonlite</td>
<td>1.6</td>
</tr>
<tr>
<td>magrittr</td>
<td>1.5</td>
</tr>
<tr>
<td>knitr</td>
<td>1.21</td>
</tr>
<tr>
<td>purrr</td>
<td>0.2.5</td>
</tr>
<tr>
<td>tm</td>
<td>0.7.6</td>
</tr>
<tr>
<td>proxy</td>
<td>0.4.22</td>
</tr>
<tr>
<td>data.table</td>
<td>1.12.0</td>
</tr>
<tr>
<td>stringr</td>
<td>1.3.1</td>
</tr>
<tr>
<td>Rook</td>
<td>1.1.1</td>
</tr>
<tr>
<td>rJava</td>
<td>0.9.10</td>
</tr>
<tr>
<td>devtools</td>
<td>2.0.1</td>
</tr>
</tbody>
</table>

### Scala in Base Engine 7

The Scala kernel is based on the **Apache Toree 0.1.x kernel**. For details on how to add more dependencies to the kernel, see [Example: Using External Packages by Adding Jars or Dependencies](#) on page 206.
Base Engine 6

This section lists the Python, R, and Scala libraries that ship with engine 6.

Python Libraries in Base Engine 6

Engine 6 ships Python 2.7.11 and 3.6.1. This section lists the libraries that ship with the Python kernels in engine 6.

Items in **bold** indicate a new version since the last release.

<table>
<thead>
<tr>
<th>Library</th>
<th>Version</th>
</tr>
</thead>
<tbody>
<tr>
<td>ipython</td>
<td>5.1.0</td>
</tr>
<tr>
<td>requests</td>
<td>2.13.0</td>
</tr>
<tr>
<td>Flask</td>
<td>0.12.0</td>
</tr>
<tr>
<td>simplejson</td>
<td>3.10.0</td>
</tr>
<tr>
<td>numpy</td>
<td>1.12.1</td>
</tr>
<tr>
<td>pandas</td>
<td>0.20.1</td>
</tr>
<tr>
<td>pandas-datreader</td>
<td>0.2.1</td>
</tr>
<tr>
<td>py4j</td>
<td>0.10.7</td>
</tr>
<tr>
<td>futures</td>
<td>2.1.4</td>
</tr>
<tr>
<td>matplotlib</td>
<td>0.8.0</td>
</tr>
<tr>
<td>seaborn</td>
<td>0.25.2</td>
</tr>
<tr>
<td>Cython</td>
<td>1.2.0</td>
</tr>
<tr>
<td>kudu-python</td>
<td></td>
</tr>
</tbody>
</table>

R Libraries in Base Engine 6

Engine 6 ships R version 3.4.1. This section lists the libraries that ship with the R kernel in engine 6.

Items in **bold** indicate a new version since the last release.

<table>
<thead>
<tr>
<th>Package</th>
<th>Version</th>
</tr>
</thead>
<tbody>
<tr>
<td>RCurl</td>
<td>1.95.4.10</td>
</tr>
<tr>
<td>caTools</td>
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</tr>
<tr>
<td>svTools</td>
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</tr>
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<td>png</td>
<td>0.1.7</td>
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<tr>
<td>RJSONIO</td>
<td>1.3.0</td>
</tr>
<tr>
<td>ggplot2</td>
<td>3.0.0</td>
</tr>
<tr>
<td>cluster</td>
<td>2.0.7.1</td>
</tr>
<tr>
<td>codetools</td>
<td>0.2.15</td>
</tr>
<tr>
<td>foreign</td>
<td>0.8.70</td>
</tr>
<tr>
<td>dplyr</td>
<td>0.7.6</td>
</tr>
<tr>
<td>httr</td>
<td>1.3.1</td>
</tr>
<tr>
<td>httpuv</td>
<td>1.4.4.2</td>
</tr>
<tr>
<td>jsonlite</td>
<td>1.5</td>
</tr>
</tbody>
</table>
### Scala in Base Engine 6

The Scala kernel is based on the [Apache Toree 0.1.x kernel](https://github.com/toree/toree). For details on how to add more dependencies to the kernel, see [Example: Using External Packages by Adding Jars or Dependencies](#) on page 206.

### Base Engine 5

This section lists the Python, R, and Scala libraries that ship with engine 5.

#### Python Libraries in Base Engine 5

Engine 5 ships **Python 2.7.11 and 3.6.1**. This section lists the libraries that ship with the Python kernels in engine 5.

<table>
<thead>
<tr>
<th>Library</th>
<th>Version</th>
</tr>
</thead>
<tbody>
<tr>
<td>ipython</td>
<td>5.1.0</td>
</tr>
<tr>
<td>requests</td>
<td>2.13.0</td>
</tr>
<tr>
<td>Flask</td>
<td>0.12.0</td>
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<td>simplejson</td>
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</tr>
<tr>
<td>numpy</td>
<td>1.12.1</td>
</tr>
<tr>
<td>pandas</td>
<td>0.20.1</td>
</tr>
<tr>
<td>pandas-datareader</td>
<td>0.2.1</td>
</tr>
<tr>
<td>py4j</td>
<td>0.10.7</td>
</tr>
<tr>
<td>futures</td>
<td>2.1.4</td>
</tr>
<tr>
<td>matplotlib</td>
<td>2.0.0</td>
</tr>
<tr>
<td>seaborn</td>
<td>0.8.0</td>
</tr>
<tr>
<td>Cython</td>
<td>0.25.2</td>
</tr>
<tr>
<td>kudu-python</td>
<td>1.2.0</td>
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</table>

#### R Libraries in Base Engine 5

Engine 5 ships **R version 3.4.1**. This section lists the libraries that ship with the R kernel in engine 5.

<table>
<thead>
<tr>
<th>Library</th>
<th>Version</th>
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</thead>
<tbody>
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</tr>
<tr>
<td>knitr</td>
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</tr>
<tr>
<td>purrr</td>
<td>0.2.5</td>
</tr>
<tr>
<td>tm</td>
<td>0.7.4</td>
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<tr>
<td>proxy</td>
<td>0.4.22</td>
</tr>
<tr>
<td>data.table</td>
<td>1.11.4</td>
</tr>
<tr>
<td>stringr</td>
<td>1.3.1</td>
</tr>
<tr>
<td>Rook</td>
<td>1.1.1</td>
</tr>
<tr>
<td>rJava</td>
<td>0.9.10</td>
</tr>
<tr>
<td>devtools</td>
<td>1.13.6</td>
</tr>
</tbody>
</table>
Items in **bold** indicate a new version since the last release.

<table>
<thead>
<tr>
<th>Package</th>
<th>Version</th>
</tr>
</thead>
<tbody>
<tr>
<td>RCurl</td>
<td>1.95.4.10</td>
</tr>
<tr>
<td>caTools</td>
<td>1.17.1</td>
</tr>
<tr>
<td>svTools</td>
<td>0.9.4</td>
</tr>
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<td>png</td>
<td>0.1.7</td>
</tr>
<tr>
<td>RJSONIO</td>
<td>1.3.0</td>
</tr>
<tr>
<td>ggplot2</td>
<td>2.2.1</td>
</tr>
<tr>
<td>cluster</td>
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<tr>
<td>codetools</td>
<td>0.2.15</td>
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<td>1.3.1</td>
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<td>httpuv</td>
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<tr>
<td>jsonlite</td>
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</tr>
<tr>
<td>magrittr</td>
<td>1.5</td>
</tr>
<tr>
<td>knitr</td>
<td><strong>1.20</strong></td>
</tr>
<tr>
<td>purrr</td>
<td><strong>0.2.5</strong></td>
</tr>
<tr>
<td>tm</td>
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<tr>
<td>proxy</td>
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<td>data.table</td>
<td><strong>1.11.4</strong></td>
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<td>stringr</td>
<td><strong>1.3.1</strong></td>
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<tr>
<td>Rook</td>
<td>1.1.1</td>
</tr>
<tr>
<td>rJava</td>
<td><strong>0.9.10</strong></td>
</tr>
<tr>
<td>devtools</td>
<td><strong>1.13.5</strong></td>
</tr>
</tbody>
</table>

**Scala in Base Engine 5**

The Scala kernel is based on the Apache Toree 0.1.x kernel. For details on how to add more dependencies to the kernel, see [Example: Using External Packages by Adding Jars or Dependencies](#) on page 206.

**Base Engine 4**

This section lists the Python, R, and Scala libraries that ship with engine 4.

**Python Libraries in Base Engine 4**

Engine 4 ships **Python 2.7.11 and 3.6.1**. This section lists the libraries that ship with the Python kernels in engine 4.

<table>
<thead>
<tr>
<th>Library</th>
<th>Version</th>
</tr>
</thead>
<tbody>
<tr>
<td>ipython</td>
<td>5.1.0</td>
</tr>
<tr>
<td>requests</td>
<td>2.13.0</td>
</tr>
<tr>
<td>Flask</td>
<td>0.12.0</td>
</tr>
<tr>
<td>Library</td>
<td>Version</td>
</tr>
<tr>
<td>------------------</td>
<td>---------</td>
</tr>
<tr>
<td>simplejson</td>
<td>3.10.0</td>
</tr>
<tr>
<td>numpy</td>
<td>1.12.1</td>
</tr>
<tr>
<td>pandas</td>
<td>0.20.1</td>
</tr>
<tr>
<td>pandas-datareader</td>
<td>0.2.1</td>
</tr>
<tr>
<td>py4j</td>
<td>0.10.4</td>
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<td>futures</td>
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<td>matplotlib</td>
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<td>seaborn</td>
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</tr>
<tr>
<td>Cython</td>
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</table>

**R Libraries in Base Engine 4**

Engine 4 ships **R version 3.4.1**. This section lists the libraries that ship with the R kernel in engine 4.

<table>
<thead>
<tr>
<th>Package</th>
<th>Version</th>
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<tbody>
<tr>
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<td>caTools</td>
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<tr>
<td>svTools</td>
<td>0.9.4</td>
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<td>0.1.7</td>
</tr>
<tr>
<td>RjsonIO</td>
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<tr>
<td>ggplot2</td>
<td>2.2.1</td>
</tr>
<tr>
<td>cluster</td>
<td>2.0.6</td>
</tr>
<tr>
<td>codetools</td>
<td>0.2.15</td>
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<tr>
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</tr>
<tr>
<td>dplyr</td>
<td>0.7.4</td>
</tr>
<tr>
<td>httr</td>
<td>1.3.1</td>
</tr>
<tr>
<td>httpuv</td>
<td>1.3.5</td>
</tr>
<tr>
<td>jsonlite</td>
<td>1.5</td>
</tr>
<tr>
<td>magrittr</td>
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<td>knitr</td>
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<tr>
<td>purrr</td>
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<td>0.7.3</td>
</tr>
<tr>
<td>proxy</td>
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</tr>
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<td>data.table</td>
<td>1.10.4.3</td>
</tr>
<tr>
<td>stringr</td>
<td>1.2.0</td>
</tr>
<tr>
<td>Rook</td>
<td>1.1.1</td>
</tr>
<tr>
<td>rJava</td>
<td>0.9.9</td>
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</tbody>
</table>
Scala in Base Engine 4

The Scala kernel is based on the Apache Torque 0.1.x kernel. For details on how to add more dependencies to the kernel, see Example: Using External Packages by Adding Jars or Dependencies on page 206.

Base Engine 3

This section lists the Python, R, and Scala libraries that ship with engine 3.

Python Libraries in Base Engine 3

Engine 3 ships Python 2.7.11 and 3.6.1. This section lists the libraries that ship with the Python kernels in engine 3.

<table>
<thead>
<tr>
<th>Library</th>
<th>Version</th>
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</thead>
<tbody>
<tr>
<td>ipython</td>
<td>5.1.0</td>
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<tr>
<td>requests</td>
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<tr>
<td>Flask</td>
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</tr>
<tr>
<td>simplejson</td>
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<tr>
<td>numpy</td>
<td>1.12.1</td>
</tr>
<tr>
<td>pandas</td>
<td>0.20.1</td>
</tr>
<tr>
<td>pandas-datareader</td>
<td>0.2.1</td>
</tr>
<tr>
<td>py4j</td>
<td>0.10.4</td>
</tr>
<tr>
<td>futures</td>
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</tr>
<tr>
<td>matplotlib</td>
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</tr>
<tr>
<td>seaborn</td>
<td>0.25.2</td>
</tr>
<tr>
<td>Cython</td>
<td>1.2.0</td>
</tr>
<tr>
<td>kudu-python</td>
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</tbody>
</table>

R Libraries in Base Engine 3

Engine 3 ships R version 3.4.1. This section lists the libraries that ship with the R kernel in engine 3.

<table>
<thead>
<tr>
<th>Package</th>
<th>Version</th>
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<tbody>
<tr>
<td>RCurl</td>
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<td>caTools</td>
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<td>svTools</td>
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<tr>
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<td>RJSONIO</td>
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<tr>
<td>ggplot2</td>
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<tr>
<td>foreign</td>
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</tr>
<tr>
<td>Package</td>
<td>Version</td>
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<td>---------</td>
<td>---------</td>
</tr>
<tr>
<td>dplyr</td>
<td>0.7.4</td>
</tr>
<tr>
<td>httr</td>
<td>1.3.1</td>
</tr>
<tr>
<td>httpuv</td>
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</tr>
<tr>
<td>jsonlite</td>
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<tr>
<td>magrittr</td>
<td>1.5</td>
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<td>knitr</td>
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<td>purrr</td>
<td>0.2.4</td>
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<td>tm</td>
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<tr>
<td>proxy</td>
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<td>rJava</td>
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<tr>
<td>devtools</td>
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</tr>
</tbody>
</table>

### Scala in Base Engine 3

The Scala kernel is based on the [Apache Toree 0.1.x kernel](https://toree.apache.org). For details on how to add more dependencies to the kernel, see [Example: Using External Packages by Adding Jars or Dependencies](#) on page 206.

### Base Engine 2

This section lists the Python, R, and Scala libraries that ship with engine 2.

#### Python 2 Libraries in Base Engine 2

Engine 2 ships **Python 2.7.11 and 3.6.1**. This section lists the libraries that ship with the Python kernels in engine 2.

<table>
<thead>
<tr>
<th>Library</th>
<th>Version</th>
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</thead>
<tbody>
<tr>
<td>ipython</td>
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<td>requests</td>
<td>2.13.0</td>
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<tr>
<td>Flask</td>
<td>0.12.0</td>
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<tr>
<td>numpy</td>
<td>1.12.1</td>
</tr>
<tr>
<td>pandas</td>
<td>0.20.1</td>
</tr>
<tr>
<td>pandas-datareader</td>
<td>0.2.1</td>
</tr>
<tr>
<td>py4j</td>
<td>0.10.4</td>
</tr>
<tr>
<td>futures</td>
<td>2.1.4</td>
</tr>
<tr>
<td>matplotlib</td>
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</tr>
<tr>
<td>seaborn</td>
<td>0.7.1</td>
</tr>
<tr>
<td>Cython</td>
<td>0.23.4</td>
</tr>
</tbody>
</table>
### R Libraries in Base Engine 2

Engine 2 ships R version 3.3.0. This section lists the libraries that ship with the R kernel in engine 2.

<table>
<thead>
<tr>
<th>Package</th>
<th>Version</th>
</tr>
</thead>
<tbody>
<tr>
<td>RCurl</td>
<td>1.95.4.8</td>
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<tr>
<td>caTools</td>
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<td>svTools</td>
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<td>RJASONIO</td>
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<tr>
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<td>codetools</td>
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<tr>
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<td>knitr</td>
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<tr>
<td>purrr</td>
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<td>data.table</td>
<td>1.10.4</td>
</tr>
<tr>
<td>stringr</td>
<td>1.2.0</td>
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<tr>
<td>Rook</td>
<td>1.1.1</td>
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<tr>
<td>rJava</td>
<td>0.9.8</td>
</tr>
<tr>
<td>devtools</td>
<td>1.13.3</td>
</tr>
</tbody>
</table>

### Scala in Base Engine 2

The Scala kernel is based on the Apache Toree 0.1.x kernel. For details on how to add more dependencies to the kernel, see Example: Using External Packages by Adding Jars or Dependencies on page 206.

### Jupyter Magic Commands

Cloudera Data Science Workbench’s Scala and Python kernels are based on Jupyter kernels. Jupyter kernels support varying magic commands that extend the core language with useful shortcuts. This section details the magic commands (magics) supported by Cloudera Data Science Workbench.

**Line magics** begin with a single %: for example, `%timeit`. **Cell magics** begin with a double %: for example, `%%bash`.

#### Python

In the default Python 2.7.11 engine, Cloudera Data Science Workbench supports most line magics, but no cell magics.

Cloudera Data Science Workbench supports the shell magic `!: for example, `!ls -alh /home/cdsw`. 
Cloudera Data Science Workbench supports the help magics `?` and `??`: for example, `?numpy` and `??numpy`. `?` displays the docstring for its argument. `??` attempts to print the source code. You can get help on magics using the `?` prefix: for example, `?%timeit`.

Cloudera Data Science Workbench supports the line magics listed at [https://ipython.org/ipython-doc/3/interactive/magics.html#line-magics](https://ipython.org/ipython-doc/3/interactive/magics.html#line-magics), with the following exceptions:

- `%colors`
- `%debug`
- `%edit`
- `%gui`
- `%history`
- `%install_default_config`
- `%install_profiles`
- `%lsmagic`
- `%macro`
- `%matplotlib`
- `%notebook`
- `%page`
- `%pastebin`
- `%pdb`
- `%prun`
- `%pylab`
- `%recall`
- `%rerun`
- `%save`
- `%sc`

**Scala**

Cloudera Data Science Workbench's Scala kernel is based on Apache Toree. It supports the line magics documented in the Apache Toree [magic tutorial](https://toree.apache.org/docs/magic-tutorial).
Using CDS 2.x Powered by Apache Spark

Apache Spark is a general purpose framework for distributed computing that offers high performance for both batch and stream processing. It exposes APIs for Java, Python, R, and Scala, as well as an interactive shell for you to run jobs.

Cloudera Data Science Workbench provides interactive and batch access to Spark 2. Connections are fully secure without additional configuration, with each user accessing Spark using their Kerberos principal. With a few extra lines of code, you can do anything in Cloudera Data Science Workbench that you might do in the Spark shell, as well as leverage all the benefits of the workbench. Your Spark applications will run in an isolated project workspace.

Cloudera Data Science Workbench’s interactive mode allows you to launch a Spark application and work iteratively in R, Python, or Scala, rather than the standard workflow of launching an application and waiting for it to complete to view the results. Because of its interactive nature, Cloudera Data Science Workbench works with Spark on YARN’s client mode, where the driver persists through the lifetime of the job and runs executors with full access to the CDH cluster resources. This architecture is illustrated the following figure:

The rest of this guide describes how to set Spark 2 environment variables, manage package dependencies, and how to configure logging. It also consists of instructions and sample code for running R, Scala, and Python projects from Spark 2.

Cloudera Data Science Workbench allows you to access the Spark History Server and even transient per-session UIs for Spark 2 directly from the workbench console. For more details, see Accessing Web User Interfaces from Cloudera Data Science Workbench on page 123.

Configuring CDS 2.x Powered by Apache Spark 2

This topic describes how to set Spark 2 environment variables, manage package dependencies for Spark 2 jobs, and how to configure logging.
Spark Configuration Files

Cloudera Data Science Workbench supports configuring Spark 2 properties on a per project basis with the spark-defaults.conf file. If there is a file called spark-defaults.conf in your project root, this will be automatically be added to the global Spark defaults. To specify an alternate file location, set the environmental variable, SPARK_CONFIG, to the path of the file relative to your project. If you’re accustomed to submitting a Spark job with key-values pairs following a --conf flag, these can also be set in a spark-defaults.conf file instead. For a list of valid key-value pairs, refer the Spark configuration reference documentation.

Administrators can set environment variable paths in the /etc/spark2/conf/spark-env.sh file.

You can also use Cloudera Manager to configure spark-defaults.conf and spark-env.sh globally for all Spark applications as follows.

**Important:** Cloudera Data Science Workbench does not automatically detect configuration changes on the CDH cluster. Therefore, any changes made to CDH services must be followed by a full reset of Cloudera Data Science Workbench. Review the associated known issue here: CDH Integration Issues.

### Configuring Global Properties Using Cloudera Manager

Configure client configuration properties for all Spark applications in spark-defaults.conf as follows:

1. Go to the Cloudera Manager Admin Console.
2. Navigate to the Spark service.
3. Click the Configuration tab.
4. Search for the Spark Client Advanced ConfigurationSnippet (Safety Valve) for spark-conf/spark-defaults.conf property.
5. Specify properties described in Application Properties. If more than one role group applies to this configuration, edit the value for the appropriate role group.
6. Click Save Changes to commit the changes.
7. Deploy the client configuration.
8. Restart Cloudera Data Science Workbench.

For more information on using a spark-defaults.conf file for Spark jobs, visit the Apache Spark 2 reference documentation.

### Configuring Spark Environment Variables Using Cloudera Manager

Configure service-wide environment variables for all Spark applications in spark-env.sh as follows:

1. Go to the Cloudera Manager Admin Console.
2. Navigate to the Spark 2 service.
3. Click the Configuration tab.
4. Search for the Spark Service Advanced ConfigurationSnippet (Safety Valve) for spark-conf/spark-env.sh property and add the paths for the environment variables you want to configure.
5. Click Save Changes to commit the changes.
6. Restart the service.
7. Deploy the client configuration.
8. Restart Cloudera Data Science Workbench.

### Managing Memory Available for Spark Drivers

By default, the amount of memory allocated to Spark driver processes is set to a 0.8 fraction of the total memory allocated for the engine container. If you want to allocate more or less memory to the Spark driver process, you can override this default by setting the spark.driver.memory property in spark-defaults.conf (as described above).
Managing Dependencies for Spark 2 Jobs

As with any Spark job, you can add external packages to the executor on startup. To add external dependencies to Spark jobs, specify the libraries you want added by using the appropriate configuration parameter in a `spark-defaults.conf` file. The following table lists the most commonly used configuration parameters for adding dependencies and how they can be used:

<table>
<thead>
<tr>
<th>Property</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td><code>spark.files</code></td>
<td>Comma-separated list of files to be placed in the working directory of each Spark executor.</td>
</tr>
<tr>
<td><code>spark.submit.pyFiles</code></td>
<td>Comma-separated list of .zip, .egg, or .py files to place on PYTHONPATH for Python applications.</td>
</tr>
<tr>
<td><code>spark.jars</code></td>
<td>Comma-separated list of local jars to include on the Spark driver and Spark executor classpaths.</td>
</tr>
<tr>
<td><code>spark.jars.packages</code></td>
<td>Comma-separated list of Maven coordinates of jars to include on the Spark driver and Spark executor classpaths. When configured, Spark will search the local Maven repo, and then Maven central and any additional remote repositories configured by <code>spark.jars.ivy</code>. The format for the coordinates are <code>groupId:artifactId:version</code>.</td>
</tr>
<tr>
<td><code>spark.jars.ivy</code></td>
<td>Comma-separated list of additional remote repositories to search for the coordinates given with <code>spark.jars.packages</code>.</td>
</tr>
</tbody>
</table>

Example `spark-defaults.conf`

Here is a sample `spark-defaults.conf` file that uses some of the Spark configuration parameters discussed in the previous section to add external packages on startup.

```bash
spark.jars.packages org.scalaj:scalaj-http_2.11:2.3.0
spark.jars my_sample.jar
spark.files data/test_data_1.csv,data/test_data_2.csv
```

- **`spark.jars.packages`**
  The `scalaj` package will be downloaded from Maven central and included on the Spark driver and executor classpaths.

- **`spark.jars`**
  The pre-existing jar, `my_sample.jar`, residing in the root of this project will be included on the Spark driver and executor classpaths.

- **`spark.files`**
  The two sample data sets, `test_data_1.csv` and `test_data_2.csv`, from the `/data` directory of this project will be distributed to the working directory of each Spark executor.

For more advanced configuration options, visit the Apache Spark 2 reference documentation.

Spark Logging Configuration

Cloudera Data Science Workbench allows you to update Spark’s internal logging configuration on a per-project basis. Spark 2 uses Apache Log4j, which can be configured through a properties file. By default, the `log4j.properties` file found in the root of your project will be appended to the existing Spark logging properties for every session and job.

To specify a custom location, set the environmental variable `LOG4J_CONFIG` to the file location relative to your project.

The Log4j documentation has more details on logging options.
Increasing the log level or pushing logs to an alternate location for troublesome jobs can be very helpful for debugging. For example, this is a log4j.properties file in the root of a project that sets the logging level to INFO for Spark jobs.

```
shell.log.level=INFO
```

PySpark logging levels should be set as follows:

```
log4j.logger.org.apache.spark.api.python.PythonGatewayServer=<LOG_LEVEL>
```

And Scala logging levels should be set as:

```
log4j.logger.org.apache.spark.repl.Main=<LOG_LEVEL>
```

### Setting Up an HTTP Proxy for Spark 2

In Cloudera Data Science Workbench clusters that use an HTTP proxy, follow these steps to support web-related actions in Spark. You must set the Spark configuration parameter extraJavaOptions on your gateway hosts.

To set up a Spark proxy:

1. Log in to Cloudera Manager.
2. Go to Spark2 > Configuration.
3. Filter the properties with Scope > Gateway and Category > Advanced.
4. Scroll down to Spark 2 Client Advanced Configuration Snippet (Safety Valve) for spark2-conf/spark-defaults.conf.
5. Enter the following configuration code, substituting your proxy host and port values:

   ```
   spark.driver.extraJavaOptions=\
   -Dhttp.proxyHost=<YOUR_HTTP_PROXY_HOST>\
   -Dhttp.proxyPort=<HTTP_PORT> \
   -Dhttps.proxyHost=<YOUR_HTTPS_PROXY_HOST> \
   -Dhttps.proxyPort=<HTTPS_PORT>
   ```
6. Click Save Changes.
7. Choose Actions > Deploy Client Configuration.

### Using Spark 2 from Python

Cloudera Data Science Workbench supports using Spark 2 from Python via PySpark.

### Setting Up a PySpark Project

**PySpark Environment Variables**

The default Cloudera Data Science Workbench engine currently includes Python 2.7.11 and Python 3.6.1. To use PySpark with lambda functions that run within the CDH cluster, the Spark executors must have access to a matching version of Python. For many common operating systems, the default system Python will not match the minor release of Python included in Data Science Workbench.

To ensure that the Python versions match, Python can either be installed on every CDH host or made available per job run using Spark’s ability to distribute dependencies. Given the size of a typical isolated Python environment and the desire to avoid repeated uploads from gateway hosts, Cloudera recommends installing Python 2.7 and 3.6 on the cluster if you are using PySpark with lambda functions.

You can install Python 2.7 and 3.6 on the cluster using any method and set the corresponding PYSPARK_PYTHON environment variable in your project. Cloudera Data Science Workbench 1.3 (and higher) include a separate environment variable for Python 3 sessions called PYSPARK3_PYTHON. Python 2 sessions continue to use the default PYSPARK_PYTHON.
variable. This will allow you to run Python 2 and Python 3 sessions in parallel without either variable being overridden by the other.

Creating and Running a PySpark Project

To get started quickly, use the PySpark template project to create a new project. For instructions, see Creating a Project on page 107.

To run a PySpark project, navigate to the project's overview page, open the workbench console and launch a Python session. For detailed instructions, see Using the Workbench on page 110.

Testing a PySpark Project in Spark Local Mode

Spark's local mode is often useful for testing and debugging purposes. Use the following sample code snippet to start a PySpark session in local mode.

```python
from pyspark.sql import SparkSession
spark = SparkSession(
    .builder(
        .appName("LocalSparkSession")
        .master("local")
    ).getOrCreate()

partitions = int(sys.argv[1]) if len(sys.argv) > 1 else 2
n = 100000 * partitions

def f(_):
    x = random() * 2 - 1
    y = random() * 2 - 1
    return 1 if x ** 2 + y ** 2 < 1 else 0

count = spark.sparkContext.parallelize(range(1, n + 1), partitions).map(f).reduce(add)
print("Pi is roughly %f" % (4.0 * count / n))
```

For more details, refer the Spark documentation: Running Spark Applications.

Example: Montecarlo Estimation

Within the template PySpark project, pi.py is a classic example that calculates Pi using the Montecarlo Estimation.

What follows is the full, annotated code sample that can be saved to the pi.py file.

```python
# Estimating \pi
#
# This PySpark example shows you how to estimate \pi in parallel
# using Monte Carlo integration.

from __future__ import print_function
import sys
from random import random
from operator import add

# Connect to Spark by creating a Spark session
from pyspark.sql import SparkSession
spark = SparkSession(
    .builder(
        .appName("PythonPi")
    ).getOrCreate()

partitions = int(sys.argv[1]) if len(sys.argv) > 1 else 2
n = 100000 * partitions

def f(_):
    x = random() * 2 - 1
    y = random() * 2 - 1
    return 1 if x ** 2 + y ** 2 < 1 else 0

# To access the associated SparkContext
count = spark.sparkContext.parallelize(range(1, n + 1), partitions).map(f).reduce(add)
print("Pi is roughly %f" % (4.0 * count / n))
```

spark.stop()
Example: Locating and Adding JARs to Spark 2 Configuration

This example shows how to discover the location of JAR files installed with Spark 2, and add them to the Spark 2 configuration.

```python
# # Using Avro data
# # This example shows how to use a JAR file on the local filesystem on
# # Spark on Yarn.
from __future__ import print_function
import os, sys
import os.path
from functools import reduce
from pyspark.sql import SparkSession
from pyspark.files import SparkFiles

# Add the data file to HDFS for consumption by the Spark executors.
!hdfs dfs -put resources/users.avro /tmp

# Find the example JARs provided by the Spark parcel. This parcel
# is available on both the driver, which runs in Cloudera Data Science Workbench, and
# the
# executors, which run on Yarn.
exampleDir = os.path.join(os.environ['SPARK_HOME'], "examples/jars")
exampleJars = [os.path.join(exampleDir, x) for x in os.listdir(exampleDir)]

# Add the Spark JARs to the Spark configuration to make them available for use.
spark = SparkSession.builder.config("spark.jars", ",".join(exampleJars))
    .appName("AvroKeyInputFormat")
    .getOrCreate()
sc = spark.sparkContext

# Read the schema.
schema = open("resources/user.avsc").read()
conf = "{avro.schema.input.key": schema }"

avro_rdd = sc.newAPIHadoopFile("/tmp/users.avro", # This is an HDFS path!
    "org.apache.avro.mapreduce.AvroKeyInputFormat",
    "org.apache.avro.mapred.AvroKey",
    "org.apache.hadoop.io.NullWritable",
    keyConverter="org.apache.spark.examples.pythonconverters.AvroWrapperToJavaConverter",
    conf=conf)
output = avro_rdd.map(lambda x: x[0]).collect()
for k in output:
    print(k)
spark.stop()
```

Example: Distributing Dependencies on a PySpark Cluster

Although Python is a popular choice for data scientists, it is not straightforward to make a Python library available on a distributed PySpark cluster. To determine which dependencies are required on the cluster, you must understand that Spark code applications run in Spark executor processes distributed throughout the cluster. If the Python code you are running uses any third-party libraries, Spark executors require access to those libraries when they run on remote executors.

This example demonstrates a way to run the following Python code (nltk_sample.py), that includes pure Python libraries (nltk), on a distributed PySpark cluster.

1. (Prerequisites)
   - Make sure the Anaconda parcel has been distributed and activated on your cluster.
   - Create a new project in Cloudera Data Science Workbench. In that project, create a new file called nltk_sample.py with the following sample script.
import os
import sys
from pyspark.sql import SparkSession

spark = SparkSession.builder.
    .appName("spark-nltk")
    .getOrCreate()

data = spark.sparkContext.textFile('1970-Nixon.txt')

def word_tokenize(x):
    import nltk
    return nltk.word_tokenize(x)

def pos_tag(x):
    import nltk
    return nltk.pos_tag([x])

words = data.flatMap(word_tokenize)
words.saveAsTextFile('nixon_tokens')

pos_word = words.map(pos_tag)
pos_word.saveAsTextFile('nixon_token_pos')

2. Go to the project you created and launch a new PySpark session.
3. Click Terminal Access and run the following command to pack the Python environment into conda.

```
conda create -n nltk_env --copy -y -q python=2.7.11 nltk numpy
```

The `--copy` option allows you to copy whole dependent packages into certain directory of a conda environment.

If you want to add extra pip packages without conda, you should copy packages manually after using `pip install`. In Cloudera Data Science Workbench, pip will install packages into the `~/.local` directory in a project.

```
pip install some-awesome-package
cp -r ~/.local/lib ~/.conda/envs/nltk_env/
```

Zip the conda environment for shipping on PySpark cluster.

```
zip -r ../../nltk_env.zip nltk_env
```

4. (Specific to NLTK) For this example, you can use NLTK data as input.

```
cd ~/nltk_data/tokenizers/
zip -r ../../tokenizers.zip *
```
5. Set spark-submit options in `spark-defaults.conf`.

```bash
spark.yarn.appMasterEnv.PYSPARK_PYTHON=./NLTK/nltk_env/bin/python
spark.yarn.appMasterEnv.NLTK_DATA=/
spark.executorEnv.NLTK_DATA=/
spark.yarn.dist.archives=nltk_env.zip#NLTK,tokenizers.zip#tokenizers,tagger.zip#taggers
```

With these settings, PySpark unzips `nltk_env.zip` into the NLTK directory. `NLTK_DATA` is the environmental variable where NLTK data is stored.

6. In Cloudera Data Science Workbench, set the `PYSPARK_PYTHON` environment variable to the newly-created environment. To do this, navigate back to the Project Overview page and click Settings > Engine > Environment Variables. Set `PYSPARK_PYTHON` to `./NLTK/nltk_env/bin/python` and click Add. Then click Save Environment.

7. Launch a new PySpark session and run the `nltk_sample.py` script in the workbench. You can test whether the script ran successfully using the following command:

```bash
!hdfs dfs -cat ./nixon_tokens/* | head -n 20
```

```
Annual
Message
to
the
Congress
on
the
State
of
the
Union
;
January
22
,
1970
Mr.
Speaker
;
Mr.
```

```
! hdfs dfs -cat nixon_token_pos/* | head -n 20
[(u'Annual', 'JJ')]
[(u'Message', 'NN')]
[(u'to', 'TO')]
[(u'the', 'DT')]
[(u'Congress', 'NNP')]
[(u'on', 'IN')]
[(u'the', 'DT')]
[(u'State', 'NNP')]
[(u'of', 'IN')]
[(u'Union', 'NN')]
[(u',', '')]
[(u'January', 'NNP')]
[(u'22', 'CD')]
[(u',', '')]
[(u'1970', 'CD')]
[(u'Mr.', 'NNP')]
[(u'Speaker', 'NN')]
[(u',', '')]
[(u'Mr.', 'NNP')]
```

---

**Using Spark 2 from R**

R users can access Spark 2 using `sparklyr`. Although Cloudera does not ship or support sparklyr, we do recommend using `sparklyr` as the R interface for Cloudera Data Science Workbench.
Installing sparklyr

Install the latest version of sparklyr as follows.

```
install.packages("sparklyr")
```

**Note:** The `spark_apply()` function requires the R Runtime environment to be pre-installed on your cluster. This will likely require intervention from your cluster administrator. For details, refer the RStudio documentation.

Connecting to Spark 2

You can connect to local instances of Spark 2 as well as remote clusters.

```
library(sparklyr)
system.time(sc <- spark_connect(master = "yarn-client"))
```

For a complete example, see *Sparklyr (R)* on page 137.

Using Spark 2 from Scala

This topic describes how to set up a Scala project for CDS 2.x Powered by Apache Spark along with a few associated tasks. Cloudera Data Science Workbench provides an interface to the Spark 2 shell (v 2.0+) that works with Scala 2.11.

Accessing Spark 2 from the Scala Engine

Unlike PySpark or Sparklyr, you can access a SparkContext assigned to the `spark` (SparkSession) and `sc` (SparkContext) objects on console startup, just as when using the Spark shell. By default, the application name will be set to `CDSW_sessionID`, where sessionid is the id of the session running your Spark code. To customize this, set the `spark.app.name` property to the desired application name in a `spark-defaults.conf` file.

`Pi.scala` is a classic starting point for calculating Pi using Monte Carlo Estimation.

This is the full, annotated code sample.

```
//Calculate pi with Monte Carlo estimation
import scala.math.random

//make a very large unique set of 1 -> n
val partitions = 2
val n = math.min(100000L * partitions, Int.MaxValue).toInt
val xs = 1 until n

//split up n into the number of partitions we can use
val rdd = sc.parallelize(xs, partitions).setName("'N values rdd'")

//generate a random set of points within a 2x2 square
val sample = rdd.map { i =>
  val x = random * 2 - 1
  val y = random * 2 - 1
  (x, y)
}.setName("'Random points rdd'")

//points w/in the square also w/in the center circle of r=1
val inside = sample.filter { case (x, y) => (x * x + y * y < 1) }.setName("'Random points inside circle'")
val count = inside.count()
```
// Area(circle)/Area(square) = inside/n => pi=4*inside/n
println("Pi is roughly " + 4.0 * count / n)

Key points to note:

• import scala.math.random

   Importing included packages works just as in the shell, and need only be done once.

• Spark context (sc).

   You can access a SparkContext assigned to the variable sc on console startup.

   val rdd = sc.parallelize(xs, partitions).setName("N values rdd")

Example: Read Files from the Cluster Local Filesystem

   Use the following command in the terminal to read text from the local filesystem. The file must exist on all hosts, and the same path for the driver and executors. In this example you are reading the file ebay-xbox.csv.

   sc.textFile("file:///tmp/ebay-xbox.csv")

Example: Using External Packages by Adding Jars or Dependencies

   External libraries are handled through line magics. Line magics in the Toree kernel are prefixed with %.

   Adding Remote Packages

   You can use Apache Toree's AddDeps magic to add dependencies from Maven central. You must specify the company name, artifact ID, and version. To resolve any transitive dependencies, you must explicitly specify the --transitive flag.

   %AddDeps org.scalaj scalaj-http_2.11 2.3.0
   import scalaj.http._
   response.body
   response.code
   response.headers
   response.cookies

   Adding Remote or Local JARs

   You can use the AddJars magic to distribute local or remote JARs to the kernel and the cluster. Using the -f option ignores cached JARs and reloads.

   %AddJar http://example.com/some_lib.jar -f
   %AddJar file:/path/to/some/lib.jar
Cloudera Data Science Workbench Administration Guide

The following topics describe some common administrative tasks that can only be performed by a Cloudera Data Science Workbench site administrator.

Monitoring Cloudera Data Science Workbench Activity

**Required Role: Site Administrator**

The **Admin > Overview** tab displays basic information about your deployment, such as the number of users signed up, the number of teams and projects created, memory used, and some average job scheduling and run times. You can also see the version of Cloudera Data Science Workbench you are currently running.

The **Admin > Activity** tab of the dashboard displays the following time series charts. These graphs should help site administrators identify basic usage patterns, understand how cluster resources are being utilized over time, and how they are being distributed among teams and users.

- **CPU** - Total number of CPUs requested by sessions running at this time.
  Note that code running inside an n-CPU session, job, experiment or model replica can access at least n CPUs worth of CPU time. Each user pod can utilize all of its host's CPU resources except the amount requested by other user workloads or Cloudera Data Science Workbench application components. For example, a 1-core Python session can use more than 1 core if other cores have not been requested by other user workloads or CDSW application components.

- **Memory** - Total memory (in GiB) requested by sessions running at this time.

- **GPU** - Total number of GPUs requested by sessions running at this time.

- **Runs** - Total number of sessions and jobs running at this time.

- **Lag** - Depicts session scheduling and startup times.
  - **Scheduling Duration**: The amount of time it took for a session pod to be scheduled on the cluster.

**Important**: The graphs and numbers on the **Admin > Activity** page do not account for any resources used by active models on the deployment. For that information, go to **Admin > Models** page.
– **Starting Duration:** The amount of time it took for a session to be ready for user input. This is the amount of time since a pod was scheduled on the cluster until code could be executed.

### Related Resources

- **Models** - [Monitoring Active Models](#) on page 160.
- **Tracking Disk Usage** - [Tracking Disk Usage on the Application Block Device](#) on page 210

### Managing the Cloudera Data Science Workbench Service in Cloudera Manager

This topic describes how to configure and manage Cloudera Data Science Workbench using Cloudera Manager. The contents of this topic only apply to CSD-based deployments. If you installed Cloudera Data Science Workbench using the RPM, the Cloudera Data Science Workbench service will not be available to you in Cloudera Manager.

### Adding the Cloudera Data Science Workbench Service

Cloudera Data Science Workbench is available as an add-on service for Cloudera Manager. To install Cloudera Data Science Workbench, you require the following files: a CSD JAR file that contains all the configuration needed to describe and manage the new Cloudera Data Science Workbench service, and the Cloudera Data Science Workbench parcel.

To install this service, first download and copy the CSD file to the Cloudera Manager Server host. Then use Cloudera Manager to distribute the Cloudera Data Science Workbench parcel to the relevant gateway hosts. You can then use Cloudera Manager's **Add Service** wizard to add the Cloudera Data Science Workbench service to your cluster.

For the complete set of instructions, see [Install Cloudera Data Science Workbench](#) on page 61.

### Roles Associated with the Cloudera Data Science Workbench Service

#### Master

Runs the Kubernetes master components on the CDSW master host.

The Master role must only be assigned to the Cloudera Data Science Workbench master host.

#### Worker

Runs the Kubernetes worker/host components on the CDSW worker hosts.

The Worker role must be assigned to all Cloudera Data Science Workbench worker hosts. **Do not assign the Master and Worker roles to the same host.** Even if you are running a single-host proof-of-concept deployment, the single Master host will be able to run user workloads just as a worker host can.

#### Docker Daemon

Runs underlying Docker processes on **all** Cloudera Data Science Workbench hosts.

The Docker Daemon role must be assigned to every Cloudera Data Science Workbench gateway host.

#### Application

Runs the Cloudera Data Science Workbench application. The Application role must only be assigned to the Cloudera Data Science Workbench master host.

The Application role requires the underlying Docker Daemon and Master/Worker roles to be up and running before the Cloudera Data Science Workbench web application can be started. If you want to restart the CDSW application, you must [restart the CDSW service](#).

Similarly, do not attempt to restart the underlying Docker Daemon role while the Master/Worker roles are still running on a host. This will result in the operation hanging indefinitely. To avoid this, always perform a full service restart.
Accessing Cloudera Data Science Workbench from Cloudera Manager

1. Log into the Cloudera Manager Admin Console.
2. Go to the CDSW service.
3. Click CDSW Web UI to visit the Cloudera Data Science Workbench web application.

Configuring Cloudera Data Science Workbench Properties

In a CSD-based deployment, Cloudera Manager allows you to configure Cloudera Data Science Workbench properties without having to directly edit any configuration file.

1. Log into the Cloudera Manager Admin Console.
2. Go to the CDSW service.
3. Click the Configuration tab.
4. Use the search bar to look for the property you want to configure. You can use Cloudera Manager to configure proxies, enable TLS, reserve the master host, and enable GPU support for Cloudera Data Science Workbench.

   If you have recently migrated from an RPM-based deployment to a CSD-based deployment, a list of the properties in cdsw.conf, along with their corresponding properties in Cloudera Manager can be found in the upgrade guide here.

5. Click Save Changes.

Starting, Stopping, and Restarting the Service

Important: On Cloudera Data Science Workbench 1.4.0 (and lower), do not stop or restart Cloudera Data Science Workbench without using the cdsw_protect_stop_restart.sh script. This is to help avoid the data loss issue detailed in TSB-346.

To start, stop, and restart the Cloudera Data Science Workbench service:

1. Log into the Cloudera Manager Admin Console.
2. On the Home > Status tab, click to the right of the CDSW service and select the action (Start, Stop, or Restart) you want to perform from the dropdown.
3. Confirm your choice on the next screen. When you see a Finished status, the action is complete.

Points to Remember

- After a restart, the Cloudera Data Science Workbench service in Cloudera Manager will display Good health even though the Cloudera Data Science Workbench web application might need a few more minutes to get ready to serve requests.
- The CDSW service must be restarted every time client configuration is redeployed to the Cloudera Data Science Workbench hosts.

Reserving the Master Host for Internal CDSW Components

Important: This feature only applies to deployments with more than one Cloudera Data Science Workbench host. Enabling this feature on single-host deployments will leave Cloudera Data Science Workbench incapable of scheduling any workloads.

Starting with version 1.4.3, Cloudera Data Science Workbench allows you to reserve the master host for running internal application components and services such as Livelog, the PostgreSQL database, and so on, while user workloads run exclusively on worker hosts.
By default, the master host runs both, user workloads as well as the application’s internal services. However, depending on the size of your CDSW deployment and the number of workloads running at any given time, it’s possible that user workloads might dominate resources on the master host. Enabling this feature will ensure that CDSW’s application components always have access to the resources they need on the master host and are not adversely affected by user workloads.

Depending on your deployment type, use one of the following sets of instructions to enable this feature:

**RPM Deployments**

To enable this feature on RPM-based deployments, go to the `/etc/cdsw/config/cdsw.conf` file and set the `RESERVE_MASTER` property to `true`.

**CSD Deployments**

On CSD-based deployments, this feature can be enabled in Cloudera Manager. Note that this feature is not yet available as a configuration property in Cloudera Manager. However, you can use an Advanced Configuration Snippet (Safety Valve) to configure this as follows:

1. Log into the Cloudera Manager Admin Console.
2. Go to the **CDSW** service.
3. Click the **Configuration** tab.
4. Use the search bar to look for the **Master Advanced Configuration Snippet (Safety Valve) for cdsw.properties** property. Add the following string to the value field:

   ```
   RESERVE_MASTER=true
   ```

5. Click **Save Changes**.
6. Restart the CDSW service to have this change go into effect.

**Managing Cloudera Data Science Workbench Worker Hosts**

You can add or remove workers from Cloudera Data Science Workbench using Cloudera Manager. For instructions, see:

- Adding a Worker Host
- Removing a Worker Host

**Health Tests**

Cloudera Manager runs a few health tests to confirm whether Cloudera Data Science Workbench and it’s components (Master and Workers) are running, and ready to serve requests.

You can choose to enable or disable individual or summary health tests, and in some cases specify what should be included in the calculation of overall health for the service, role instance, or host. See Configuring Monitoring Settings for more information.

**Tracking Disk Usage on the Application Block Device**

This section demonstrates how to use Cloudera Manager to chart disk usage on the Application block device over time, and to create a trigger to notify cluster administrators when free space on the block device falls below a certain threshold. The latter is particularly important because once the Application block device runs out of memory, Cloudera Data Science Workbench will stop launching any new sessions or jobs. Advance notifications will give administrators a chance to expand the block device or cleanup existing data before Cloudera Data Science Workbench users run into any problems.

**Create a Chart to Track Disk Usage on the Application Block Device**

The following steps use Cloudera Manager’s Chart Builder to track disk usage on the Application Block Device (mounted to `/var/lib/cdsw` on the CDSW master host) over time.
1. Log into the Cloudera Manager Admin Console.

2. Click Charts > Chart Builder.

3. Enter a tsquery that charts memory usage on the block device. For example, the following tsquery creates a chart to track unallocated memory on the Application block device.

```
select capacity_free where mountpoint="/var/lib/cdsw" and category=FILESYSTEM and hostname="<CDSW_Master_hostname>"
```

Alternatively, you could use the following tsquery to track the amount of memory already in use on the block device.

```
select capacity, capacity_used where mountpoint="/var/lib/cdsw" and category=FILESYSTEM and hostname="<CDSW_Master_hostname>"
```

Make sure you insert the hostname for your master host as indicated in the queries.

4. Click Build Chart. You should see a preview of the chart below.

5. Click Save.

6. Enter a name for the chart.

7. Select Add chart to another dashboard. From the dropdown list of available System Dashboards, select CDH Cloudera Data Science Workbench Status Page.

8. Click Save Chart. If you navigate back to the CDSW service page, you should now see the new chart on this page.

For more details about Cloudera Manager’s Chart Builder, see the following topic in the Cloudera Manager documentation: Charting Time Series Data.

---

Create a Trigger to Notify Cluster Administrators when Free Space Runs Low

The following steps create a trigger to alert Cloudera Manager cluster administrators when free space on the Application Block Device has fallen below a specific threshold.

1. Log in to Cloudera Manager and go to the CDSW service page.

2. Click Create Trigger.

3. Give the trigger a name.

4. Modify the Expression field to include a condition for the trigger to fire. For example, if the trigger should fire when unallocated memory on the Application Block Device falls below 250GB, the expression should be:

```
IF (select capacity_free where mountpoint="/var/lib/cdsw" and category=FILESYSTEM and hostname="<CDSW_Master_hostname>" and LAST (capacity_free) < 250GB) DO health:concerning
```
On the right hand side of the page, you should see a preview of the query you have entered and a chart that displays the result of the query as in the following sample image. Note that if the query is incorrect or incomplete you will not see the preview on the right.

5. Click **Create Trigger**. If you navigate back to the CDSW service page, you should now see the new trigger in the list of Health Tests.

For more details about Triggers, refer the following topic in the Cloudera Manager documentation: [Triggers](#).

### Creating Diagnostic Bundles

Diagnostic data for Cloudera Data Science Workbench is now available as part of the Cloudera Manager diagnostic bundle. For details on usage and diagnostic data collection in Cloudera Data Science Workbench, see [Data Collection in Cloudera Data Science Workbench](#) on page 212.

### Data Collection in Cloudera Data Science Workbench

Cloudera Data Science Workbench collects usage and diagnostic data in two ways, both of which can be controlled by system administrators.

#### Usage Tracking

Cloudera Data Science Workbench collects aggregate usage data by sending limited tracking events to Google Analytics and Cloudera servers. No customer data or personal information is sent as part of these bundles.

#### Disable Usage Tracking

1. Log in to Cloudera Data Science Workbench as a site administrator.
2. Click **Admin**.
3. Click the **Settings** tab.
4. Uncheck **Send usage data to Cloudera**.

In addition to this, Cloudera Manager also collects and sends anonymous usage information using Google Analytics to Cloudera. If you are running a CSD-based deployment and want to disable data collection by Cloudera Manager, see [Managing Anonymous Usage Data Collection](#).

### Diagnostic Bundles

Diagnostic bundles are used to aid in debugging issues filed with Cloudera Support. They can be created using either Cloudera Manager (only for CSD-based deployments), or the command line. This section also provides details on the information collected by Cloudera Data Science workbench as part of these bundles.
Using Cloudera Manager

If you are working on a CSD-based deployment, Cloudera Data Science Workbench logs and diagnostic data are available as part of the diagnostic bundles created by Cloudera Manager. By default, Cloudera Manager is configured to collect diagnostic data weekly and to send it to Cloudera automatically. You can schedule the frequency of data collection on a daily, weekly, or monthly schedule, or even disable the scheduled collection of data. To learn how to configure the frequency of data collection or disable it entirely, see Diagnostic Data Collection by Cloudera Manager.

You can also manually trigger a collection and transfer of diagnostic data to Cloudera at any time. For instructions, see Manually Collecting and Sending Diagnostic Data to Cloudera.

Note: If Cloudera Data Science Workbench data is missing from Cloudera Manager diagnostic bundles, it might be due to this known issue.

Using the Command Line

Diagnostic bundles can be created by system administrators using the `cdsw logs` command. By default, sensitive information will be redacted from the log files in the bundle. This is the bundle that you should attach to any case opened with Cloudera Support. The filename of this generated bundle will be of the form, `cdsw-logs-$hostname-$date-$time.redacted.tar.gz`.

If you want to turn off redaction of log files, you can use the `-x | --skip-redaction` option as demonstrated below. This diagnostic bundle is only meant for internal use. It should be retained at least for the duration of the support case, in case any critical information was redacted. However, it can be shared with Cloudera at your discretion.

```bash
cdsw logs --skip-redaction
```

The filename of this bundle will be of the form, `cdsw-logs-$hostname-$date-$time.tar.gz`.

The contents of both archives are stored in text and can easily be inspected by system administrators. Both forms are designed to be easily diff-able.

Information Collected in Diagnostic Bundles

Cloudera Data Science Workbench diagnostic bundles collect the following information:

- System information such as hostnames, operating system, kernel modules and settings, available hardware, and system logs.
- Cloudera Data Science Workbench version, status information, configuration, and the results of install-time validation checks.
- Details about file systems, devices, and mounts in use.
- CDH cluster configuration, including information about Java, Kerberos, installed parcels, and CDH services such as Spark 2.
- Network configuration and status, including interfaces, routing configuration, and reachability.
- Status information for system services such as Docker, Kubernetes, NFS, and NTP.
- Listings for processes, open files, and network sockets.
- Reachability, configuration, and logs for Cloudera Data Science Workbench application components.
- Hashed Cloudera Data Science Workbench user names.
- Information about Cloudera Data Science Workbench sessions, including type, engine, ownership, termination status, and performance data.
**Cloudera Data Science Workbench Email Notifications**

**Required Role:** Site Administrator

Go to the **Admin > Settings** tab to specify an email address for outbound invitations and job notifications.

By default, all emails are sent from noreply@your-cdsw-domain. However, if your SMTP domain is different from the Cloudera Data Science Workbench domain, or it does not allow spoofing, you will need to explicitly specify the email address at the No Reply Email field.

Cloudera Data Science Workbench sends email notifications when you add collaborators to a project, share a project with a colleague, and for job status updates (email recipients are configured per-job). Emails are not sent when you create a new project. Email preferences cannot currently be configured at an individual user level.

**Managing License Keys for Cloudera Data Science Workbench**

Cloudera Data Science Workbench requires a Cloudera Enterprise license. To obtain a Cloudera Enterprise license, either fill in this form, or call 866-843-7207. Note that only one license key can be used at a time.

After an initial trial period of 60 days, you must upload a license key to continue to use Cloudera Data Science Workbench.

**Trial License**

Cloudera Data Science Workbench is fully functional during a 60-day, non-renewable trial period. The trial period starts when you create your first user.

If 60 days or fewer remain on the license, a badge in the lower left corner of the dashboard displays the number of days remaining. The initial trial period is 60 days, so the remaining days are always displayed during the trial period.

When a trial license expires, functionality is limited as follows:

- A warning banner notifies you that the license has expired and suggests you contact the site administrator or upload a license key.
- You cannot create new users, projects, sessions, or jobs.
- Existing users can log in and view their projects and files.
- You cannot run existing jobs.

At this point you can obtain a Cloudera Enterprise license and upload it to Cloudera Data Science Workbench using the steps described below. Cloudera Data Science Workbench will then go back to being fully functional.

**Cloudera Enterprise License**

When an Enterprise license expires, a warning banner displays, but all product features remain fully functional.

Contact Cloudera Support to receive an updated license.

**Uploading License Keys**

To upload the license key:

1. Go to **Admin > License**.
2. Click **Upload License**.
3. Select the license file to be uploaded and click **Upload**.
Cluster Management

This section describes some common tasks and guidelines related to cluster management for Cloudera Data Science Workbench.

For a basic overview of how Cloudera Data Science Workbench fits onto a Cloudera Manager and CDH cluster, refer to the Architecture Overview on page 16.

Backup and Disaster Recovery for Cloudera Data Science Workbench

All application data for Cloudera Data Science Workbench, including project files and database state, is stored on the master host at /var/lib/cdsw. Given typical access patterns, it is strongly recommended that /var/lib/cdsw be stored on a dedicated SSD block device or SSD RAID configuration. Because application data is not replicated to HDFS or backed up by default, site administrators must enable a backup strategy to meet any disaster recovery scenarios.

Cloudera strongly recommends both regular backups and backups before upgrades and is not responsible for any data loss.

Creating a Backup

1. **Cloudera Data Science Workbench 1.4.2 or lower**

   Do not stop or restart Cloudera Data Science Workbench without using the `cdsw_protect_stop_restart.sh` script. This is to help avoid the data loss issue detailed in TSB-346.

   Run the script on your master host and stop Cloudera Data Science Workbench (instructions below) only when instructed to do so by the script. Then proceed with step 2 of this process.

2. **Cloudera Data Science Workbench 1.4.3 or higher**

   Depending on your deployment, use one of the following sets of instructions to stop the application.

   To stop Cloudera Data Science Workbench:

   - **CSD** - Log in to Cloudera Manager. On the Home > Status tab, click to the right of the CDSW service and select Stop from the dropdown. Wait for the action to complete.
   - OR
   - **RPM** - Run the following command on the master host:

     ```bash
cdsw stop
     ```

3. To create the backup, run the following command on the master host:

   ```bash
tar cvzf cdsw.tar.gz /var/lib/cdsw/*
   ```

   **Note:** Using tar to create the backup preserves important file metadata such as file ownership. Other methods of copying/saving files might not preserve this information. This metadata is required for tasks such as migrating CDSW to another cluster.

4. (Optional) If needed, the following command can be used to unpack the tar bundle.

   ```bash
tar xvzf cdsw.tar.gz -C /var/lib/cdsw
   ```
Cloudera Data Science Workbench Scaling Guidelines

New hosts can be added and removed from a Cloudera Data Science Workbench deployment without interrupting any jobs already scheduled on existing hosts. Therefore, it is rather straightforward to increase capacity based on observed usage. At a minimum, Cloudera recommends you allocate at least 1 CPU core and 2 GB of RAM per concurrent session or job. CPU can burst above a 1 CPU core share when spare resources are available. Therefore, a 1 CPU core allocation is often adequate for light workloads. Allocating less than 2 GB of RAM can lead to out-of-memory errors for many applications.

As a general guideline, Cloudera recommends hosts with RAM between 60GB and 256GB, and between 16 and 48 cores. This provides a useful range of options for end users. SSDs are strongly recommended for application data storage.

For some data science and machine learning applications, users can collect a significant amount of data in memory within a single R or Python process, or use a significant amount of CPU resources that cannot be easily distributed into the CDH cluster. If individual users frequently run larger workloads or run workloads in parallel over long durations, increase the total resources accordingly. Understanding your users’ concurrent workload requirements or observing actual usage is the best approach to scaling Cloudera Data Science Workbench.

Ports Used By Cloudera Data Science Workbench

Cloudera Data Science Workbench runs on gateway hosts in a CDH cluster. As such, Cloudera Data Science Workbench acts as a gateway and requires full connectivity to CDH services (Impala, Spark 2, etc.) running on the cluster. Additionally, in the case of Spark 2, CDH cluster hosts will require access to the Spark driver running on a set of random ports (20050-32767) on Cloudera Data Science Workbench hosts.

Internally, the Cloudera Data Science Workbench master and worker hosts require full connectivity with no firewalls. Externally, end users connect to Cloudera Data Science Workbench exclusively through a web server running on the master host, and therefore do not need direct access to any other internal Cloudera Data Science Workbench or CDH services.

This information has been summarized in the following table.

<table>
<thead>
<tr>
<th>Components</th>
<th>Details</th>
</tr>
</thead>
<tbody>
<tr>
<td>Communication with the CDH cluster</td>
<td><strong>CDH -&gt; Cloudera Data Science Workbench</strong></td>
</tr>
<tr>
<td></td>
<td>The CDH cluster must have access to the Spark driver that runs on Cloudera Data Science Workbench hosts, on a set of randomized ports in the range, <strong>20050-32767</strong>.</td>
</tr>
<tr>
<td></td>
<td><strong>Cloudera Data Science Workbench -&gt; CDH</strong></td>
</tr>
<tr>
<td></td>
<td>As a gateway service, Cloudera Data Science Workbench must have access to all the ports used by CDH and Cloudera Manager.</td>
</tr>
<tr>
<td>Communication with the Web Browser</td>
<td>The Cloudera Data Science Workbench web application is available at port <strong>80</strong>. HTTPS access is available over port <strong>443</strong>.</td>
</tr>
</tbody>
</table>

Managing Cloudera Data Science Workbench Hosts

This topic describes how to perform some common tasks related to managing Cloudera Data Science Workbench hosts.

Migrating a Deployment to a New Set of Hosts

The following topics describe how to migrate a Cloudera Data Science Workbench deployment to a new set of gateway hosts.

- **Migrating a CSD Deployment** on page 219
Adding a Worker Host

**Note:** Worker hosts are not required for a fully-functional Cloudera Data Science Workbench deployment. For proof-of-concept deployments, you can deploy a 1-host cluster with just a Master host. The Master host can run user workloads just as a worker host can.

**Using Cloudera Manager**

Perform the following steps to add a new worker host to Cloudera Data Science Workbench.

1. Log in to the Cloudera Manager Admin Console.
2. **Add a new host to your cluster.** Make sure this is a gateway host and you are not running any services on this host.
3. Assign the HDFS, YARN, and Spark 2 gateway roles to the new host. For instructions, refer the Cloudera Manager documentation at Adding a Role Instance.

**Note:** If you are using Spark 2.2 (or higher), review JDK 8 Requirement for Spark 2.2 (or higher) on page 53 for any additional steps you might need to perform to configure JAVA_HOME on the new nodes.

4. Go to the Cloudera Data Science Workbench service.
5. Click the Instances tab.
6. Click **Add Role Instances**.
7. Assign the Worker and Docker Daemon roles to the new host. Click **Continue**.
8. Review your changes and click **Continue**. The wizard finishes by performing any actions necessary to add the new role instances. **Do not start the new roles at this point. You must run the Prepare Node command as described in the next steps before the roles are started.**
9. The new host must have the following packages installed on it.

```
nfs-utils
libseccomp
lvm2
bridge-utils
libtool-ltdl
iptables
rsync
policycoreutils-python
selinux-policy-base
selinux-policy-targeted
ntp
ebtables
bind-utils
mmap-ncat
openssl
e2fsprogs
redhat-lsb-core
socat
```

You must either manually install these packages now, or, allow Cloudera Manager to install them in the next step.

If you choose the latter, make sure that Cloudera Manager has the permission needed to install the required packages. To do so, go to the Cloudera Data Science Workbench service and click **Configuration.** Search for the **Install Required Packages** property and make sure it is enabled.

10. **Click Instances** and select the new host. From the list of available actions, select the **Prepare Node** command to install the required packages on the new node.
11. On the **Instances** page, select the new role instances and click **Actions for Selected > Start.**
Using Packages

On an RPM deployment, the procedure to add a worker host to an existing deployment is the same as that required when you first install Cloudera Data Science Workbench on a worker. For instructions, see Installing Cloudera Data Science Workbench on a Worker Host.

Removing a Worker Host

Using Cloudera Manager

Perform the following steps to remove a worker host from Cloudera Data Science Workbench.

1. Log into the Cloudera Manager Admin Console.
2. Click the Instances tab.
3. Select the Docker Daemon and Worker roles on the host to be removed from Cloudera Data Science Workbench.
4. Select Actions for Selected > Stop and click Stop to confirm the action. Click Close when the process is complete.
5. On the Instances page, re-select the Docker Daemon and Worker roles that were stopped in the previous step.
6. Select Actions for Selected > Delete and then click Delete to confirm the action.

Using Packages

To remove a worker host:

1. On the master host, run the following command to delete the worker host:

```
kubectl delete node <worker_host_domain_name>
```

2. Reset the worker host.

```
cdsw reset
```

Changing the Domain Name

Cloudera Data Science Workbench allows you to change the domain of the web console.

Using Cloudera Manager

1. Log into the Cloudera Manager Admin Console.
2. Go to the Cloudera Data Science Workbench service.
3. Click the Configuration tab.
4. Search for the Cloudera Data Science Workbench Domain property and modify the value to reflect the new domain.
5. Click Save Changes.
6. Restart the Cloudera Data Science Workbench service to have the changes go into effect.

Using Packages

1. Open /etc/cdsw/config/cdsw.conf and set the DOMAIN variable to the new domain name.

```
DOMAIN="cdsw.<your_new_domain>.com"
```

2. Run the following commands to have the new domain name go into effect.

```
cdsw reset
cdsw init
```
Migrating a Deployment to a New Set of Hosts

This section describes how to migrate a Cloudera Data Science Workbench deployment to a new set of gateway hosts.

Migrating a CSD Deployment

This section describes how to migrate a CSD-based Cloudera Data Science Workbench service to a new set of gateway hosts.

Add and Set Up the New Hosts

1. **Add new hosts to your cluster as needed.** Make sure they are gateway hosts that have been assigned gateway roles for HDFS, YARN, and Spark 2. Do not run any other services on these hosts.

2. Set up the new hosts as per the Cloudera Data Science Workbench hardware requirements listed [here](#).
   - **Disable Untrusted SSH Access** on page 60 on the new hosts.
   - **Configure Block Devices** on page 60 on the new hosts.

Stop the CDSW Service

**Important:** On Cloudera Data Science Workbench 1.4.0 (and lower), **do not** stop or restart Cloudera Data Science Workbench without using the `cdsw_protect_stop_restart.sh` script. This is to help avoid the data loss issue detailed in [TSB-346](#).

Use Cloudera Manager to stop all roles of the CDSW service.

1. Log into the Cloudera Manager Admin Console.
2. On the Home > Status tab, click to the right of the CDSW service and select **Stop** from the dropdown.
3. Confirm your choice on the next screen. When you see a **Finished** status, the action is complete.

Backup Application Data

In Cloudera Data Science Workbench all stateful data is stored on the master host at `/var/lib/cdsw`. Backup the contents of this directory before you begin the migration process.

To create the backup, run the following command on the master host:

```bash
tar cvzf cdsw.tar.gz /var/lib/cdsw/*
```

Delete CDSW Roles from Existing Hosts

1. Log into the Cloudera Manager Admin Console.
2. Go to the CDSW service.
3. Click the **Instances** tab.
4. Select all the role instances.
5. Select Actions for Selected > **Delete**. Click **Delete** to confirm the deletion.

Move Backup to the New Master

Copy the backup taken previously to the host that will be the new Cloudera Data Science Workbench master. Unpack the contents of the backup into `/var/lib/cdsw`.

```bash
tar xvzf cdsw.tar.gz -C /var/lib/cdsw
```
Update DNS Records for the New Master

Update your DNS records with the IP address for the new master host.

Add Role Instances for the New Hosts

1. Log into the Cloudera Manager Admin Console.
2. Go to the CDSW service.
3. Click the Instances tab.
4. Click Add Role Instances. Assign the Cloudera Data Science Workbench Master, Application, and Docker Daemon roles to the new master host. If you want to configure worker hosts, assign the Cloudera Data Science Workbench Worker and Docker Daemon roles to the new workers.

**Important:** Do not assign the Master and Worker roles to the same host. Even if you only have one host for Cloudera Data Science Workbench, the Master can automatically perform the functions of a Worker host as needed.

5. Click Continue. On the Review Changes page, review the configuration changes to be applied. The wizard finishes by performing any actions necessary to add the new role instances.

**Do not start the new roles at this point. You must run the Prepare Node command as described in the next step before the roles are started.**

Run the Prepare Node command on the New Hosts

The new hosts must have the following packages installed on it.

- nfs-utils
- libseccomp
- lvm2
- bridge-utils
- libtool-ltdl
- iptables
- rsync
- policycoreutils-python
- selinux-policy-base
- selinux-policy-targeted
- ntp
- ebtables
- bind-utils
- nmap-ncat
- openssl
- e2fsprogs
- redhat-lsb-core
- socat

You can either manually install these packages now, or, allow Cloudera Manager to install them as part of the Prepare Node command later in this step.

If you choose the latter, make sure that Cloudera Manager has the permissions needed to install the required packages. To do so, go to the CDSW service and click Configuration. Search for the Install Required Packages property and make sure it is enabled.

Then run the Prepare Node command on the new hosts.

1. Go to the CDSW service.
2. Click Instances.
3. Select all the role instances.
4. Select Actions for Selected > Prepare Node. This will install the required set of packages on all the new hosts.

Start the CDSW Service

1. Log into the Cloudera Manager Admin Console.
2. On the **Home > Status** tab, click to the right of the **CDSW** service and select **Start** from the dropdown.

3. Confirm your choice on the next screen. When you see a **Finished** status, the action is complete.

### Migrating an RPM Deployment

This section describes how to migrate an RPM-based Cloudera Data Science Workbench service to a new set of gateway hosts.

#### Add and Set Up the New Hosts

1. **Add new hosts to your cluster as needed.** Make sure they are gateway hosts that have been assigned gateway roles for HDFS, YARN, and Spark 2. Do not run any other services on these hosts.

2. Set up the new hosts as per the Cloudera Data Science Workbench hardware requirements listed [here](#).
   - Disable **Untrusted SSH Access** on page 60 on the new hosts.
   - Configure **Block Devices** on page 60 on the new hosts.

#### Stop Cloudera Data Science Workbench

**Important:** On Cloudera Data Science Workbench 1.4.0 (and lower), **do not** stop or restart Cloudera Data Science Workbench without using the `cdsw_protect_stop_restart.sh` script. This is to help avoid the data loss issue detailed in TSB-346.

Run the following command on the master host to stop Cloudera Data Science Workbench.

```
cdsw stop
```

#### Backup Application Data

In Cloudera Data Science Workbench all stateful data is stored on the master host at `/var/lib/cdsw`. Backup the contents of this directory before you begin the migration process.

To create the backup, run the following command on the **master** host:

```
tar cvzf cdsw.tar.gz /var/lib/cdsw/*
```

#### Remove Cloudera Data Science Workbench from Existing Hosts

Run the following commands on the existing master and any worker hosts you want to migrate.

```
cdsw reset
yum remove cloudera-data-science-workbench
```

#### Move Backup to New Master

Copy the backup taken in the previous step to the host that will be the new Cloudera Data Science Workbench master. Unpack the contents of the backup into `/var/lib/cdsw`.

```
tar xvzf cdsw.tar.gz -C /var/lib/cdsw
```

#### Update DNS Records for the New Master

Update your DNS records with the IP address for the new master host.
Install Cloudera Data Science Workbench on New Master Host

For instructions, see Installing Cloudera Data Science Workbench 1.5.x Using Packages on page 67.

Rollback Cloudera Data Science Workbench to an Older Version

All stateful data for Cloudera Data Science Workbench is stored in the /var/lib/cdsw directory on the Master host. The contents of this directory are forward compatible, which is what allows for upgrades. However, they are not backward compatible. Therefore, to rollback Cloudera Data Science Workbench to a previous version, you must have a backup of the /var/lib/cdsw directory, taken prior to the last upgrade.

In general, the steps required to restore a previous version of Cloudera Data Science Workbench are:

1. Depending on your deployment, either uninstall the RPM or deactivate the current CDSW parcel in Cloudera Manager.
2. On the master host, restore the backup copy you have of /var/lib/cdsw. Note that any changes made after this backup will be lost.
3. Install a version of Cloudera Data Science Workbench that is equal to or greater than the version of the /var/lib/cdsw backup.

Uninstalling Cloudera Data Science Workbench

This topic describes how to uninstall Cloudera Data Science Workbench from a cluster.

CSD Deployments

1. Log in to the Cloudera Manager Admin Console.
2. Important: On Cloudera Data Science Workbench 1.4.0 (and lower), do not stop or restart Cloudera Data Science Workbench without using the cdsw_potect_stop_restart.sh script. This is to help avoid the data loss issue detailed in TSB-346.

   On the Home > Status tab, click
   to the right of the CDSW service and select Stop from the dropdown and confirm that you want to stop the service.

3. (Strongly Recommended) On the master host, backup the contents of the /var/lib/cdsw directory. This is the directory that stores all your application data.

   To create the backup, run the following command on the master host:
   ```
   tar cvzf cdsw.tar.gz /var/lib/cdsw/*
   ```

4. Go back to Cloudera Manager and click Hosts > Parcels.
5. Go to the CDSW parcel and click Deactivate.
6. Select Deactivate Only from the list.
7. Click the
   to the right of the Activate button and select Remove From Hosts.
8. Click OK to confirm.
9. Go back to the Home > Status page and click
   to the right of the CDSW service. Select Delete from the dropdown and confirm that you want to delete the service.
10. Remove all your user data that is stored in /var/lib/cdsw on the master host from the deployment. This step will permanently remove all user data.

```bash
sudo rm -Rf /var/lib/cdsw
```

For more details on how to uninstall Cloudera Manager and it's components, see Uninstalling Cloudera Manager and Managed Software.

### RPM Deployments

1. **Important:** On Cloudera Data Science Workbench 1.4.0 (and lower), **do not** stop or restart Cloudera Data Science Workbench without using the `cdsw_protect_stop_restart.sh` script. This is to help avoid the data loss issue detailed in **TSB-346**.

   Run the following command on the master host to stop Cloudera Data Science Workbench.

   ```bash
cdsw stop
```

2. **(Strongly Recommended)** On the master host, backup the contents of the /var/lib/cdsw directory. This is the directory that stores all your application data.

   To create the backup, run the following command on the master host:

   ```bash
tar cvzf cdsw.tar.gz /var/lib/cdsw/*
```

3. To uninstall Cloudera Data Science Workbench, run the following commands on the master host and all the worker hosts.

   ```bash
cdsw reset
yum remove cloudera-data-science-workbench
```

4. Remove all your user data that is stored in /var/lib/cdsw on the master host from the deployment. This step will permanently remove all user data.

   ```bash
sudo rm -Rf /var/lib/cdsw
```
Cloudera Data Science Workbench Security Guide

This topic provides an overview of the Cloudera Data Science Workbench security model and describes how Cloudera Data Science Workbench leverages the security and governance capabilities of your Cloudera Enterprise cluster to deliver a secure, robust, collaborative data science platform for the enterprise.

Security Model

Cloudera Data Science Workbench uses Kubernetes to schedule and manage Docker containers. All Cloudera Data Science Workbench components (web application, PostgreSQL database, Livelog, and so on) execute inside Docker containers. These are represented on the left-hand side of the diagram. Similarly, the environments that users operate in (via sessions, jobs, experiments, models), also run within isolated Docker containers that we call engines.

This architecture allows Cloudera Data Science Workbench to leverage the isolation properties of Docker to achieve notable security benefits.

**Benefits of the Docker Isolation Model** - Docker containers share the underlying host operating system, but are isolated from the rest of the host and from each other. Each container gets its own:

- **Isolated File System**: The Docker container does not see the host file system, but instead sees only the filesystem provided by the container and any host volumes that you have explicitly mounted into the container. This means a user launching a Cloudera Data Science Workbench session will only have access to the project files, and any specific host volumes you have chosen to mount into the session engine. They will not otherwise have access to the underlying host filesystem.

- **Isolated Process Namespace**: Docker containers cannot affect any processes running either on the host operating system or in other containers. Cloudera Data Science Workbench creates a new container each time a session/job/experiment/model is launched. This means user workloads can run in complete isolation from each other.

For more details on the Docker security model, see [Docker Security Overview](#).
Wildcard DNS Subdomain Requirement

When you first set up Cloudera Data Science Workbench, you are asked to create a wildcard DNS entry for the Cloudera Data Science Workbench domain. Cloudera Data Science Workbench uses these wildcard subdomains (*.cdsw.<your_domain>.com) to route HTTP requests to engines and services launched by users.

Every time users launch workloads (session/job/experiment/model) on Cloudera Data Science Workbench, a new engine is created for each workload. These engines are isolated Docker containers where users can execute code. Each engine is assigned its own unique, randomly-generated ID, which is saved to the CDSW_ENGINE_ID variable. This ID is also used to create a unique subdomain for each engine. These subdomains are of the form: <CDSW_ENGINE_ID>.cdsw.<your_domain>.com.

Assigning a unique subdomain to each engine allows Cloudera Data Science Workbench to:

- Securely expose interactive session services, such as visualizations, the terminal, and web UIs such as TensorBoard, Shiny, Plotly, and so on;
- Prevent cross-site scripting (XSS) attacks by securely isolating user-generated content from the Cloudera Data Science Workbench application.

It is important to note that because there is no limit to the number of workloads (i.e. engines) users can launch, Cloudera Data Science Workbench requires the ability to randomly generate large numbers of engine IDs (and their subdomains) on-demand. Therefore, creating a wildcard DNS subdomain is essential for Cloudera Data Science Workbench to function successfully.

Additionally, if you want to enable TLS for your deployment, your TLS certificate will need to include both, the Cloudera Data Science Workbench domain, as well as the wildcard for all first-level subdomains. This is required so that your browser can trust communications with the <CDSW_ENGINE_ID>.cdsw.<your_domain>.com subdomains.

Authentication

Authentication occurs in various forms in the Cloudera Data Science Workbench application. These are:

- **User Login**: Occurs when users log in to the Cloudera Data Science Workbench Web UI and authenticate themselves to the application. Cloudera Data Science Workbench works with the following authentication back-ends: the local CDSW database, LDAP/AD, and SAML. Using either LDAP or SAML is recommended, as it eases the administrative burden of managing identity in Cloudera Data Science Workbench.

- **SSH Key Authentication**: Each user account is assigned an SSH key pair for use in sessions. This SSH key pair can be used to authenticate to an external version control system such as Git. Only the public key appears in the UI; the private key is loaded into user sessions when launched.

- **Hadoop Cluster Authentication**: Authentication to the underlying CDH/HDP cluster is handled via Kerberos. To authenticate themselves to the cluster, users must provide a Kerberos principal and keytab/password. These credentials are stored in the internal Cloudera Data Science Workbench database. Note that Cloudera Data Science Workbench user sessions are only provided with the Kerberos credential-cache file which does not store the user's password. For more details, see Hadoop Authentication with Kerberos for Cloudera Data Science Workbench on page 236.

- **API Authentication**: Each user account is assigned an API Key that can be used for authentication when communicating with the Cloudera Data Science Workbench API. For more details, see Cloudera Data Science Workbench Jobs API on page 165.

- **Per-Model Authentication**: Each model is assigned a unique Access Key. This access key serves as an authentication token that allows users to make calls to the model once it has been deployed. For details, see Model Access Key.
Authorization

This section describes how Cloudera Data Science Workbench handles authorization for users attempting to connect to the attached CDH/HDP cluster, and how Cloudera Data Science Workbench users and teams are granted access to projects.

Cluster Authorization

Cloudera Data Science Workbench connects to your cluster like any other client. It relies on your cluster’s built-in security layer to enforce strong authentication and authorization for connections from Cloudera Data Science Workbench.

Cloudera Data Science Workbench users attempting to connect to a secure CDH or HDP cluster must first authenticate themselves using their Kerberos credentials. Once a user has been successfully authenticated, Cloudera Data Science Workbench then depends on your cluster’s built-in authorization tools (such as HDFS ACLs, Sentry, Ranger, and so on) to enforce their own existing access control rules when a Cloudera Data Science Workbench user attempts to access data on the cluster.

For example, let’s assume you are running a secure CDH cluster with Impala (secured by Sentry). On this cluster, only the `impala-admin` group has been granted the required Sentry privileges to modify data in Impala. If a Cloudera Data Science Workbench user runs a Python job that attempts to modify data in an Impala table, Sentry privileges will be enforced. That is, if the user is not a member of the `impala-admin` group, the access request will be denied.

Similarly, if a Cloudera Data Science Workbench user is attempting to submit Spark jobs on secure HDP clusters, Ranger policies (via the respective HDFS and YARN plugins) will be enforced to decide whether the user has the permission to access the requested directories in HDFS, and whether they can submit jobs to the Spark queue.

This means as long as your cluster components have been secured with access control rules (via ACLs or Sentry or Ranger) that forbid specific users/groups from performing certain actions on the cluster, then all access attempts from Cloudera Data Science Workbench users will also be subject to those rules.

User Role Authorization

Cloudera Data Science Workbench has one major specialized user role across the deployment: site administrators. Site administrators are superusers who have complete access to all activity on your Cloudera Data Science Workbench deployment. This includes all the configuration settings, projects (private and public), and workloads running on the deployment.

When LDAP or SAML based authentication is used, it is possible to restrict the users who can log into Cloudera Data Science Workbench based on their LDAP/SAML group membership. Additionally, you can specify LDAP/SAML groups that should automatically be granted site administrator privileges when they log in to Cloudera Data Science Workbench.

Apart from site administrators, every other Cloudera Data Science Workbench user is granted access to projects either as a project collaborator or as part of a team. Continue reading to find out how these project and team permissions interact with each other.

Access Control for Teams and Projects

When a team or project is created, the Team/Project Admin role is assigned to the user who created it. Other Team/Project Admins can be assigned later but there must always be at least one user assigned as Admin for the team or project. Team and project administrators then decide what level of access other users are granted per-team or per-project.

Project Access Levels

Users who are explicitly added to a project are referred to as project collaborators. Project collaborators can be assigned one of the following levels of access:

- **Viewer** - Read-only access to code, data, and results.
- **Contributor** - Can view, edit, create, and delete files and environmental variables, run sessions/experiments/jobs/models and execute code in running jobs. Additionally, Contributors can set the default engine for the project.
- **Admin** - Has complete access to all aspects of the project. This includes the ability to add new collaborators, and delete the entire project.

**Team Access Levels**

Users who are explicitly added to a team are referred to as team members. Team members can be assigned one of the following levels of access:

- **Viewer** - Read-only access to team projects. Cannot create new projects within the team but can be added to existing ones.
- **Contributor** - Write-level access to all team projects with Team or Public visibility. Can create new projects within the team. They can also be added to existing team projects.
- **Admin** - Has complete access to all team projects, can add new team members, and modify team account information.

**Project Visibility Levels**

Projects can be created either in your personal context, or in a team context. Furthermore, projects can be created with one of the following visibility levels:

- **Private** - Private projects can be created either in your personal context, or in a team context. They can only be accessed by project collaborators.
- **Team** - Team projects can only be created in a team context. They can be viewed by all members of the team.
- **Public** - Public projects can be created either in your personal context, or in a team context. They can be viewed by all authenticated Cloudera Data Science Workbench users.

It is important to remember that irrespective of the visibility level of the project, site administrators will always have complete Admin-level access to all projects on Cloudera Data Science Workbench. Additionally, depending on the visibility level of the project, and the context in which it was created, a few other users/team members might also have Contributor or Admin-level access to your project by default.

Use the following table to find out who might have default access to your projects on Cloudera Data Science Workbench.

<table>
<thead>
<tr>
<th>Project Visibility</th>
<th>Access Levels for Cloudera Data Science Workbench Users</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Private Visibility</strong></td>
<td><strong>Private Projects Created in Personal Context</strong></td>
</tr>
<tr>
<td></td>
<td>The following user roles will have access to private projects in your personal context:</td>
</tr>
<tr>
<td></td>
<td><strong>Admin Access</strong></td>
</tr>
<tr>
<td></td>
<td>• Site Administrators</td>
</tr>
<tr>
<td></td>
<td>• Project Admins</td>
</tr>
<tr>
<td></td>
<td><strong>Contributor Access</strong></td>
</tr>
<tr>
<td></td>
<td>• Collaborators explicitly added to the project and given Contributor access.</td>
</tr>
<tr>
<td></td>
<td><strong>Viewer Access</strong></td>
</tr>
<tr>
<td></td>
<td>• Viewers explicitly added to the project and given Viewer access.</td>
</tr>
<tr>
<td></td>
<td><strong>Private Projects Created in a Team Context</strong></td>
</tr>
<tr>
<td></td>
<td>For private projects created within a team context, project-level permissions granted by Project Admins will take precedence over team-level permissions. The only exception to this rule are users who are Team Admins. Team Admins will always have Admin-level access to all projects within their team context, irrespective of the access level granted to them per-project.</td>
</tr>
</tbody>
</table>
| | The following user roles will have access to private projects created within a team context:
### Project Visibility

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<td><strong>Admin Access</strong></td>
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<tr>
<td>• Site Administrators</td>
</tr>
<tr>
<td>• Project Admins</td>
</tr>
<tr>
<td>• Team Admins</td>
</tr>
<tr>
<td><strong>Contributor Access</strong></td>
</tr>
<tr>
<td>• Collaborators explicitly added to the project and given Contributor access.</td>
</tr>
<tr>
<td><strong>Viewer Access</strong></td>
</tr>
<tr>
<td>• Viewers explicitly added to the project and given Viewer access.</td>
</tr>
</tbody>
</table>

### Team Visibility

<table>
<thead>
<tr>
<th>Team Projects</th>
</tr>
</thead>
</table>

Projects with Team visibility can only be created in a team context. For team projects, both team access levels and project access levels must be taken into consideration to determine who has access to these projects.

**Points to note:**

- Team members do not need to be explicitly added as project collaborators to have access to a team project. While you can explicitly invite specific team members to collaborate on your project, it is important to remember that all team members will have some level of access to your project.
  - The project Collaborators page does not list all team members; it only lists those you have explicitly added as collaborators. However, team members will still have access to all team projects. By default, their level of access to the projects is the same as their level of access to the team.
  - Project Admins cannot downgrade access levels for team members. If project-level permissions don’t match up to team-level permissions, team permissions will take precedence.
    - For example, if you add a Team Contributor as a collaborator to a team project, but only give them Project Viewer permission, the user will still have Contributor-level access to the project. Similarly, Team Admins will always have Admin-level access to all projects within their team context, irrespective of the access level granted to them per-project.
  - Project Admins also cannot upgrade access levels for team members with Viewer-level access to the team. That is, Team Viewers cannot be given Contributor or Admin access to any team projects.

The following user roles will have access to team projects on Cloudera Data Science Workbench:

<table>
<thead>
<tr>
<th><strong>Admin Access</strong></th>
</tr>
</thead>
<tbody>
<tr>
<td>• Site Administrators</td>
</tr>
<tr>
<td>• Team Admins</td>
</tr>
<tr>
<td>• Project Admins</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th><strong>Contributor Access</strong></th>
</tr>
</thead>
<tbody>
<tr>
<td>• All team members with Contributor access.</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th><strong>Viewer Access</strong></th>
</tr>
</thead>
<tbody>
<tr>
<td>• All team members with Viewer access.</td>
</tr>
</tbody>
</table>

### Public Visibility

<table>
<thead>
<tr>
<th>Public Projects Created in Personal Context</th>
</tr>
</thead>
</table>

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<table>
<thead>
<tr>
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<th>Access Levels for Cloudera Data Science Workbench Users</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>The following user roles will have access to public projects on Cloudera Data Science Workbench:</td>
</tr>
<tr>
<td></td>
<td><strong>Admin Access</strong></td>
</tr>
<tr>
<td></td>
<td>• Site Administrators</td>
</tr>
<tr>
<td></td>
<td>• Project Admins</td>
</tr>
<tr>
<td></td>
<td><strong>Contributor Access</strong></td>
</tr>
<tr>
<td></td>
<td>• Collaborators explicitly added to the project and given Contributor access.</td>
</tr>
<tr>
<td></td>
<td><strong>Viewer Access</strong></td>
</tr>
<tr>
<td></td>
<td>• All authenticated CDSW users.</td>
</tr>
</tbody>
</table>

**Public Projects Created in a Team Context**

The following user roles will have access to public projects created in team contexts:

**Admin Access**

• Site Administrators
• Project Admins
• Team Admins

**Contributor Access**

• All team members with Contributor access.

The team/project access rules and nuances described in the Team section apply here as well.

**Viewer Access**

• All authenticated CDSW users.

---

**Note: Restricting Access to Active Sessions**

Users with Admin or Contributor-level permissions on projects have access to all of the project’s active sessions and can execute commands within these active sessions. Cloudera Data Science Workbench (1.4.3 and higher) includes a feature that allows site administrators to restrict this ability by allowing only session creators to execute commands within their own active sessions. For details on how to enable this, see Restricting Access to Active Sessions.

---

**Wire Encryption**

**External Communications**

Cloudera Data Science Workbench uses HTTP and WebSockets (WS) to support interactive connections to the Cloudera Data Science Workbench web application. However, these connections are not secure by default.

For secure, encrypted communication, Cloudera Data Science Workbench can be configured to use a TLS termination proxy to handle incoming connection requests. The termination proxy server will decrypt incoming connection requests and forward them to the Cloudera Data Science Workbench web application.

The Cloudera Data Science Workbench documentation describes two different approaches to TLS termination: internal and external TLS termination. Both provide a secure TLS connection between users and Cloudera Data Science Workbench. If you require more control over the TLS protocol and cipher suite, we recommend external termination. Both approaches require TLS certificates that list both, the Cloudera Data Science Workbench domain, as well as a
wildcard for all first-level subdomains. For example, if the Cloudera Data Science Workbench domain is cdsw.<your_domain>.com, then the TLS certificate must include both cdsw.<your_domain>.com and *.cdsw.<your_domain>.com.

**Browser Security**

Cloudera Data Science Workbench also allows you to customize the HTTP headers accepted by Cloudera Data Science Workbench. The list of security headers enabled by default can be found in the documentation here: [HTTP Headers](#). Disabling these features could leave your Cloudera Data Science Workbench deployment vulnerable to clickjacking, cross-site scripting (XSS), or any other injection attacks.

**Internal Communications**

Internal communications between some Cloudera Data Science Workbench components are protected by mutually authenticated TLS.

The underlying Kubernetes cluster has a root Certificate Authority (CA) that is used to validate certificates for internal Kubernetes components. For details, refer the Kubernetes documentation here: [TLS certificates](#).

**Cloudera Data Science Workbench Gateway Host Security**

The [Cloudera Data Science Workbench master host](#) stores all the critical, stateful, persistent data for a Cloudera Data Science Workbench deployment. This data includes your deployment secrets, such as, Kerberos credentials, encrypted passwords, SSH and API keys, and so on. While the Cloudera Data Science Workbench worker hosts do not store the same secrets, they also store sensitive information. Therefore, protecting the master and worker hosts is extremely important. Cloudera recommends the following security best practices for all the Cloudera Data Science Workbench hosts:

- Disable untrusted SSH access to the Cloudera Data Science Workbench hosts. Cloudera Data Science Workbench assumes that users only access the gateway hosts through the web application. Users with SSH access to a Cloudera Data Science Workbench host can gain full access to the cluster, including access to other users' workloads. Therefore, untrusted (non-sudo) SSH access to Cloudera Data Science Workbench hosts must be disabled to ensure a secure deployment.

- Tightly control root access via sudo.

- Uninstall or disable any other unnecessary services running on the Cloudera Data Science Workbench gateway hosts.

- Keep the hosts' operating system updated to avoid security vulnerabilities.

- Monitor user login activity on the system.

**Host Mounts**

Cloudera Data Science Workbench allows site administrators to expose part of the host's file system into users' engine containers at runtime. This is done using the [host mounts](#) feature. It is worth noting that directories mounted using this feature are then available to all projects across the deployment. Use this feature with great care and ensure that no sensitive information is mounted.

**Base Engine Image Security**

The base engine image is a Docker image that contains all the building blocks needed to launch a Cloudera Data Science Workbench session and run a workload. It consists of kernels for Python, R, and Scala, and some common third-party libraries and packages that can be used to run common data analytics operations. Additionally, to provide a flexible, collaborative environment, data science teams can install their own preferred data science packages, just as they would on their local computers, to run workloads on data in the associated Hadoop cluster.
Enabling TLS/SSL for Cloudera Data Science Workbench

Cloudera Data Science Workbench uses HTTP and WebSockets (WS) to support interactive connections to the Cloudera Data Science Workbench web application. However, these connections are not secure by default. This topic describes how you can use TLS/SSL to enforce secure encrypted connections, using HTTPS and WSS (WebSockets over TLS), to the Cloudera Data Science Workbench web application.

Specifically, Cloudera Data Science Workbench can be configured to use a TLS termination proxy to handle incoming connection requests. The termination proxy server will decrypt incoming connection requests and forward them to the Cloudera Data Science Workbench web application. A TLS termination proxy can be internal or external.

Internal Termination

An internal termination proxy will be run by Cloudera Data Science Workbench's built-in load balancer, called the ingress controller, on the master host. The ingress controller is primarily responsible for routing traffic and load balancing between Cloudera Data Science Workbench's web service backend. Once configured, as shown in the instructions that follow, it will start terminating HTTPS traffic as well. The primary advantage of internal termination approach is simplicity.

External Termination

External TLS termination can be provided through a number of different approaches. Common examples include:

- Load balancers, such as the AWS Elastic Load Balancer
- Modern firewalls
- Reverse web proxies, such as nginx
- VPN appliances supporting TLS/SSL VPN

Organizations that require external termination will often have standardized on single approach for TLS. The primary advantage of this approach is that it allows such organizations to integrate with Cloudera Data Science Workbench without violating their IT department's policies for TLS. For example, with an external termination proxy, Cloudera Data Science Workbench does not need access to the TLS private key.

Load balancers and proxies often require a URL they can ping to validate the status of the web service backend. For instance, you can configure a load balancer to send an HTTP GET request to /internal/load-balancer/health-ping. If the response is 200 (OK), that means the backend is healthy. Note that, as with all communication to the web backend from the load balancer when TLS is terminated externally, this request should be sent over HTTP and not HTTPS.

Note that any terminating load balancer must provide the following header fields so that Cloudera Data Science Workbench can detect the IP address and protocol used by the client:

- X-Forwarded-For (client's IP address),
- X-Forwarded-Proto (client's requested protocol, i.e. HTTPS),
- X-Forwarded-Host (the "Host" header of the client's original request).

See Configuring HTTP Headers for Cloudera Data Science Workbench on page 244 for more details on how to customize HTTP headers required by Cloudera Data Science Workbench.

Related topic: Troubleshooting TLS/SSL Errors on page 251
Private Key and Certificate Requirements

The TLS certificate issued by your CA must list both, the Cloudera Data Science Workbench, as well as a wildcard for all first-level subdomains. For example, if the Cloudera Data Science Workbench domain is `cdsw.company.com`, then the TLS certificate must include both `cdsw.company.com` and `*.cdsw.company.com`.

Creating a Certificate Signing Request (CSR) and Key/Certificate Pair

Use the following steps to create a Certificate Signing Request (CSR) to submit to your CA. Then, create a private key/certificate pair that can be used to authenticate incoming communication requests to Cloudera Data Science Workbench.

**Important:** Make sure you use `openssl`, and not `keytool`, to perform these steps. Keytool does not support a wildcard Subject Alternative Name (SAN) and cannot create flat files.

1. Create a `cdsw.cnf` file and populate it with the required configuration parameters including the SAN field values.

   ```
   vi cdsw.cnf
   ```


3. Modify the following sections and save the `cdsw.cnf` file:

   ```
   [ CA_default ]
   default_md = sha2
   
   [ req ]
   default_bits = 2048
   distinguished_name = req_distinguished_name
   req_extensions = req_ext
   
   [ req_distinguished_name ]
   countryName = Country Name (2 letter code)
   stateOrProvinceName = State or Province Name (full name)
   localityName = Locality Name (eg, city)
   organizationName = Organization Name (eg, company)
   commonName = Common Name (e.g. server FQDN or YOUR name)
   
   [ req_ext ]
   subjectAltName = @alt_names
   
   [alt_names]
   DNS.1 = *.cdsw.company.com
   DNS.2 = cdsw.company.com
   ```

   **Key points to note:**
   
   - The domains set in the `DNS.1` and `DNS.2` entries above must match the DOMAIN set in `cdsw.conf`.
   - The `default_md` parameter must be set to `sha256` at a minimum. Older hash functions such as SHA1 are deprecated and will be rejected by browsers, either currently or in the very near future.
   - The `commonName` (CN) parameter will be ignored by browsers. You must use Subject Alternative Names.

4. Run the following command to generate the CSR.

   ```
   openssl req -out cert.csr -newkey rsa:2048 -nodes -keyout private.key -config cdsw.cnf
   ```

   This command generates the private key and the CSR in one step. The `-nodes` switch disables encryption of the private key (which is not supported by Cloudera Data Science Workbench at this time).

5. Use the CSR and private key generated in the previous step to request a certificate from the CA. If you have access to your organization’s internal CA or PKI, use the following command to request the certificate. If you do not have
access, or are using a third-party/commercial CA, use your organization’s respective internal process to submit the request.

```bash
openssl x509 -req -days 365 -in cert.csr -CA ca.crt -CAkey ca.key -CAcreateserial -out <your_tls_cert>.crt -sha256 -extfile cdsw.cnf -extensions req_ext
```

6. Run the following command to verify that the certificate issued by the CA lists both the required domains, `cdsw.company.com` and `*.cdsw.company.com`, under X509v3 Subject Alternative Name.

```bash
openssl x509 -in <your_tls_cert>.crt -noout -text
```

You should also verify that a valid hash function is being used to create the certificate. For SHA-256, the value under Signature Algorithm will be `sha256WithRSAEncryption`.

Configuring Internal Termination

Depending on your deployment (CSD or RPM), use one of the following sets of instructions to configure internal termination.

**CSD Deployments**

To enable internal termination, configure the following properties in the CDSW service in Cloudera Manager.

1. Log in to the Cloudera Manager Admin Console.
2. Navigate to the CDSW service and click **Configuration**.
3. Search for the following properties and configure as required.
   - **Enable TLS** - When enabled, this property enforces HTTPS and WSS connections. The server will now redirect any HTTP request to HTTPS and generate URLs with the appropriate protocol.
   - **TLS Key for Internal Termination** - Set to the path of the TLS private key.
   - **TLS Certificate for Internal Termination** - Set to the path of the TLS certificate.
   
   Certificates and keys must be in PEM format.

4. Click **Save Changes**.
5. Restart the CDSW service.

**RPM Deployments**

To enable internal termination, configure the following properties in `cdsw.conf` (on all Cloudera Data Science Workbench hosts).

- **TLS_ENABLE** - When set to `true`, this property enforces HTTPS and WSS connections. The server will now redirect any HTTP request to HTTPS and generate URLs with the appropriate protocol.
- **TLS_KEY** - Set to the path of the TLS private key.
- **TLS_CERT** - Set to the path of the TLS certificate.

Certificates and keys must be in PEM format.

You can configure these properties either as part of the installation process or after the fact. If you make any changes to `cdsw.conf` after installation is complete, make sure to **restart the master and worker hosts as needed**.

Configuring External Termination

Depending on your deployment (CSD or RPM), use one of the following sets of instructions to configure external termination.

**CSD Deployments**

To enable external termination, configure the following property in the CDSW service in Cloudera Manager.
1. Log in to the Cloudera Manager Admin Console.
2. Navigate to the CDSW service and click Configuration.
3. Search for the following properties and configure as required.
   - **Enable TLS** - When enabled, this property enforces HTTPS and WSS connections. The server will now redirect any HTTP request to HTTPS and generate URLs with the appropriate protocol.
     
     The **TLS Key for Internal Termination** and **TLS Certificate for Internal Termination** properties must be left blank.

4. Click Save Changes.
5. Restart the CDSW service.

**RPM Deployments**

To enable external termination, configure the following property in `cdsw.conf` (on all Cloudera Data Science Workbench hosts).

- **TLS_ENABLE** - When set to true, this property enforces HTTPS and WSS connections. The server will now redirect any HTTP request to HTTPS and generate URLs with the appropriate protocol.

  The **TLS_KEY** and **TLS_CERT** properties must be left blank.

You can configure this property either as part of the installation process or after the fact. If you make any changes to `cdsw.conf` after installation is complete, make sure to restart the master and worker hosts as needed.

**Known Issues and Limitations**

- Communication within the Cloudera Data Science Workbench cluster is not encrypted.

- Cloudera Data Science Workbench does not support encrypted private keys with internal TLS termination. If you require an encrypted private key, use external TLS termination with a terminating proxy that does support encrypted private keys.

- Troubleshooting can be difficult because browsers do not typically display helpful security errors with WebSockets. Often they will just silently fail to connect.

- **Self-signed certificates**

  In general, browsers do not support self-signed certificates for WSS. Your certificate must be signed by a Certificate Authority (CA) that your users’ browsers will trust. Cloudera Data Science Workbench will not function properly if browsers silently abort WebSockets connections.

  If you are using a TLS certificate that has been used to sign itself, and is not signed by a CA in the trust store, then the browser will display a dialog asking if you want to trust the certificate provided by Cloudera Data Science Workbench. This means you are using a self-signed certificate, which is **not supported and will not work**. In this case WSS connections will likely be aborted silently, regardless of your response (Ignore/Accept) to the dialog.

  As long as you have a TLS certificate signed by a CA certificate in the trust store, it will be supported and will work with Cloudera Data Science Workbench. For example, if you need to use a certificate signed by your organization’s internal CA, make sure that all your users import your root CA certificate into their machine’s trust store. This can be done using the Keychain Access application on Macs or the Microsoft Management Console on Windows.
Configuring Cloudera Data Science Workbench Deployments Behind a Proxy

If your deployment is behind an HTTP or HTTPS proxy, you must configure the hostname of the proxy you are using in Cloudera Data Science Workbench as follows.

```
HTTP_PROXY="<http://proxy_host>:<proxy-port>"
HTTPS_PROXY="<http://proxy_host>:<proxy-port>"
```

Depending on your deployment, use one of the following methods to configure the proxy in Cloudera Science Workbench:

- **CSD** - Set the **HTTP Proxy** or **HTTPS Proxy** properties in the Cloudera Manager's CDSW service.
- **RPM** - Set the **HTTP_PROXY** or **HTTPS_PROXY** properties in `/etc/cdsw/config/cdsw.conf` on all Cloudera Data Science Workbench gateway hosts.

**Intermediate Proxy:** If you are using an intermediate proxy such as Cntlm to handle NTLM authentication, add the Cntlm proxy address to these fields.

```
HTTP_PROXY="http://localhost:3128"
HTTPS_PROXY="http://localhost:3128"
```

**Supporting a TLS-Enabled Proxy Server:**

If the proxy server uses TLS encryption to handle connection requests, you will need to add the proxy’s root CA certificate to your host’s store of trusted certificates. This is because proxy servers typically sign their server certificate with their own root certificate. Therefore, any connection attempts will fail until the Cloudera Data Science Workbench host trusts the proxy’s root CA certificate. If you do not have access to your proxy’s root certificate, contact your Network / IT administrator.

To enable trust, perform the following steps on the master and worker hosts.

1. Copy the proxy’s root certificate to the trusted CA certificate store (`ca-trust`) on the Cloudera Data Science Workbench host.

   ```
   cp /tmp/<proxy-root-certificate>.crt /etc/pki/ca-trust/source/anchors/
   ```

2. Use the following command to rebuild the trusted certificate store.

   ```
   update-ca-trust extract
   ```

3. If you will be using custom engine images that will be pulled from a Docker repository, add the proxy’s root certificates to a directory under `/etc/docker/certs.d`. For example, if your Docker repository is at `docker.repository.mycompany.com`, create the following directory structure:

   ```
   /etc/docker/certs.d
   |-- docker.repository.mycompany.com  # Directory named after Docker repository
   |   |-- <proxy-root-certificate>.crt  # Docker-related root CA certificates
   ```

   This step is not required with the standard engine images because they are included in the Cloudera Data Science Workbench RPM.

4. Re-initialize Cloudera Data Science Workbench to have this change go into effect.

   ```
   cdsw init
   ```
Configure Hostnames to be Skipped from the Proxy

Starting with version 1.4, if you have defined a proxy in the `HTTP_PROXY(S)` or `ALL_PROXY` properties, Cloudera Data Science Workbench automatically appends the following list of IP addresses to the `NO_PROXY` configuration. Note that this is the minimum required configuration for this field.

```
*127.0.0.1,localhost,100.66.0.1,100.66.0.2,100.66.0.3,  
100.66.0.4,100.66.0.5,100.66.0.6,100.66.0.7,100.66.0.8,100.66.0.9,  
100.66.0.10,100.66.0.11,100.66.0.12,100.66.0.13,100.66.0.14,  
100.66.0.15,100.66.0.16,100.66.0.17,100.66.0.18,100.66.0.19,  
100.66.0.20,100.66.0.21,100.66.0.22,100.66.0.23,100.66.0.24,  
100.66.0.25,100.66.0.26,100.66.0.27,100.66.0.28,100.66.0.29,  
100.66.0.30,100.66.0.31,100.66.0.32,100.66.0.33,100.66.0.34,  
100.66.0.35,100.66.0.36,100.66.0.37,100.66.0.38,100.66.0.39,  
100.66.0.40,100.66.0.41,100.66.0.42,100.66.0.43,100.66.0.44,  
100.66.0.45,100.66.0.46,100.66.0.47,100.66.0.48,100.66.0.49,  
100.66.0.50,100.77.0.10,100.77.0.128,100.77.0.129,100.77.0.130,  
100.77.0.131,100.77.0.132,100.77.0.133,100.77.0.134,100.77.0.135,  
100.77.0.136,100.77.0.137,100.77.0.138,100.77.0.139*
```

This list includes `127.0.0.1`, `localhost`, and any private Docker registries and HTTP services inside the firewall that Cloudera Data Science Workbench users might want to access from the engines.

To configure any additional hostnames that should be skipped from the proxy, use one of the following methods depending on your deployment:

- **On a CSD deployment**, use the Cloudera Manager CDSW service's `No Proxy` property to specify a comma-separated list of hostnames.
- **On an RPM deployment**, configure the `NO_PROXY` field in `cdsw.conf` on all Cloudera Data Science Workbench hosts.

Hadoop Authentication with Kerberos for Cloudera Data Science Workbench

Cloudera Data Science Workbench users can authenticate themselves using Kerberos against the cluster KDC defined in the host’s `/etc/krb5.conf` file. Cloudera Data Science Workbench does not assume that your Kerberos principal is always the same as your login information. Therefore, you will need to make sure Cloudera Data Science Workbench knows your Kerberos identity when you sign in.

To authenticate against your cluster’s Kerberos KDC, go to the top-right dropdown menu, click **Account settings > Hadoop Authentication**, and enter your Kerberos principal. To authenticate, either enter your password or click **Upload Keytab** to upload the keytab file directly to Cloudera Data Science Workbench. Once successfully authenticated, Cloudera Data Science Workbench uses your stored credentials to ensure that you are secure when running your workloads.

When you authenticate with Kerberos, Cloudera Data Science Workbench will store your keytab in an internal database. When you subsequently run an engine, the keytab is used by a Cloudera Data Science Workbench sidecar container to generate ticket-granting tickets for use by your code. Ticket-granting tickets allow you to access resources such as Spark, Hive, and Impala, on Kerberized CDH clusters.

While you can view your current ticket-granting ticket by typing `klist` in an engine terminal, there is no way for you or your code to view your keytab. This prevents malicious code and users from stealing your keytab.
Important:

- If the `/etc/krb5.conf` file is not available on all Cloudera Data Science Workbench hosts, authentication will fail.
- If you do not see the **Hadoop Authentication** tab, make sure you are accessing your personal account’s settings from the top right menu. If you have selected a team account, the tab will not be visible when accessing the Team Settings from the left sidebar.
- When you upload a Kerberos keytab to authenticate yourself to the CDH cluster, Cloudera Data Science Workbench might display a fleeting error message (‘cancelled’) in the bottom right corner of the screen, even if authentication was successful. This error message can be ignored.

**UI Behavior for Non-Kerberized Clusters**

The contents of the **Hadoop Authentication** tab change depending on whether the cluster is kerberized. For a secure cluster with Kerberos enabled, the **Hadoop Authentication** tab displays a **Kerberos** section with fields to enter your Kerberos principal and username. However, if Cloudera Data Science Workbench cannot detect a `krb5.conf` file on the host, it will assume the cluster is not kerberized, and the **Hadoop Authentication** tab will display **Hadoop Username Override** configuration instead.

For a non-kerberized cluster, by default, your Hadoop username will be set to your Cloudera Data Science Workbench username. To override this default and set an alternative `HADOOP_USER_NAME`, go to the **Hadoop Username Override** setting at **Account settings > Hadoop Authentication**.

**Important:** In version 1.4.0, the Hadoop username on non-kerberized clusters defaults to `cdsw` - not your Cloudera Data Science Workbench username. This is a known issue and has been fixed in version 1.4.2 (and higher).

If the **Hadoop Authentication** tab is incorrectly displaying Kerberos configuration fields for a non-kerberized cluster, make sure the `krb5.conf` file is not present on the host running Cloudera Data Science Workbench. If you do find any instances of `krb5.conf` on the host, depending on your deployment, perform one of the following sets of actions:

- On CSD deployments, go to Cloudera Manager and stop the CDSW service. Remove the `krb5.conf` file(s) from the Cloudera Data Science Workbench gateway host, and then start the CDSW in Cloudera Manager.

  OR

- On RPM deployments, run `cdsw stop`, remove the `krb5.conf` file(s) from the Cloudera Data Science Workbench gateway host, and run `cdsw start`.

You should now see the expected **Hadoop Username Override** configuration field.

**Limitations**

- Cloudera Data Science Workbench only supports Active Directory and MIT KDCs. PowerBroker-equipped Active Directory is not supported.
- Cloudera Data Science Workbench does not support the use of Kerberos plugin modules in `krb5.conf`.

**Configuring External Authentication with LDAP and SAML**

Cloudera Data Science Workbench supports user authentication against its internal local database, and against external services such as Active Directory, OpenLDAP-compatible directory services, and SAML 2.0 Identity Providers. By default, Cloudera Data Science Workbench performs user authentication against its internal local database. This topic describes the signup process for the first user, how to configure authentication using LDAP, Active Directory or SAML 2.0, and
an optional workaround that allows site administrators to bypass external authentication by logging in using the local database in case of misconfiguration.

User Signup Process

The first time you visit the Cloudera Data Science Workbench web console, the first account that you sign up with is a local administrator account. If in the future you intend to use external services for authentication, Cloudera recommends you use exclusive username & email combinations, rather than site administrators’ work email addresses, for both the first site administrator account, and any other local accounts created before switching to external authentication. If the username/email combinations are not unique, an email address might end up being associated with different usernames, one for the external authentication service provider and one for a local Cloudera Data Science Workbench account. This will prevent the user from logging into Cloudera Data Science Workbench with their credentials for the external authentication service.

The link to the signup page is only visible on the login page when the authentication type is set to local. When you enable external services for authentication, signing up through the local database is disabled, and user accounts are automatically created upon their first successful login.

Optionally, site administrators can use a Require invitation to sign up flag under the Admin > Settings tab to require invitation tokens for account creation. When this flag is enabled, only users that are invited by site administrators can login to create an account, regardless of the authentication type.

Important: If you forget the original credentials, or make a mistake with LDAP or SAML configuration, you can use the workaround described in Debug Login URL on page 244.

When you switch to using external authentication methods such as LDAP or SAML 2.0, user accounts will be automatically created upon each user’s first successful login. Cloudera Data Science Workbench will extract user attributes such as username, email address and full name from the authentication responses received from the LDAP server or SAML 2.0 Identity Provider and use them to create the user accounts.

Configuring LDAP/Active Directory Authentication

Cloudera Data Science Workbench supports both search bind and direct bind operations to authenticate against an LDAP or Active Directory directory service. The search bind authentication mechanism performs an ldapsearch against the directory service, and binds using the found Distinguished Name (DN) and password provided. The direct bind authentication mechanism binds to the LDAP server using a username and password provided at login.

You can configure Cloudera Data Science Workbench to use external authentication methods by clicking the Admin link on the left sidebar and selecting the Security tab. Select LDAP from the list to start configuring LDAP properties.

LDAP General Settings

- **LDAP Server URI**: Required. The URI of the LDAP/Active Directory server against which Cloudera Data Science Workbench should authenticate. For example, ldaps://ldap.company.com:636.
- **Use Direct Bind**: If checked, the username and password provided at login are used with the LDAP Username Pattern for binding to the LDAP server. If unchecked, Cloudera Data Science Workbench uses the search bind mechanism and two configurations, LDAP Bind DN and LDAP Bind Password, are required to perform the ldapsearch against the LDAP server.
- **LDAP Bind DN**: Required when using search bind. The DN to bind to for performing ldapsearch. For example, cn=admin,dc=company,dc=com.
- **LDAP Bind Password**: Required when using search bind. This is the password for the LDAP Bind DN.
- **LDAP Search Base**: Required. The base DN from which to search for the provided LDAP credentials. For example, ou=Engineering,dc=company,dc=com.
- **LDAP User Filter**: Required. The LDAP filter for searching for users. For example, 
  
  ((&(sAMAccountName={0})(objectclass=person))). The {0} placeholder will be replaced with the username provided at login.
• **LDAP User Username Attribute**: Required. The case-sensitive username attribute of the LDAP directory service. This is used by Cloudera Data Science Workbench to perform the bind operation and extract the username from the response. Common values are `uid`, `sAMAccountName`, or `userPrincipalName`.

<table>
<thead>
<tr>
<th>General Settings</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>LDAP Server URI</strong> *</td>
</tr>
<tr>
<td>ldaps://ldap.company.com</td>
</tr>
<tr>
<td><strong>Use Direct Bind</strong></td>
</tr>
<tr>
<td>By default, Cloudera Data Science Workbench searches for the users by binding to the provided <strong>LDAP Bind DN</strong> and <strong>LDAP Bind Password</strong>. When checked, Cloudera Data Science Workbench will attempt to search for the user by binding to the user-provided username and password. In such case, please make sure the user has permissions to search for themselves in the LDAP server.</td>
</tr>
<tr>
<td><strong>LDAP Bind DN</strong> *</td>
</tr>
<tr>
<td>CN=Test1 Person,OU=People,DC=company,DC=com</td>
</tr>
<tr>
<td><strong>LDAP User Search Base</strong> *</td>
</tr>
<tr>
<td>OU=People,DC=company,DC=com</td>
</tr>
<tr>
<td><strong>LDAP User Search Filter</strong></td>
</tr>
<tr>
<td>objectClass=person</td>
</tr>
<tr>
<td><strong>LDAP User Username Attribute</strong> *</td>
</tr>
<tr>
<td>sAMAccountName</td>
</tr>
</tbody>
</table>

When you select **Use Direct Bind**, Cloudera Data Science Workbench performs a direct bind to the LDAP server using the LDAP Username Pattern with the credentials provided on login (not **LDAP Bind DN** and **LDAP Bind Password**).

By default, Cloudera Data Science Workbench performs an LDAP search using the bind DN and credentials specified for the **LDAP Bind DN** and **LDAP Bind Password** configurations. It searches the subtree, starting from the base DN specified for the **LDAP Search Base** field, for an entry whose attribute specified in **LDAP User Username Attribute**, has the same value as the username provided on login. Cloudera Data Science Workbench then validates the user-provided password against the DN found as a result of the search.

**LDAP Over SSL (LDAPS)**

To support secure communication between Cloudera Data Science Workbench and the LDAP/Active Directory server, Cloudera Data Science Workbench might require a CA certificate to be able to validate the identity of the LDAP/Active Directory service.

- **CA Certificate**: If the certificate of your LDAP/Active Directory service was signed by a trusted or commercial Certificate Authority (CA), it is not necessary to upload the CA certificate here. However, if your LDAP/Active Directory certificate was signed by a self-signed CA, you must upload the self-signed CA certificate to Cloudera Data Science Workbench in order to use LDAP over SSL (LDAPS).

**LDAP Group Settings**

**Important**: Due to a known issue in Cloudera Data Science Workbench 1.5.x, LDAP group search fails when Active Directory returns escape characters as part of the distinguished name (DN).

In addition to the general LDAP settings, you can use the following group settings to restrict the access to Cloudera Data Science Workbench to certain groups in LDAP:

- **LDAP Group Search Base**: The base distinguished name (DN) where Cloudera Data Science Workbench will search for groups.

- **LDAP Group Search Filter**: The LDAP filter that Cloudera Data Science Workbench will use to determine whether a user is affiliated to a group.
A group object in LDAP or Active Directory typically has one or more **member** attributes that stores the DNs of users in the group. If **LDAP Group Search Filter** is set to **member=(0)**, Cloudera Data Science Workbench will automatically substitute the **(0)** placeholder for the DN of the authenticated user.

- **LDAP User Groups**: A list of LDAP groups whose users have access to Cloudera Data Science Workbench. When this property is set, only users that successfully authenticate themselves AND are affiliated to at least one of the groups listed here, will be able to access Cloudera Data Science Workbench.

  If this property is left empty, all users that can successfully authenticate themselves to LDAP will be able to access Cloudera Data Science Workbench.

- **LDAP Full Administrator Groups**: A list of LDAP groups whose users are automatically granted the **site administrator** role on Cloudera Data Science Workbench.

  The **LDAP User Groups** and **LDAP Full Administrator Groups** properties work independently. While the former specifies which groups can access Cloudera Data Science Workbench, the latter specifies a *subset* of groups that should also be granted site administrator privileges. This means to take advantage of both features, you must make sure that any groups listed in **LDAP Full Administrator Groups** are also listed in **LDAP User Groups**.

  If you want to restrict access to Cloudera Data Science Workbench to members of a group whose DN is:

  ```
  CN=CDSWUsers,OU=Groups,DC=company,DC=com
  ```

  And automatically grant site administrator privileges to members of a group whose DN is:

  ```
  CN=CDSWAdmins,OU=Groups,DC=company,DC=com
  ```

  Add the CNs of both groups to the following settings in Cloudera Data Science Workbench:

  - **LDAP User Groups**: CDSWUsers, CDSWAdmins
  - **LDAP Full Administrator Groups**: CDSWAdmins

**Figure 5: Example**

### How Login Works with LDAP Group Settings Enabled

With LDAP Group settings enabled, the login process in Cloudera Data Science Workbench works as follows:

1. **Authentication with LDAP**
When an unauthenticated user first accesses Cloudera Data Science Workbench, they are sent to the login page where they can login by providing a username and password.

Cloudera Data Science Workbench will search for the user by binding to the LDAP Bind DN and verify the username/password credentials provided by the user.

2. Authorization Check for Access to Cloudera Data Science Workbench

If the user is authenticated successfully, Cloudera Data Science Workbench will then use the LDAP Group Search Filter to search for all groups the user is affiliated to, in the DN provided by LDAP Group Search Base.

The list of LDAP groups the user belongs to is then compared to the pre-authorized list of groups specified in the LDAP User Groups property. If there is at least one match, this user will be allowed to access Cloudera Data Science Workbench.

3. Check for Site Administrator Privileges

If the user also belongs to an LDAP group that has been pre-authorized for site administrator privileges in the LDAP Full Administrator Groups property, Cloudera Data Science Workbench will automatically grant the site admin role to this user upon login.

Test LDAP Configuration

You can test your LDAP/Active Directory configuration by entering your username and password in the Test LDAP Configuration section. This form simulates the user login process and allows you to verify the validity of your LDAP/Active Directory configuration without opening a new window.

Before using this form, make sure you click Update to save the LDAP configuration you want to test.

Configuring SAML Authentication

Cloudera Data Science Workbench supports the Security Assertion Markup Language (SAML) for Single Sign-on (SSO) authentication; in particular, between an identity provider (IDP) and a service provider (SP). The SAML specification defines three roles: the principal (typically a user), the IDP, and the SP. In the use case addressed by SAML, the principal (user agent) requests a service from the service provider. The service provider requests and obtains an identity assertion from the IDP. On the basis of this assertion, the SP can make an access control decision—in other words it can decide whether to perform some service for the connected principal.

The primary SAML use case is called web browser single sign-on (SSO). A user with a user agent (usually a web browser) requests a web resource protected by a SAML SP. The SP, wanting to know the identity of the requesting user, issues an authentication request to a SAML IDP through the user agent. In the context of this terminology, Cloudera Data Science Workbench operates as a SP.

Cloudera Data Science Workbench supports both SP- and IDP-initiated SAML 2.0-based SSO. Its Assertion Consumer Service (ACS) API endpoint is for consuming assertions received from the Identity Provider. If your Cloudera Data Science Workbench domain root were cdsw.company.com, then this endpoint would be available at http://cdsw.company.com/api/v1/saml/acs. SAML 2.0 metadata is available at http://cdsw.company.com/api/v1/saml/metadata for IDP-initiated SSO. Cloudera Data Science Workbench uses HTTP Redirect Binding for authentication requests and expects to receive responses from HTTP POST Binding.

When Cloudera Data Science Workbench receives the SAML responses from the Identity Provider, it expects to see at least the following user attributes in the SAML responses:

- The unique identifier or username. Valid attributes are:
  - uid
  - urn:oid:0.9.2342.19200300.100.1.1

- The email address. Valid attributes are:
  - mail
  - email
  - urn:oid:0.9.2342.19200300.100.1.3
The common name or full name of the user. Valid attributes are:

- `cn`
- `urn:oid:2.5.4.3`

In the absence of the `cn` attribute, Cloudera Data Science Workbench will attempt to use the following user attributes, if they exist, as the full name of the user:

- The first name of the user. Valid attributes are:
  - `givenName`
  - `urn:oid:2.5.4.42`
- The last name of the user. Valid attributes are:
  - `sn`
  - `urn:oid:2.5.4.4`

Configuration Options

Use the following properties to configure SAML authentication and authorization in Cloudera Data Science Workbench. For an overview of the login process, see How Login Works with SAML Group Settings Enabled on page 243.

Cloudera Data Science Workbench Settings

- **Entity ID**: Required. A globally unique name for Cloudera Data Science Workbench as a Service Provider. This is typically the URI.
- **NameID Format**: Optional. The name identifier format for both Cloudera Data Science Workbench and Identity Provider to communicate with each other regarding a user. Default: `urn:oasis:names:tc:SAML:1.1:nameid-format:emailAddress`.
- **Authentication Context**: Optional. SAML authentication context classes are URIs that specify authentication methods used in SAML authentication requests and authentication statements. Default: `urn:oasis:names:tc:SAML:2.0:ac:classes:PasswordProtectedTransport`.

Signing SAML Authentication Requests

- **CDSW Private Key for Signing Authentication Requests**: Optional. If you upload a private key, you must upload a corresponding certificate as well so that the Identity Provider can use the certificate to verify the authentication requests sent by Cloudera Data Science Workbench. You can upload the private key used for both signing authentication requests sent to Identity Provider and decrypting assertions received from the Identity Provider.
- **CDSW Certificate for Signature Validation**: Required if the Cloudera Data Science Workbench Private Key is set, otherwise optional. You can upload a certificate in the PEM format for the Identity Provider to verify the authenticity of the authentication requests generated by Cloudera Data Science Workbench. The uploaded certificate is made available at the `http://cdsw.company.com/api/v1/saml/metadata` endpoint.

SAML Assertion Decryption

Cloudera Data Science Workbench uses the following properties to support SAML assertion encryption & decryption.

- **CDSW Certificate for Encrypting SAML Assertions**: Must be configured on the Identity Provider so that Identity Provider can use it for encrypting SAML assertions for Cloudera Data Science Workbench.
- **CDSW Private Key for Decrypting SAML Assertions**: Used to decrypt the encrypted SAML assertions.

Identity Provider

- **Identity Provider SSO URL**: Required. The entry point of the Identity Provider in the form of URI.
• **Identity Provider Signing Certificate**: Optional. Administrators can upload the X.509 certificate of the Identity Provider for Cloudera Data Science Workbench to validate the incoming SAML responses.

Cloudera Data Science Workbench extracts the Identity Provider SSO URL and Identity Provider Signing Certificate information from the uploaded Identity Provider Metadata file. Cloudera Data Science Workbench also expects all Identity Provider metadata to be defined in a `<md:EntityDescriptor>` XML element with the namespace "urn:oasis:names:tc:SAML:2.0:metadata", as defined in the SAMLMeta-xsd schema.

For on-premises deployments, you must provide a certificate and private key, generated and signed with your trusted Certificate Authority, for Cloudera Data Science Workbench to establish secure communication with the Identity Provider.

**Authorization**

When you’re using SAML 2.0 authentication, you can use the following properties to restrict the access to Cloudera Data Science Workbench to certain groups of users:

- **SAML Attribute Identifier for User Role**: The Object Identifier (OID) of the user attribute that will be provided by your identity provider for identifying a user’s role/affiliation. You can use this field in combination with the following **SAML User Groups** property to restrict access to Cloudera Data Science Workbench to only members of certain groups.

  For example, if your identity provider returns the `OrganizationalUnitName` user attribute, you would specify the OID of the `OrganizationalUnitName`, which is `urn:oid:2.5.4.11`, as the value for this property.

- **SAML User Groups**: A list of groups whose users have access to Cloudera Data Science Workbench. When this property is set, only users that are successfully authenticated AND are affiliated to at least one of the groups listed here, will be able to access Cloudera Data Science Workbench.

  For example, if your identity provider returns the `OrganizationalUnitName` user attribute, add the value of this attribute to the **SAML User Groups** list to restrict access to Cloudera Data Science Workbench to that group.

  If this property is left empty, all users that can successfully authenticate themselves will be able to access Cloudera Data Science Workbench.

- **SAML Full Administrator Groups**: A list of groups whose users are automatically granted the site administrator role on Cloudera Data Science Workbench.

  The **SAML User Groups** and **SAML Full Administrator Groups** properties work independently. While the former specifies which groups can access Cloudera Data Science Workbench, the latter specifies a *subset* of groups that should also be granted site administrator privileges. This means to take advantage of both features, you must make sure that any groups listed in **SAML Full Administrator Groups** are also listed in **SAML User Groups**.

**How Login Works with SAML Group Settings Enabled**

With SAML Group settings enabled, the login process in Cloudera Data Science Workbench works as follows:

1. **Authentication by Identity Provider**

   When an unauthenticated user accesses Cloudera Data Science Workbench, they are first sent to the identity provider’s login page, where the user can login as usual.

   Once successfully authenticated by the identity provider, the user is sent back to Cloudera Data Science Workbench along with a SAML assertion that includes, amongst other things, a list of the user’s attributes.

2. **Authorization Check for Access to Cloudera Data Science Workbench**

   Cloudera Data Science Workbench will attempt to look up the value of the **SAML Attribute Identifier for User Role** in the SAML assertion and check to see whether that value, which could be one or more group names, exists in the **SAML User Groups** whitelist. The user will only be granted access if the user is affiliated to at least one of the groups listed in **SAML User Groups**.

3. **Check for Site Administrator Privileges**

   The **SAML User Groups** and **SAML Full Administrator Groups** properties work independently. While the former specifies which groups can access Cloudera Data Science Workbench, the latter specifies a *subset* of groups that should also be granted site administrator privileges. This means to take advantage of both features, you must make sure that any groups listed in **SAML Full Administrator Groups** are also listed in **SAML User Groups**.
If the user also belongs to a group that is listed in the **SAML Full Administrator Groups** property, Cloudera Data Science Workbench will automatically grant the site admin role to this user upon login.

**Debug Login URL**

When using external authentication, such as LDAP, Active Directory or SAML 2.0, even a small mistake in authentication configurations in either Cloudera Data Science Workbench or the Identity Provider could potentially block all users from logging in.

Cloudera Data Science Workbench provides an optional fallback debug login URL for site administrators to log in against the local database with their username/password created during the signup process before changing the external authentication method. The debug login URL is `http://cdsw.company.com/login?debug=1`. If you do not remember the original password, you can reset it by going directly to `http://cdsw.company.com/forgot-password`. When configured to use external authentication, the link to the forgot password page is disabled on the login page for security reasons.

**Disabling the Debug Login Route**

Optionally, the debug login route can be disabled to prevent users from accessing Cloudera Data Science Workbench via local database when using external authentication. In case of external authentication failures, when the debug login route is disabled, root access to the master host is required to re-enable the debug login route.

Contact Cloudera Support for more information.

**Configuring HTTP Headers for Cloudera Data Science Workbench**

**Required Role:** Site Administrator

Cloudera Data Science Workbench 1.4.2 (and higher) include three properties that allow you to customize the HTTP headers accepted by Cloudera Data Science Workbench. They are available under the site administrator panel at Admin > Security.

**Important:** Any changes to the following properties require a full restart of Cloudera Data Science Workbench. For CSD deployments, go to Cloudera Manager and restart the CDSW service. For RPM deployments, run `cdsw restart` on the master host.

**Enable HTTP Security Headers**

When **Enable HTTP security headers** is enabled, the following HTTP headers will be included in HTTP responses from servers:

- X-XSS-Protection
- X-DNS-Prefetch-Control
- X-Frame-Options
- X-Download-Options
- X-Content-Type-Options

This property is **enabled by default**.

Disabling this property could leave your Cloudera Data Science Workbench deployment vulnerable to clickjacking, cross-site scripting (XSS), or any other injection attacks.

**Enable HTTP Strict Transport Security (HSTS)**

**Note:** Without TLS/SSL enabled, configuring this property will have no effect on your browser.
When both TLS/SSL and this property (Enable HTTP Strict Transport Security (HSTS)) are enabled, Cloudera Data Science Workbench will inform your browser that it should never load the site using HTTP. Additionally, all attempts to access Cloudera Data Science Workbench using HTTP will automatically be converted to HTTPS.

This property is disabled by default.

If you ever need to downgrade to back to HTTP, use the following sequence of steps: First, deactivate this checkbox to disable HSTS and restart Cloudera Data Science Workbench. Then, load the Cloudera Data Science Workbench web application in each browser to clear the respective browser's HSTS setting. Finally, disable TLS/SSL across the cluster. Following this sequence should help avoid a situation where users get locked out of their accounts due to browser caching.

Enable Cross-Origin Resource Sharing (CORS)

Most modern browsers implement the Same-Origin Policy, which restricts how a document or a script loaded from one origin can interact with a resource from another origin. When the Enable cross-origin resource sharing property is enabled on Cloudera Data Science Workbench, web servers will include the Access-Control-Allow-Origin: * HTTP header in their HTTP responses. This gives web applications on different domains permission to access the Cloudera Data Science Workbench API through browsers.

This property is enabled by default.

If this property is disabled, web applications from different domains will not be able to programmatically communicate with the Cloudera Data Science Workbench API through browsers.

SSH Keys

This topic describes the different types of SSH keys used by Cloudera Data Science Workbench, and how you can use those keys to authenticate to an external service such as GitHub.

Personal Key

Cloudera Data Science Workbench automatically generates an SSH key pair for your user account. You can rotate the key pair and view your public key on your user settings page. It is not possible for anyone to view your private key.

Every console you run has your account’s private key loaded into its SSH-agent. Your consoles can use the private key to authenticate to external services, such as GitHub. For instructions, see Adding SSH Key to GitHub on page 245.

Team Key

Like Cloudera Data Science Workbench users, each Cloudera Data Science Workbench team has an associated SSH key. You can access the public key from the team's account settings. Click Account, then select the team from the drop-down menu at the upper right corner of the page.

Team SSH keys provide a useful way to give an entire team access to external resources such as databases or GitHub repositories (as described in the next section). When you launch a console in a project owned by a team, you can use that team's SSH key from within the console.

Adding SSH Key to GitHub

If you want to use GitHub repositories to create new projects or collaborate on projects, use the following instructions to add your Cloudera Data Science Workbench SSH public key to your GitHub account:

1. Sign in to Cloudera Data Science Workbench.
2. Go to the upper right drop-down menu and switch context to the account whose key you want to add.
3. On the left sidebar, click Settings.
4. Go to the SSH Keys tab and copy your public SSH key.
5. Sign in to your GitHub account and add the Cloudera Data Science Workbench key copied in the previous step to your GitHub account. For instructions, refer the GitHub documentation on adding SSH keys to GitHub.
SSH Tunnels

**Important:** SSH tunnels do not work in Cloudera Data Science Workbench 1.4.0. This issue has been fixed in Cloudera Data Science Workbench 1.4.2 (and higher).

In some environments, external databases and data sources reside behind restrictive firewalls. A common pattern is to provide access to these services using a bastion host with only the SSH port open. This introduces complexity for end users who must manually set up SSH tunnels to access data. Cloudera Data Science Workbench provides a convenient way to connect to such resources.

From the Project > Settings > Tunnels page, you can use your SSH key to connect Cloudera Data Science Workbench to an external database or cluster by creating an SSH tunnel. If you create an SSH tunnel to an external server in one of your projects, then all engines that you run in that project are able to connect securely to a port on that server by connecting to a local port. The encrypted tunnel is completely transparent to the user or code.

To create an automatic SSH tunnel:

1. Open the Project Settings page.
2. Open the Tunnels tab.
3. Click New Tunnel.
4. Enter the server IP Address or DNS hostname.
5. Enter your username on the server.
6. Enter the local port that should be proxied, and to which remote port on the server.

Then, on the remote server, configure SSH to accept password-less logins using your individual or team SSH key. Often, you can do so by appending the SSH key to the file `/home/username/.ssh/authorized_keys`.
Troubleshooting Cloudera Data Science Workbench

Use one or more of the following courses of action to start debugging issues with Cloudera Data Science Workbench.

- Check the status of the application.
  
  ```
  cdsw status
  ```

- SSH to your master host and run the following host validation command to check that the key services are running:
  
  ```
  cdsw validate
  ```

- Make sure your Cloudera Data Science Workbench configuration is correct.
  
  **CSD Deployments**
  
  Log into Cloudera Manager and review configuration for the CDSW service.

  **RPM Deployments**
  
  ```
  cat /etc/cdsw/config/cdsw.conf
  ```

The following sections describe solutions to potential problems and error messages you may encounter while installing, configuring or using Cloudera Data Science Workbench.

Understanding Installation Warnings

This section describes solutions to some warnings you might encounter during the installation process.

**Preexisting iptables rules not supported**

```
WARNING: Cloudera Data Science Workbench requires iptables, but does not support preexisting iptables rules.

Kubernetes makes extensive use of iptables. However, it’s hard to know how pre-existing iptables rules will interact with the rules inserted by Kubernetes. Therefore, Cloudera recommends you run the following commands to clear all pre-existing rules before you proceed with the installation.

```

```
sudo iptables -P INPUT ACCEPT
sudo iptables -P FORWARD ACCEPT
sudo iptables -P OUTPUT ACCEPT
sudo iptables -t nat -F
sudo iptables -t mangle -F
sudo iptables -F
sudo iptables -X
```

The warning can be ignored after you clear the pre-existing rules or are sure that there are no pre-existing iptables rules.

**Remove the entry corresponding to /dev/xvdc from /etc/fstab**

Cloudera Data Science Workbench installs a custom filesystem on its Application and Docker block devices. These filesystems will be used to store user project files and Docker engine images respectively. Therefore, Cloudera Data Science Workbench requires complete access to the block devices. To avoid losing any existing data, make sure the block devices allocated to Cloudera Data Science Workbench are reserved only for the workbench.
Linux sysctl kernel configuration errors

Kubernetes and Docker require non-standard kernel configuration. Depending on the existing state of your kernel, this might result in sysctl errors such as:

- `sysctl net.bridge.bridge-nf-call-iptables must be set to 1`

This is because the settings in `/etc/sysctl.conf` conflict with the settings required by Cloudera Data Science Workbench. Cloudera cannot make a blanket recommendation on how to resolve such errors because they are specific to your deployment. Cluster administrators may choose to either remove or modify the conflicting value directly in `/etc/sysctl.conf`, remove the value from the conflicting configuration file, or even delete the module that is causing the conflict.

To start diagnosing the issue, run the following command to see the list of configuration files that are overwriting values in `/etc/sysctl.conf`.

```
SYSTEMD_LOG_LEVEL=debug /usr/lib/systemd/systemd-sysctl
```

You will see output similar to:

```
Parsing /usr/lib/sysctl.d/00-system.conf
Parsing /usr/lib/sysctl.d/50-default.conf
Parsing /etc/sysctl.d/99-sysctl.conf
Overwriting earlier assignment of net/bridge/bridge-nf-call-ip6tables in file '/etc/sysctl.d/99-sysctl.conf'.
Overwriting earlier assignment of net/bridge/bridge-nf-call-ip6tables in file '/etc/sysctl.d/99-sysctl.conf'.
Overwriting earlier assignment of net/bridge/bridge-nf-call-ip6tables in file '/etc/sysctl.d/99-sysctl.conf'.
Parsing /etc/sysctl.d/k8s.conf
Overwriting earlier assignment of net/bridge/bridge-nf-call-iptables in file '/etc/sysctl.d/k8s.conf'.
Parsing /etc/sysctl.d/k8s.conf
Overwriting earlier assignment of net/bridge/bridge-nf-call-ip6tables in file '/etc/sysctl.d/k8s.conf'.
Overwriting earlier assignment of net/bridge/bridge-nf-call-ip6tables in file '/etc/sysctl.d/k8s.conf'.
Setting 'net/ipv4/conf/all/promote_secondaries' to '1'
Setting 'net/ipv4/conf/default/promote_secondaries' to '1'
...
```

`/etc/sysctl.d/k8s.conf` is the configuration added by Cloudera Data Science Workbench. Administrators must make sure that no other file is overwriting values set by `/etc/sysctl.d/k8s.conf`.

CDH parcels not found at /opt/cloudera/parcels

There are two possible reasons for this warning:

- If you are using a custom parcel directory, you can ignore the warning and proceed with the installation. Once the Cloudera Data Science Workbench is running, set the path to the CDH parcel in the admin dashboard. See Configuring the Engine Environment on page 174.
- This warning can be an indication that you have not added gateway roles to the Cloudera Data Science Workbench hosts. In this case, do not ignore the warning. Exit the installer and go to Cloudera Manager to add gateway roles to the cluster. See Configure Gateway Hosts Using Cloudera Manager on page 67.

CDSW docker daemons fail to start

CDSW docker daemons fail to start with the following error:

```
Error starting daemon: error initializing graphdriver: devmapper: Unable to take ownership of thin-pool (docker-thinpool) that already has used data blocks.
```
This issue occurs when the block devices you specified for the **Docker Block Device** field already have data on them. This is a safeguard to prevent block devices from being wiped inadvertently. Note that resolving this issue involves deleting data from the block devices.

To resolve this issue, perform the following steps:

1. Verify that it is okay to delete the data on the block device.
2. SSH to the Cloudera Data Science Workbench master host.
3. Run the following script:
   ```bash
   /opt/cloudera/parcels/CDSW/scripts/teardown-docker.sh
   ```
4. In the Cloudera Manager Admin Console, select the Cloudera Data Science Workbench service.
5. On the **Instances** tab, select the Docker Daemons.
6. Click **Actions for Selected (n) > Prepare Node**.
7. Start the Cloudera Data Science Workbench service by clicking **Actions > Start**.

### User Process Limit

During host validation, you may encounter the following warning message:

```
{WARN} Cloudera Data Science Workbench recommends that all users have a max-user-processes limit of at least 65536.
```

This message appears if the user process limit is under 65536. You can increase the user process limit by adding the following line to the `/etc/security/limits.conf` file:

```
ulimit -u 65536
```

Set this configuration on every Cloudera Data Science Workbench host. You can also edit `/etc/security/limits.conf` to configure the user process limit.

### Open Files Limit

During host validation, you may encounter the following warning message:

```
{WARN} Cloudera Data Science Workbench recommends that all users have a max-open-files limit set to 1048576.
```

This message appears if the open files limit is under 1048576. Note that on HDP clusters, the open file limit recommendation is 10000 at a minimum. Cloudera recommends a higher limit for clusters with Cloudera Data Science Workbench.

You can configure the file limit with the following command:

```
ulimit -n 1048576
```

Set this configuration on every Cloudera Data Science Workbench host. You can also edit `/etc/security/limits.conf` to configure the open files limit.

### Disable SE Linux

During installation, you may encounter the following message:

```
Please disable SELinux by setting SELINUX=disabled|permissive in /etc/selinux/config, then reboot or using setenforce 0 command.
```

---

Troubleshooting Cloudera Data Science Workbench
SELinux enforces additional control policies for what a user, process, or daemon can do. If SELinux is enabled or not in permissive mode, Cloudera Data Science Workbench may not have the proper permissions to run.

To resolve this issue, you must change the SELinux mode on every host by doing one of the following:

- Edit the configuration file for SELinux and set it to disabled or permissive. Note that if you set SELinux to permissive mode, events such as access denials will be logged, but the denial will not be enforced. You can find the SELinux configuration file in the following location: /etc/selinux/config.
- Run the following command: `setenforce 0`. This command disables SELinux completely.

Failed to run Kernel memory slabs check

Users might see the following error message in Cloudera Manager after upgrading to Cloudera Data Science Workbench 1.4.2.

```
Bad: Failed to run Kernel memory slabs check
```

This error is an indication that Cloudera Data Science Workbench hosts were not rebooted as part of the upgrade to version 1.4.2. The host reboot is required to fix a Red Hat kernel slab leak issue that was discovered in Cloudera Data Science Workbench 1.4.0. For more information, see: [Red Hat Only] Host Reboot Required for Upgrades from Cloudera Data Science Workbench 1.4.0 on page 44.

To proceed, stop Cloudera Data Science Workbench and reboot all Cloudera Data Science Workbench hosts. As a precaution, you might want to consult your cluster/IT administrator before you start rebooting hosts. Once all hosts have rebooted, restart Cloudera Data Science Workbench.

If that does not fix the issue, contact Cloudera Support.

Error Encountered Trying to Load Images when Initializing Cloudera Data Science Workbench

Here are some sample error messages you might see when initializing Cloudera Data Science Workbench:

```
Error encountered while trying to load images.: 1

Unable to load images from [/etc/cdsw/images/cdsw_<version>.tar.gz].: 1

Error processing tar file(exit status 1): write /../..tar: no space left on device
```

These errors are an indication that the root volume is running out of space when trying to initialize Cloudera Data Science Workbench. During the initialization process, the Cloudera Data Science Workbench installer temporarily decompresses the engine image file located in /etc/cdsw/images to the /var/lib/docker/tmp/ directory.

If you have previously partitioned the root volume (which should be at least 100 GB), make sure you allocate at least 20 GB to /var/lib/docker/tmp so that the installer can proceed without running out of space.

404 Not Found Error

The 404 Not Found error might appear in the browser when you try to reach the Cloudera Data Science Workbench web application.

This error is an indication that your installation of Cloudera Data Science Workbench was successful, but there was a mismatch in the domain configured in cdsw.conf and the domain referenced in the browser. To fix the error, go to /etc/cdsw/config/cdsw.conf and check that the URL you supplied for the DOMAIN property matches the one you
are trying to use to reach the web application. This is the wildcard domain dedicated to Cloudera Data Science Workbench, not the hostname of the master host.

If this requires a change to cdsw.conf, after saving the changes run cdsw reset followed by cdsw init.

Troubleshooting Kerberos Errors

HDFS commands fail with Kerberos errors even though Kerberos authentication is successful in the web application

If Kerberos authentication is successful in the web application, and the output of klist in the engine reveals a valid-looking TGT, but commands such as hdfs dfs -ls / still fail with a Kerberos error, it is possible that your cluster is missing the Java Cryptography Extension (JCE) Unlimited Strength Jurisdiction Policy File. The JCE policy file is required when Red Hat uses AES-256 encryption. This library should be installed on each cluster host and will live under $JAVA_HOME. For more information, see Using AES-256 Encryption.

Cannot find renewable Kerberos TGT

Cloudera Data Science Workbench runs its own Kerberos TGT renewer which produces non-renewable TGT. However, this confuses Hadoop's renewer which looks for renewable TGTs. If the Spark 2 logging level is set to WARN or lower, you may see exceptions such as:

```

16/12/24 16:41:23 WARN security.UserGroupInformation: PrivilegedActionException as:user@CLOUDERA.LOCAL (auth:KERBEROS) cause:javax.security.sasl.SaslException: GSS initiate failed [Caused by GSSException: No valid credentials provided (Mechanism level: Failed to find any Kerberos tgt)]
```

This is not a bug. Spark 2 workloads will not be affected by this. Access to Kerberized resources should also work as expected.

Troubleshooting TLS/SSL Errors

This section describes some common issues with TLS configuration on Cloudera Data Science Workbench. Common errors include:

- Cloudera Data Science Workbench initialisation fails with an error such as:

```
Error preparing server: tls: failed to parse private key
```

- Your browser reports that the Cloudera Data Science Workbench web application is not secure even though you have enabled TLS settings as per Enabling TLS/SSL for Cloudera Data Science Workbench on page 231.

Possible Causes and Solutions

- **Certificate does not include the wildcard domain** - Confirm that the TLS certificate issued by your CA lists both, the Cloudera Data Science Workbench domain, as well as a wildcard for all first-level subdomains. For example, if your Cloudera Data Science Workbench domain is cdsw.company.com, then the TLS certificate must include both cdsw.company.com and *.cdsw.company.com.

- **Path to the private key and/or certificate is incorrect** - Confirm that the path to the private key file is correct by comparing the path and file name to the values for TLS_KEY and/or TLS_CERT in cdsw.conf or Cloudera Manager. For example:

```
TLS_CERT="/path/to/cert.pem"
TLS_KEY="/path/to/private.key"
```
Troubleshooting Cloudera Data Science Workbench

- **Private key file does not have the right permissions** - The private key file must have read-only permissions. Set it as follows:

  `chmod 444 private.key`

- **Private key is encrypted** - Cloudera Data Science Workbench does not support encrypted private keys. Check to see if your private key is encrypted:

  `cat private.key`

  -----BEGIN RSA PRIVATE KEY-----
  Proc-Type: 4,ENCRYPTED
  DEK-Info: DES-EDE3-CBC,11556F53E4A2824A

  If the private key is encrypted as shown above, use the following steps to decrypt it:

  1. Make a backup of the private key file.

     `mv private.key private.key.encrypted`

  2. Decrypt the backup private key and save the file to `private.key`. You will be asked to enter the private key password.

     `openssl rsa -in private.key.encrypted -out private.key`

- **Private key and certificate are not related** - Check to see if the private key matches the public key in the certificate.

  1. Print a hash of the private key modulus.

     `openssl rsa -in private.key -noout -modulus | openssl md5`

     `(stdin)= 7a8d72ed61bb4be3c1f59e4f0161c023`

  2. Print a hash of the public key modulus.

     `openssl x509 -in cert.pem -noout -modulus | openssl md5`

     `(stdin)= 7a8d72ed61bb4be3c1f59e4f0161c023`

  If the md5 hash output of both keys is different, they are not related to each other, and will not work. You must revoke the old certificate, [regenerate a new private key and Certificate Signing Request (CSR)](https://example.com), and then apply for a new certificate.

Troubleshooting Issues with Workloads

This section describes some potential issues data scientists might encounter once the application is running workloads.

**404 error in Workbench after starting an engine**

This is typically caused because a wildcard DNS subdomain was not set up before installation. While the application will largely work, the engine consoles are served on subdomains and will not be routed correctly unless a wildcard DNS entry pointing to the master host is properly configured. You might need to wait 30-60 minutes until the DNS entries propagate. For instructions, see [Set Up a Wildcard DNS Subdomain](https://example.com) on page 60.
Engines cannot be scheduled due to lack of CPU or memory

A symptom of this is the following error message in the Workbench: "Unschedulable: No node in the cluster currently has enough CPU or memory to run the engine."

Either shut down some running sessions or jobs or provision more hosts for Cloudera Data Science Workbench.

Workbench prompt flashes red and does not take input

The Workbench prompt flashing red indicates that the session is not currently ready to take input.

Cloudera Data Science Workbench does not currently support non-REPL interaction. One workaround is to skip the prompt using appropriate command-line arguments. Otherwise, consider using the terminal to answer interactive prompts.

PySpark jobs fail due to HDFS permission errors

```
```

(Required for CDH 5 and CDH 6) To be able to use Spark 2, each user must have their own /home directory in HDFS. If you sign in to Hue first, these directories will automatically be created for you. Alternatively, you can have cluster administrators create these directories.

```
hdfs dfs -mkdir /user/<username>
hdfs dfs -chown <username>:<username> /user/<username>
```

PySpark jobs fail due to Python version mismatch

```
Exception: Python in worker has different version 2.6 than that in driver 2.7, PySpark cannot run with different minor versions
```

One solution is to install the matching Python 2.7 version on all the cluster hosts. Another, more recommended solution is to install the Anaconda parcel on all CDH cluster hosts. Cloudera Data Science Workbench Python engines will use the version of Python included in the Anaconda parcel which ensures Python versions between driver and workers will always match. Any library paths in workloads sent from drivers to workers will also match because Anaconda is present in the same location across all hosts. Once the parcel has been installed, set the PYSPARK_PYTHON environment variable in the Cloudera Data Science Workbench Admin dashboard. Alternatively, you can use Cloudera Manager to set the path.

Jobs fail due to incorrect JAVA_HOME on HDP

Commands, such as hdfs commands, and jobs fail with an error similar to the following message:

```
ERROR: JAVA_HOME /usr/lib/jvm/java does not exist.
```

The JAVA_HOME path you configure for Cloudera Data Science Workbench in cdsw.conf must match the JAVA_HOME configured by hadoop-env.sh for the HDP cluster. After you update JAVA_HOME in cdsw.conf, you must restart Cloudera Data Science Workbench. For more information, see Changes to cdsw.conf.

Troubleshooting Issues with Models and Experiments

See the following topics:

- [Debugging Issues with Experiments](#) on page 145
- [Debugging Issues with Models](#) on page 162
This topic describes the commands available with the Cloudera Data Science Workbench command line utility, \texttt{cdsw}. Running \texttt{cdsw} without any arguments will print a brief description of each command.

**Start, Stop, Restart for CSD Deployments:** The commands available for a CSD-based deployment are only a subset of those available for an RPM deployment. For example, the CLI for CSD deployments does not have commands such as \texttt{cdsw start}, \texttt{stop}, and \texttt{restart} available. Instead, these actions must be executed through the Cloudera Data Science Workbench service in Cloudera Manager. For instructions, see Starting, Stopping, and Restarting the Service on page 209.

> **Important:** On Cloudera Data Science Workbench 1.4.0 (and lower), do not stop or restart Cloudera Data Science Workbench without using the \texttt{cdsw_protect_stop_restart.sh} script. This is to help avoid the data loss issue detailed in TSB-346.

All of the following commands can be used in an RPM-based deployment. Those available for CSD-based deployments have been marked in the table.

<table>
<thead>
<tr>
<th>Command</th>
<th>Description and Usage</th>
</tr>
</thead>
<tbody>
<tr>
<td>\texttt{cdsw init}</td>
<td>Initializes and bootstraps the master host. Use this command to start Cloudera Data Science Workbench.  &lt;br/&gt; Also see, Additional Usage Notes on page 255</td>
</tr>
<tr>
<td>\texttt{cdsw start}</td>
<td>Run on the master host to start application components.</td>
</tr>
<tr>
<td>\texttt{cdsw stop}</td>
<td>Run on the master host to stop application components.</td>
</tr>
<tr>
<td>\texttt{cdsw restart}</td>
<td>Run on the master host to restart application components.  &lt;br/&gt; To restart a worker host, use \texttt{cdsw reset}, followed by \texttt{cdsw join}. These commands have been explained further in this topic.</td>
</tr>
<tr>
<td>\texttt{cdsw reset}</td>
<td>De-registers and resets a host.  &lt;br/&gt; On a worker host, this command will remove the worker from the cluster.  &lt;br/&gt; On the master host, this command will bring down the application and effectively tear down the Cloudera Data Workbench deployment.</td>
</tr>
<tr>
<td>\texttt{cdsw join}</td>
<td>Initializes a worker host. After a worker host has been added, run this command on the worker host to add it to the Cloudera Data Science Workbench cluster.  &lt;br/&gt; This registers the worker hosts with the master, and increases the available pool of resources for workloads.</td>
</tr>
<tr>
<td>\texttt{cdsw status}</td>
<td>Displays the current status of the application.  &lt;br/&gt; Starting with version 1.4, you can use \texttt{cdsw status -v} or \texttt{cdsw status --verbose} for more detailed output.  &lt;br/&gt; The \texttt{cdsw status} command is not supported on worker hosts.</td>
</tr>
<tr>
<td>\texttt{cdsw validate}</td>
<td>Performs diagnostic checks for common errors that might be preventing the application from running as expected.  &lt;br/&gt; This command should typically be run as the first step to troubleshooting any problems with the application, as indicated by \texttt{cdsw status}.</td>
</tr>
<tr>
<td>Command</td>
<td>Description and Usage</td>
</tr>
<tr>
<td>-------------</td>
<td>-----------------------------------------------------------------------------------------------------------------------------------------------------</td>
</tr>
</tbody>
</table>
| `cdsw logs` | Creates a tarball with diagnostic information for your Cloudera Data Science Workbench installation.  
If you file a case with Cloudera Support, run this command on each host and attach the resulting bundle to the case.  
For more details on the information collected in these bundles, see Data Collection in Cloudera Data Science Workbench on page 212. |
| `cdsw version` | Displays the version number and type of Cloudera Data Science Workbench deployment (RPM or CSD). |
| `cdsw help` | Displays the inline help options for the Cloudera Data Science Workbench CLI. |

**Additional Usage Notes**

**Note:** These notes apply only to RPM-based deployments. In case of CSD-based deployments where you cannot directly modify `cdsw.conf`, Cloudera Manager will prompt you if the Cloudera Data Science Workbench service needs to be restarted.

**Changes to `cdsw.conf`:** Make sure `cdsw.conf` is consistent across all Cloudera Data Science Workbench hosts. Any changes made to the file must be copied over to all the other hosts.

- **Master Host** - Changes to the `JAVA_HOME`, `MASTER_IP`, `DOCKER_BLOCK_DEVICES`, and `APPLICATION_BLOCK_DEVICE` parameters in `cdsw.conf` require a re-initialization of the master host.

  ```
  cdsw reset  
  cdsw init
  ```

  Changes to other `cdsw.conf` parameters such as domain name changes, or TLS and HTTP proxy changes, require a restart of the application components.

  ```
  cdsw restart
  ```

- **Worker Host** - Changes to `cdsw.conf` on a worker host, require a restart of the worker host as follows:

  ```
  cdsw reset  
  cdsw join
  ```
Cloudera Data Science Workbench FAQs

Where can I get a sample project to try out Cloudera Data Science Workbench?

Cloudera Data Science Workbench ships with sample project templates that you can use to try running workloads. These are currently available in Python, R, and Scala. See Create a Project from a Built-in Template on page 98.

What are the software and hardware requirements for Cloudera Data Science Workbench?

For detailed information on the software and hardware required to successfully run Cloudera Data Science Workbench, see Cloudera Data Science Workbench 1.5.x Requirements and Supported Platforms on page 52.

Can I run Cloudera Data Science Workbench on hosts shared with other Hadoop services?

No. Cloudera does not support running Cloudera Data Science Workbench on non-dedicated hosts. Running other services on Cloudera Data Science Workbench hosts can lead to unreliable execution of workloads and difficult to debug out-of-memory errors.

How does Cloudera Data Science Workbench use Docker and Kubernetes?

Cloudera Data Science Workbench uses Docker and Kubernetes to manage containers. Currently, Cloudera Data Science Workbench only supports the versions of Docker and Kubernetes that are shipped with each release. Upgrading Docker, or Kubernetes, or running on third-party Kubernetes clusters is not supported.

Cloudera does not support Kubernetes or Docker for running any other workloads beyond those on Cloudera Data Science Workbench.

Can I run Cloudera Data Science Workbench on my own Kubernetes cluster?

This is not supported.

Does Cloudera Data Science Workbench support REST API access?

Starting with version 1.1.0, Cloudera Data Science Workbench supports a Jobs REST API that lets you orchestrate jobs from 3rd party workflow tools. For more details, see Cloudera Data Science Workbench Jobs API on page 165.

Other means of API access to Cloudera Data Science Workbench are not supported at this time.

How do I contact Cloudera for issues regarding Cloudera Data Science Workbench?

Cloudera Support

If you are a Cloudera customer, you can register for an account to create a support ticket at the support portal.

Before you log a support ticket, run the following command on the master host to create a tarball with diagnostic information for your Cloudera Data Science Workbench installation.

cdsw logs
Attach the resulting bundle to the support case you create.

**Cloudera Community**

Register for the [Cloudera Community forums](https://www.cloudera.com) and post your questions or feedback on the [Cloudera Data Science Workbench board](https://www.cloudera.com).
Cloudera Data Science Workbench Glossary

Terms related to Cloudera Data Science Workbench:

**Cloudera Data Science Workbench**
Cloudera Data Science Workbench is a product that enables fast, easy, and secure self-service data science for the enterprise. It allows data scientists to bring their existing skills and tools, such as R, Python, and Scala, to securely run computations on data in Hadoop clusters.

**site administrator**
A Cloudera Data Science Workbench user with all-access permissions. Site administrators can add or disable users/teams, monitor and manage resource usage, secure access to the deployment, and more. The Site Administration dashboard is only accessible to site administrators.

**API**
Cloudera Data Science Workbench exposes a limited REST API that allows you to schedule existing jobs from third-party workflow tools.

**cluster**
Refers to the CDH cluster managed by Cloudera Manager, including the gateway hosts that are running Cloudera Data Science Workbench.

**context**
Cloudera Data Science Workbench uses the notion of contexts to separate your personal account from any team accounts you belong to. This gives you leave to run experiments in your own personal context, while you can simultaneously collaborate with others in your organization within a team context.

**engine**
In Cloudera Data Science Workbench, engines are responsible for running R, Python, and Scala code written by users and for facilitating access to the CDH cluster. Each engine functions as an isolated virtual machine, customized to have all the necessary dependencies to access the CDH cluster while keeping each project’s environment entirely isolated. The only artifacts that remain after an engine runs is a log of the analysis and any files that were generated or modified inside the project’s filesystem, which is mounted to each engine at `/home/cdsw`.

**experiment**
Experiments are batch executed workloads that help facilitate model training in Cloudera Data Science Workbench.

**gateway host**
On a Cloudera Manager cluster, a gateway host is one that has been assigned a gateway role for a CDH service. Such a host will receive client configuration for that CDH service even though the host does not have any role instances for that service running on it.

Cloudera Data Science Workbench runs on dedicated gateway hosts on a CDH cluster. These hosts are assigned gateway roles for the Spark and HDFS services so that Cloudera Data Science Workbench has the client configuration required to access the CDH cluster.

**job**
Jobs are sessions that can be scheduled in advance and do not need to be launched manually each time.

**Livelog**
Cloudera Data Science Workbench allows users to work interactively with R, Python, and Scala from their browser and display results in realtime. This realtime state is stored in an internal database, called Livelog.

**master**
A typical Cloudera Data Science Workbench deployment consists of 1 master host and zero or more worker hosts. The master host keeps track of all critical, persistent, and stateful application data within Cloudera Data Science Workbench.
model

Model is a high level abstract term that is used to describe several possible incarnations of objects created during the model deployment process in Cloudera Data Science Workbench. You should note that 'model' does not always refer to a specific artifact. More precise terms (as defined in the documentation) should be used whenever possible.

pipeline

A series of jobs that depend on each other and must therefore be executed in a specific pre-defined sequence.

project

Projects hold the code, configuration, and libraries needed to reproducibly run data analytics workloads. Each project is independent, ensuring users can work freely without interfering with one another or breaking existing workloads.

session

A session is an interactive environment where you can run exploratory analysis in R, Python, and Scala.

team

A group of trusted users who are collaborating on a project in Cloudera Data Science Workbench.

terminal

Cloudera Data Science Workbench allows terminal access to actively running engines. The terminal can be used to move project files around, run Git commands, access the YARN and Hadoop CLIs, or install libraries that cannot be installed directly from the engine.

web application

Refers to the Cloudera Data Science Workbench web application running at cdsw.<your_domain>.com.

workbench

The console in the web application that is used to launch interactive sessions and run exploratory data analytic workloads. It consists of two panes, a navigable filesystem and editor on the left, and an interactive command prompt on the right.

worker

Worker hosts are transient hosts that can be added or removed from a Cloudera Data Science Workbench deployment depending on the number of users and workloads you are running.

Related Topics

- [Cloudera Enterprise Glossary](#)
- [Cloudera Director Glossary](#)
Appendix: Apache License, Version 2.0

**SPDX short identifier: Apache-2.0**

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