R on Databricks Compendium

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What is this?



♦ Under Development

i This is not intended to be an exhaustive guide, it's currently a place for me to document and collate useful information regarding R on Databricks.

There aren't many definitive examples of how to use R and Databricks together - hopefully the content here will serve as a useful resource.

Part I

MIflow

1 Log R Models to Unity Catalog

Currently {mlflow} doesn't support directly logging R models to Unity Catalog. This section will cover why, and then how to overcome each roadblock.

1.1 Unity Catalog Model Requirements

For models to be logged into Unity Catalog they **must** have a model signature. The Model signature defines the schema for model inputs/outputs.

Typically when using python this would be inferred via model input examples. Input examples are optional but strongly recommended.

The documentation discusses signature enforcement, currently this isn't implemented for R. Therefore you can decide if the signature is a dummy value for the sake of moving forward, or correct to clearly communicate the behaviour of the model.

! Important

It's important to clarify that for python the signature is enforced at time of inference *not* when registering the model to Unity Catalog.

The signature correctness is not validated when registering the model, it just has to be syntactically valid.

So, let's look at the existing code to log models in the crate flavour:

```
mlflow_save_model.crate <- function(model, path, model_spec=list(), ...) {
  if (dir.exists(path)) unlink(path, recursive = TRUE)
    dir.create(path)

  serialized <- serialize(model, NULL)

  saveRDS(
    serialized,
    file.path(path, "crate.bin")
  )</pre>
```

```
model_spec$flavors <- append(model_spec$flavors, list(
    crate = list(
    version = "0.1.0",
    model = "crate.bin"
    )
))
mlflow_write_model_spec(path, model_spec)
model_spec
}</pre>
```

- 1 Create the directory to save the model if it doesn't exist, if it does, empty it
- 2 Serialise the model, which is an object of class crate (from {carrier} package)
- (3) Save the serialised model via saveRDS to the directory as crate.bin
- 4 Define the model specification, this contains metadata required ensure reproducibility. In this case it's only specifying a version and what file the model can be found within.

The missing puzzle piece is the definition of a signature. Instead of explicitly adding code to the crate flavour itself, we'll take advantage of the model_spec parameter.

That means we can focus on mlflow::mlflow_log_model directly, we'd need to adjust the code as follows:

```
mlflow_log_model <- function(model, artifact_path, ...) {</pre>
                                                                                 (1)
  temp_path <- fs::path_temp(artifact_path)</pre>
  model_spec <- mlflow_save_model(</pre>
    model, path = temp_path,
    model_spec = list(
                                                                                 (2)
      utc_time_created = mlflow_timestamp(),
      run_id = mlflow_get_active_run_id_or_start_run(),
      artifact_path = artifact_path,
      flavors = list()
    ),
  ...)
  res <- mlflow log artifact(path = temp path, artifact path = artifact path)
  tryCatch({
    mlflow:::mlflow_record_logged_model(model_spec)
  error = function(e) {
```

- (1) Add a new parameter signature
- (2) Propagate signature to the model_spec parameter when invoking mlflow::mlflow_save_model

Benefit of this method is that all model flavors will inherit the capability to log a signature.

1.2 Working Through the Solution

To keep things simple we'll be logging a "model" (a function which divides by two).

```
half <- function(x) x / 2
half(1:10)</pre>
```

```
[1] 0.5 1.0 1.5 2.0 2.5 3.0 3.5 4.0 4.5 5.0
```

Without any changes, a simplified example of logging to {mlflow} would look like:

(1) As discussed earlier, this is where things start to go awry with respect to Unity Catalog

1.2.1 Patching mlflow_log_model

Note

Technically, patching mlflow_log_model isn't the only way to achieve this fix - you could modify the yaml after it's written.

I won't be showing that method as It's just as tedious and can change depending on the model flavour (with respect to where artifacts may reside), patching is more robust.

```
mlflow_log_model <- function(model, artifact_path, signature = NULL, ...) { (1)</pre>
  format_signature <- function(signature) {</pre>
                                                                               (2)
    lapply(signature, function(x) {
      jsonlite::toJSON(x, auto_unbox = TRUE)
    })
  }
  temp_path <- fs::path_temp(artifact_path)</pre>
  model_spec <- mlflow_save_model(model, path = temp_path, model_spec = list(</pre>
    utc_time_created = mlflow:::mlflow_timestamp(),
    run_id = mlflow:::mlflow_get_active_run_id_or_start_run(),
    artifact_path = artifact_path,
    flavors = list(),
    signature = format_signature(signature)
                                                                               (3)
  ), ...)
  res <- mlflow_log_artifact(path = temp_path, artifact_path = artifact_path)</pre>
    mlflow:::mlflow_record_logged_model(model_spec)
  },
  error = function(e) {
    warning(
      paste("Logging model metadata to the tracking server has failed, possibly due to older
            "server version. The model artifacts have been logged successfully.",
            "In addition to exporting model artifacts, MLflow clients 1.7.0 and above",
            "attempt to record model metadata to the tracking store. If logging to a",
            "mlflow server via REST, consider upgrading the server version to MLflow",
            "1.7.0 or above.", sep=" ")
    )
  })
```

```
res
}

# overriding the function in the existing mlflow namespace
assignInNamespace("mlflow_log_model", mlflow_log_model, ns = "mlflow")
```

- (1) signature has been added to function parameters, it's defaulting to NULL so that existing code won't break
- 2 Adding format_signature function so don't need to write JSON by hand, adding this within function for simplicity
- 3 signature is propagated to mlflow_save_model's model_spec parameter which will write a valid signature

1.2.2 Logging Model with a Signature

```
with(mlflow_start_run(), {
    # typically you'd do more modelling related activities here
    model <- carrier::crate(~half(.x))
    signature <- list(
        inputs = list(list(type = "double", name = "x")),
        outputs = list(list(type = "double"))
    )
    mlflow_log_model(model, "model", signature = signature)
})</pre>
```

- (1) Explicitly defining a signature, a list that contains input and outputs, each are lists of lists respectively
- (2) Passing defined signature to the now patched mlflow_log_model function

1.2.3 Registration to Unity Catalog

Now that the prerequisite of adding a model signature has been satisfied there is one last hurdle to overcome, registering to Unity Catalog.

The hurdle is due to {mlflow} not having been updated yet to support registration to Unity Catalog directly. The easiest way to overcome this is to simply register the run via python.

For example:

```
import mlflow
mlflow.set_registry_uri("databricks-uc")

catalog = "main"
schema = "default"
model_name = "my_model"
run_uri = "runs:/<run_id>/model"

mlflow.register_model(run_uri, f"{catalog}.{schema}.{model_name}")
```

(1) You'll need to either get the run_uri programmatically or copy it manually

To do this with R you'll need to make a series of requests to Unity Catalog endpoints for registering model, the specific steps are:

1. (Optional) Create a new model in Unity Catalog

- POST request on /api/2.0/mlflow/unity-catalog/registered-models/create
 - name: 3 tiered namespace (e.g. main.default.my_model)

2. Create a version for the model

- POST request on /api/2.0/mlflow/unity-catalog/model-versions/create
 - name: 3 tiered namespace (e.g. main.default.my_model)
 - source: URI indicating the location of the model artifacts
 - run_id: run_id from tracking server that generated the model
 - run_tracking_server_id: Workspace ID of run that generated the model
- This will return storage_location and version

3. Copy model artifacts

• Need to copy the artifacts to storage_location from step (2)

4. Finalise the model version

- POST request on /api/2.0/mlflow/unity-catalog/model-versions/finalize
 - name: 3 tiered namespace (e.g. main.default.my_model)
 - version: version returned from step (2)

It's considerably easier to just use Python to register the model at this time.

1.3 Fixing mlflow

Ideally this page wouldn't exist and {mlflow} would support Unity Catalog. Hopefully sometime soon I find the time to make a pull request myself - until then this serves as a guide.

2 Model Serving



Model serving on Databricks is well documented for Python. It is a common misconception that you cannot serve R models on Databricks, you can!

The amount of effort is going to depend heavily on the model flavour. In most occasions the amount of effort is not worth it - but that decision is not mine to make, I will show you how and you can decide for yourself.

2.1 Serving Overview



Work In Progress

2.2 XGBoost



Work In Progress

XGBoost is relatively straightforward since it the artifacts generated when saving the model are universally understood across each language the library is available in.

2.3 ONNX



Work In Progress

2.4 Crate



Work In Progress

Part II Package Management

3 Faster Package Installs

You may have noticed that when installing packages in the notebook it can take a while. It could be minutes, hours in extreme cases, to install the suite of packages your project requires. This is especially tedious if you need to do this every time a job runs, or each morning when your cluster is started.

Clusters are ephemeral and by default have no persistent storage, therefore installed packages will not be available on restart.

By default Databricks installs packages from CRAN. CRAN does not provide pre-compiled binaries for Linux (Databricks clusters' underlying virtual machines are Linux, Ubuntu specifically).

Posit to save the day! Posit provides a public package manager that has all packages from CRAN (and Bioconductor!). There is a helpful wizard to get started.

With our new found knowledge we can make installing R packages within Databricks significantly faster. There are multiple ways to solve this, each differing slightly, but fundamentally the same.

3.1 Setting Repo within Notebook

The quickest method is to follow the wizard and adjust the repos option:

```
# need to set the user agent string otherwise installs will be slow
# e.g. selecting Ubuntu 22.04 in wizard
options(
   HTTPUserAgent = sprintf("R/%s R (%s)", getRversion(), paste(getRversion(), R.version["platerepos = "https://packagemanager.posit.co/cran/__linux__/jammy/latest"))
```

This works well but not all versions of the Databricks Runtime use the same version of Ubuntu.

It's easier to detect the Ubuntu release code name dynamically:

```
release <- system("lsb_release -c --short", intern = T)

# need to set the user agent string otherwise installs will be slow
options(
   HTTPUserAgent = sprintf("R/%s R (%s)", getRversion(), paste(getRversion(), R.version["platerepos = paste0("https://packagemanager.posit.co/cran/__linux__/", release, "/latest")
)</pre>
```

1 system is used to run the command to retrieve the release code name

The downside of this method is that it requires every notebook to adjust the repos and HTTPUserAgent options.

3.2 Cluster Environment Variable & Init Script

Databricks clusters allow specification of environment variables, there is a specific variable (DATABRICKS_DEFAULT_R_REPOS) that can be set to adjust the default repository for the entire cluster.

You can again refer to the wizard, the environment variables section of cluster should be:

```
DATABRICKS_DEFAULT_R_REPOS=<posit-package-manager-url-goes-here>
```

Unfortunately this isn't as dynamic as the first option and you still need to set the HTTPUserAgent in Rprofile.site via an init script.

The init script will be:

```
#!/bin/bash
# Append changes to Rprofile.site
cat <<EOF >> "/etc/R/Rprofile.site"
options(
   HTTPUserAgent = sprintf("R/%s R (%s)", getRversion(), paste(getRversion(), R.version["plate"))
EOF
```

! Important

Due to how Databricks starts up the R shell for notebook sessions it's not straightforward to adjust the repos option in an init script alone.

 ${\tt DATABRICKS_DEFAULT_R_REPOS} \ \ {\tt is} \ \ {\tt referenced} \ \ {\tt as} \ \ {\tt part} \ \ {\tt of} \ \ {\tt the} \ \ {\tt startup} \ \ {\tt process} \ \ {\tt after}$

Rprofile.site is executed and will override any earlier attempt to adjust repos. Therefore you'll need to use both the init script and the environment variable configuration.

3.3 Setting Repo for Cluster Library

i Note

Similar to setting <code>DATABRICKS_DEFAULT_R_REPOS</code> this requires the <code>HTTPUserAgent</code> also to be set and it's unlikely to be helpful other than for it's purpose of installing a package to make it available for all cluster users.

Cluster libraries can install R packages and support specification of the repository.

4 Persisting Packages



{renv}

♦ Under Development

Part III Data Engineering

6 {dbplyr} & {odbc}

♦ Under Development

Part IV Miscellaneous

7 {htmlwidgets} in notebooks

When you try to view a {htmlwidget} based visualisation (e.g. {leaflet}) in a Databricks notebook you'll find there is no rendered output by default.

The Databricks documentation details how to get this working but requires specification of the workspace URL explicitly.

{brickster} has a helper function that simplifies enabling {htmlwidgets}:

```
remotes::install_github("zacdav-db/brickster")
brickster::notebook_enable_htmlwidgets()
```

The function itself is straightforward, here is code (simplified version of brickster::notebook_enable_htmlwidge) that doesn't require installing {brickster}:

```
enable_htmlwidgets <- function(height = 450) {</pre>
  # new option to control default widget height, default is 450px
  options(db_htmlwidget_height = height)
                                                                                (1)
 system("apt-get --yes install pandoc", intern = T)
                                                                                (2)
 if (!base::require("htmlwidgets")) {
                                                                                (3)
    utils::install.packages("htmlwidgets")
 # new method will fetch height based on new option, or default to 450px
 new_method <- function(x, ...) {</pre>
                                                                                (4)
    x$height <- getOption("db_htmlwidget_height", 450)
    file <- tempfile(fileext = ".html")</pre>
    htmlwidgets::saveWidget(x, file = file)
    contents <- as.character(rvest::read_html(file))</pre>
    displayHTML(contents)
 }
 utils::assignInNamespace("print.htmlwidget", new_method, ns = "htmlwidgets") (5)
  invisible(list(default_height = height, print = new_method))
```

}

- (1) The height of the htmlwidget output is controlled via an option (db_htmlwidget_height), this allows the height to be adjusted without re-running the function
- 2 Installing pandoc as it's required to use htmlwidgets::saveWidget
- 3 Ensure that {htmlwidgets} is installed
- (4) Function that writes the widget to a temporary file as a self-contained html file and then reads the contents and presents via displayHTML
- (5) Override the htmlwidgets::print.htmlwidget method

8 OAuth {odbc}

When using {odbc} to connect to Databricks clusters and SQL warehouses you'll likely have used a personal access token (PAT). It's not uncommon for workspace administrators to disable the use of PATs.

If you are unable to create a PAT you are still able to connect to Databricks but you'll need to use OAuth (either M2M or U2M).

User-to-machine (U2M) is typically what you'd want to use. Good news, the Databricks ODBC driver supports both since 2.7.5.

8.1 U2M Example

Note

OAuth U2M or OAuth 2.0 browser-based authentication works only with applications that run locally. It does not work with server-based or cloud-based applications.

When running this code you should be prompted to login to the workspace or you'll see a window that says "success". You can close the window and continue working in R.

```
library(odbc)
library(DBI)

con <- DBI::dbConnect(
    drv = odbc::databricks(),
    httpPath = "/sql/1.0/warehouses/<warehouse-id>",
    workspace = "<workspace-name>.cloud.databricks.com",
    authMech = 11,
    auth_flow = 2
)
```

- (1) {odbc} recently added odbc::databricks() to simplify connecting to Databricks (requires version >=1.4.0)
- (2) The httpPath can be found in the 'Connection Details' tab of a SQL warehouse

- (3) workspace refers to the workspace URL, also found in 'Connection Details' tab as 'Server hostname'
- 4 The docs mention setting AuthMech to 11 and Auth_Flow to 2