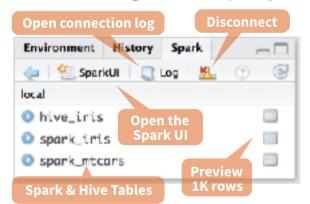
# Data Science in Spark with Sparklyr:: CHEAT SHEET

### Intro

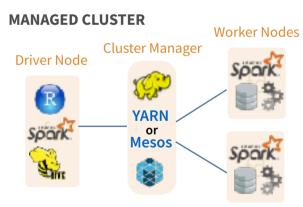
**sparklyr** is an R interface for **Apache Spark**™, it provides a complete **dplyr** backend and the option to query directly using Spark SQL statement. With sparklyr, you can orchestrate distributed machine learning using either Spark's MLlib or H2O Sparkling Water.

Starting with version 1.044, RStudio Desktop, Server and Pro include integrated support for the sparklyr package. You can create and manage connections to Spark clusters and local Spark instances from inside the IDE.

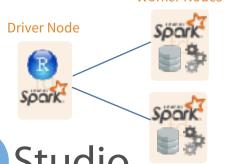
#### **RStudio Integrates with sparklyr**



### **Cluster Deployment**



#### **STAND ALONE CLUSTER** Worker Nodes



## Data Science Toolchain with Spark + sparklyr

#### **Import**

- Export an R DataFrame
- Read a file
- Read existing Hive table

R for Data Science, Grolemund & Wickham

**LOCAL MODE** (No cluster required)

1. Install a local version of Spark:

ON A MESOS MANAGED CLUSTER

sc <- spark connect (master = "local")</pre>

1. Install RStudio Server or Pro on one of the

2. Locate path to the cluster's Spark directory

spark connect(master="[mesos URL]",

version = "1.6.2", spark\_home =

1. The Livy REST application should be

sc <- spark\_connect(method = "livy",</pre>

spark\_install ("2.0.1")

2. Open a connection

existing nodes

3. Open a connection

[Cluster's Spark path])

**USING LIVY** (Experimental)

running on the cluster

master = "http://host:port")

2. Connect to the cluster

**Getting Started** 

- dplyr verb
- Direct Spark SQL (DBI)

#### SDF function (Scala API)

#### Wrangle

### R for plotting Model

**Transform** 

Transformer

function

- Spark MLlib
- **H20 Extension**

**Understand** 

### Using sparklyr

**Communicate** 

Collect data

Share plots,

documents,

and apps

into R



A brief example of a data analysis using Apache Spark, R and sparklyr in local mode

library(sparklyr); library(dplyr); library(ggplot2); library(tidyr); **Install Spark locally** set.seed(100)

spark install("2.0.1")

**Connect to local version** 

import iris <- copy to(sc, iris, "spark iris", overwrite = TRUE)

sc <- spark connect(master = "local")</pre>

Copy data to Spark memory

partition iris <- sdf partition(</pre> import iris,training=0.5, testing=0.5)

model\_iris <- tidy\_iris %>%

pred\_iris <- sdf\_predict(</pre>

test\_iris <- tbl(sc,"spark\_iris\_test")

**Partition** 

Spark table

sdf register(partition iris, c("spark\_iris\_training","spark\_iris\_test"))

tidy\_iris <- tbl(sc,"spark\_iris\_training") %>%

select(Species, Petal\_Length, Petal\_Width)

ml\_decision\_tree(response="Species",

features=c("Petal Length","Petal Width"))

Create a hive metadata for each partition

#### **ON A SPARK STANDALONE CLUSTER**

- 2. Install a local version of Spark: spark\_install (version = "2.0.1")
- 3. Open a connection host:port", version = "2.0.1",

### **ON A YARN MANAGED CLUSTER**

**Visualize** 

Collect data into

- 1. Install RStudio Server or RStudio Pro on one of the existing nodes, preferably an edge node
- 2. Locate path to the cluster's Spark Home Directory, it normally is "/usr/lib/spark"
- 3. Open a connection spark connect(master="yarn-client", version = "1.6.2", spark home = [Cluster's Spark path])

- 1. Install RStudio Server or RStudio Pro on one of the existing nodes or a server in the same LAN
  - spark\_connect(master="spark:// spark home = spark home dir())

### **Tuning Spark**

#### **EXAMPLE CONFIGURATION**

config <- spark config() config\$spark.executor.cores <- 2 config\$spark.executor.memory <- "4G" sc <- spark connect (master="yarn-client", config = config, version = "2.0.1")

#### **IMPORTANT TUNING PARAMETERS** with defaults

- spark.yarn.am.cores
- spark.executor.instances
- spark.executor.memory 1g

spark.executor.cores 1

- sparklyr.shell.driver-memory
- spark.yarn.am.memory 512m spark.executor.extraJavaOptions
- spark.network.timeout 120s spark.executor.heartbeatInterval 10s
  - sparklyr.shell.executor-memory

model\_iris, test\_iris) %>% Bring data back into R memory for plotting

pred iris %>%

collect

inner\_join(data.frame(prediction=0:2, lab=model\_iris\$model.parameters\$labels)) %>% ggplot(aes(Petal\_Length, Petal\_Width, col=lab)) + geom\_point()

spark disconnect(sc)

**Disconnect** 

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### Reactivity

#### **COPY A DATA FRAME INTO SPARK**

sdf\_copy\_to(sc, iris, "spark\_iris")

sdf\_copy\_to(sc, x, name, memory, repartition,
overwrite)

#### **IMPORT INTO SPARK FROM A FILE**

Arguments that apply to all functions:

sc, name, path, options = list(), repartition = 0, memory = TRUE, overwrite = TRUE

CSV

spark\_read\_csv( header = TRUE,
columns = NULL, infer\_schema = TRUE,
delimiter = ",", quote = "\"", escape = "\\",
charset = "UTF-8", null\_value = NULL)

**JSON** 

spark\_read\_json()

PARQUET spark\_read\_parquet()

#### **SPARK SQL COMMANDS**

DBI::dbWriteTable(sc, "spark\_iris", iris)

DBI::dbWriteTable(conn, name, value)

#### FROM A TABLE IN HIVE

my\_var <- tbl\_cache(sc, name=
"hive\_iris")</pre>

tbl\_cache(sc, name, force = TRUE)
Loads the table into memory

my\_var <- **dplyr::tbl**(sc, name= "hive\_iris")

dplyr::**tbl(**scr, ...)

Creates a reference to the table without loading it into memory

### Visualize & Communicate

#### **DOWNLOAD DATA TO R MEMORY**

r\_table <- **collect**(my\_table) plot(Petal\_Width~Petal\_Length, data=r\_table)

dplyr::**collect(**x**)** 

Download a Spark DataFrame to an R DataFrame

sdf\_read\_column(x, column)

Returns contents of a single column to R

#### **SAVE FROM SPARK TO FILE SYSTEM**

Arguments that apply to all functions: x, path

CSV

spark\_read\_csv( header = TRUE,
delimiter = ",", quote = "\"", escape = "\\",
charset = "UTF-8", null\_value = NULL)

**JSON** 

spark\_read\_json(mode = NULL)

PARQUET spark\_read\_parquet(mode = NULL)

# Wrangle

#### **SPARK SOL VIA DPLYR VERBS**

Translates into Spark SQL statements

my\_table <- my\_var %>%
 filter(Species=="setosa") %>%
 sample\_n(10)

#### **DIRECT SPARK SQL COMMANDS**

my\_table <- DBI::**dbGetQuery**( sc , "SELECT \* FROM iris LIMIT 10")

DBI::dbGetQuery(conn, statement)

#### **SCALA API VIA SDF FUNCTIONS**

sdf mutate(.data)

Works like dplyr mutate function

sdf\_partition(x, ..., weights = NULL, seed =
sample (.Machine\$integer.max, 1))
sdf\_partition(x, training = 0.5, test = 0.5)

sdf\_register(x, name = NULL)

Gives a Spark DataFrame a table name

sdf\_sample(x, fraction = 1, replacement =
TRUE, seed = NULL)

sdf\_sort(x, columns)

Sorts by >=1 columns in ascending order

sdf\_with\_unique\_id(x, id = "id")

sdf\_predict(object, newdata)

Spark DataFrame with predicted values

#### **ML TRANSFORMERS**

**ft\_binarizer**(my\_table,input.col="Petal\_Le ngth", output.col="petal\_large", threshold=1.2)

Arguments that apply to all functions: x, input.col = NULL, output.col = NULL

ft\_binarizer(threshold = 0.5)
Assigned values based on threshold

ft\_bucketizer(splits)

Numeric column to discretized column

ft\_discrete\_cosine\_transform(inverse
= FALSE)

Time domain to frequency domain

ft\_elementwise\_product(scaling.col)
Element-wise product between 2 cols

 $ft\_index\_to\_string()$ 

Index labels back to label as strings

ft\_one\_hot\_encoder()

Continuous to binary vectors

**ft\_quantile\_discretizer(**n.buckets=5L**)**Continuous to binned categorical
values

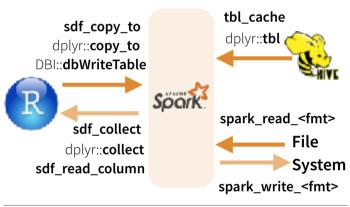
ft\_sql\_transformer(sql)

ft\_string\_indexer( params = NULL)
 Column of labels into a column of label
 indices.

ft vector assembler()

Combine vectors into single row-vector

### Reading & Writing from Apache Spark



### **Extensions**

Create an R package that calls the full Spark API & provide interfaces to Spark packages.

#### **CORE TYPES**

spark\_connection() Connection between R and the
Spark shell process

spark\_jobj() Instance of a remote Spark object
spark\_dataframe() Instance of a remote Spark
DataFrame object

#### **CALL SPARK FROM R**

invoke() Call a method on a Java object
invoke\_new() Create a new object by invoking a
constructor

invoke\_static() Call a static method on an object

#### MACHINE LEARNING EXTENSIONS

ml\_create\_dummy\_variables() ml\_options()
ml\_prepare\_dataframe() ml\_model()
ml\_prepare response features intercept()

### Model (MLlib)

ml\_decision\_tree(my\_table,
 response = "Species", features =
 c("Petal\_Length", "Petal\_Width"))



ml\_als\_factorization(x, user.column = "user",
 rating.column = "rating", item.column = "item",
 rank = 10L, regularization.parameter = 0.1, iter.max = 10L,
 ml.options = ml\_options())

ml\_generalized\_linear\_regression(x, response, features,
 intercept = TRUE, family = gaussian(link = "identity"), iter.max =
 100L, ml.options = ml\_options())

ml\_kmeans(x, centers, iter.max = 100, features = dplyr::tbl\_vars(x),
 compute.cost = TRUE, tolerance = 1e-04, ml.options = ml\_options())

 $ml_lda(x, features = dplyr::tbl_vars(x), k = length(features), alpha = (50/k) + 1, beta = 0.1 + 1, ml.options = ml_options())$ 

ml\_linear\_regression(x, response, features, intercept = TRUE, alpha = 0, lambda = 0, iter.max = 100L, ml.options = ml\_options()) Same options for: ml\_logistic\_regression

ml\_multilayer\_perceptron(x, response, features, layers, iter.max =
 100, seed = sample(.Machine\$integer.max, 1), ml.options =
 ml\_options())

ml\_naive\_bayes(x, response, features, lambda = 0, ml.options =
 ml\_options())

ml\_one\_vs\_rest(x, classifier, response, features, ml.options =
 ml\_options())

ml\_pca(x, features = dplyr::tbl\_vars(x), ml.options = ml\_options())

ml\_random\_forest(x, response, features, max.bins = 32L,
 max.depth = 5L, num.trees = 20L, type = c("auto", "regression",
 "classification"), ml.options = ml\_options())

ml\_survival\_regression(x, response, features, intercept =
 TRUE,censor = "censor", iter.max = 100L, ml.options = ml\_options())

ml\_binary\_classification\_eval(predicted\_tbl\_spark, label, score,
 metric = "areaUnderROC")

ml\_classification\_eval(predicted\_tbl\_spark, label, predicted\_lbl,
 metric = "f1")

ml\_tree\_feature\_importance(sc, model)





