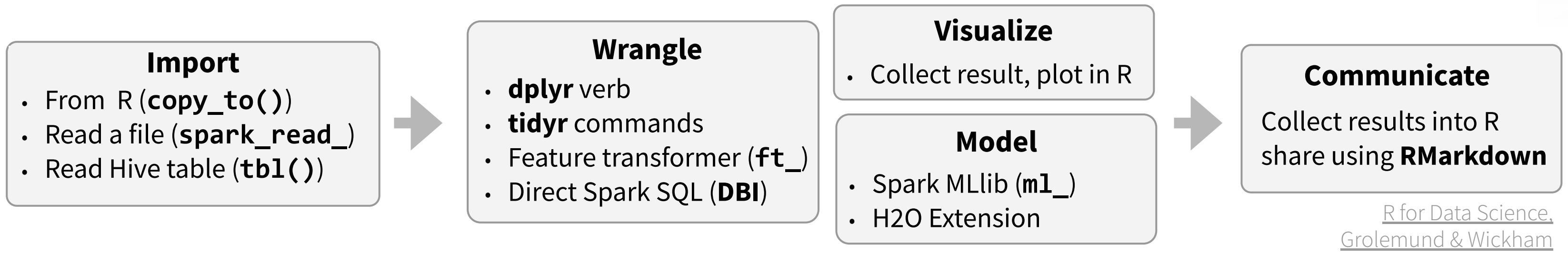
Data Science in Spark with *sparklyr* : : CHEAT SHEET

Intro

***sparklyr* is an R interface for Apache Spark™.** It enables us to write all of our analysis code in R, but have the actual processing happen inside Spark clusters. Easily manipulate and model large-scale using R and Spark via *sparklyr.*


Import
- From R (**copy_to()**)
- Read a file (**spark_read_**)
- Read Hive table (**tbl()**)

Wrangle
- **dplyr** verb
- **tidyr** commands
- Feature transformer (**ft_**)
- Direct Spark SQL (**DBI**)

Visualize
- Collect result, plot in R

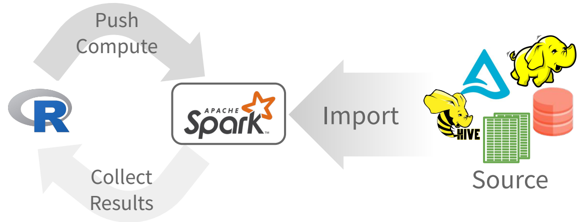
Model
- Spark MLlib (**ml_**)
- H2O Extension

Communicate

Collect results into R share using **RMarkdown**

R for Data Science, Grolemund & Wickham

Import



Import data into Spark, not R

READ A FILE INTO SPARK

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()**
TEXT	**spark_read_text()**
ORC	**spark_read_orc()**
LIBSVM	**spark_read_libsvm()**
DELTA	**spark_read_delta()**
AVRO	**spark_read_avro()**

R DATA FRAME INTO SPARK

dplyr::**copy_to**(dest, df, name**)**

Apache Arrow accelerates data transfer between R and Spark. To use, simply load the library

```
library(sparklyr)
library(arrow)
```

FROM A TABLE IN HIVE

dplyr::**tbl**(scr, …) - Creates a reference to the table without loading it into memory

Wrangle

DPLYR VERBS

Translates into Spark SQL statements

```
copy_to(sc, mtcars) %>%
      mutate(trm = ifelse(am == 0,
                "auto", "man")) %>%
      group_by(trm) %>%
      summarise_all(mean)
```

TIDYR

pivot_longer() - Collapse several columns into two.

pivot_wider() - Expand two columns into several.

nest() / unnest() - Convert groups of cells into list-columns, and vice versa.

unite() / separate() - Split a single column into several columns, and vice versa.

fill() - Fill NA with the previous value

FEATURE TRANSFORMERS

ft_binarizer() - Assigned values based on threshold

ft_bucketizer() - Numeric column to discretized column

ft_count_vectorizer() - Extracts a vocabulary from document

ft_discrete_cosine_transform() - 1D discrete cosine transform of a real vector

ft_elementwise_product() - Element-wise product between 2 cols

ft_hashing_tf() - Maps a sequence of terms to their term frequencies using the hashing trick.

ft_idf() - Compute the Inverse Document Frequency (IDF) given a collection of documents.

ft_imputer() - Imputation estimator for completing missing values, uses the mean or the median of the columns.

ft_index_to_string() - Index labels back to label as strings

ft_interaction() - Takes in Double and Vector columns and outputs a flattened vector of their feature interactions.

ft_max_abs_scaler() - Rescale each feature individually to range [-1, 1]

ft_min_max_scaler() - Rescale each feature to a common range [min, max] linearly

ft_ngram() - Converts the input array of strings into an array of n-grams

ft_bucketed_random_projection_lsh()
ft_minhash_lsh() - Locality Sensitive Hashing functions for Euclidean distance and Jaccard distance (MinHash)

ft_normalizer() - Normalize a vector to have unit norm using the given p-norm

ft_one_hot_encoder()- Continuous to binary vectors

ft_pca() - Project vectors to a lower dimensional space of top k principal components.

ft_quantile_discretizer() - Continuous to binned categorical values.

ft_regex_tokenizer() - Extracts tokens either by using the provided regex pattern to split the text.

ft_robust_scaler() - Removes the median and scales according to standard scale.

ft_standard_scaler() - Removes the mean and scaling to unit variance using column summary statistics

ft_stop_words_remover() - Filters out stop words from input

ft_string_indexer() - Column of labels into a column of label indices.

ft_tokenizer() - Converts to lowercase and then splits it by white spaces

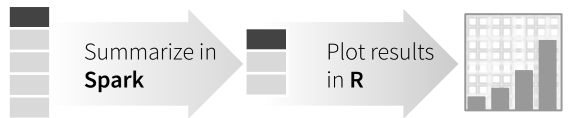
ft_vector_assembler() - Combine vectors into single row-vector

ft_vector_indexer() - Indexing categorical feature columns in a dataset of Vector

ft_vector_slicer() - Takes a feature vector and outputs a new feature vector with a subarray of the original features

ft_word2vec() - Word2Vec transforms a word into a code

Visualize



DPLYR + GGPLOT2

```
copy_to(sc, mtcars) %>%
  group_by(cyl) %>%
  summarise(mpg_m = mean(mpg)) %>%
  collect() %>%
  ggplot() +
    geom_col(aes(cyl, mpg_m))
```

Summarize in Spark
Collect results in R
Create plot

RStudio

RStudio® is a trademark of RStudio, Inc. • • info@rstudio.com • 844-448-1212 • rstudio.com • Learn more at **spark.rstudio.com** • sparklyr 1.7.5 • Updated: 2022-02

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Modeling

REGRESSION

ml_linear_regression() - Linear regression.

ml_aft_survival_regression() - Parametric survival regression model named accelerated failure time (AFT) model

ml_generalized_linear_regression() - GLM

ml_isotonic_regression() - Currently implemented using parallelized pool adjacent violators algorithm. Only univariate (single feature) algorithm supported

ml_random_forest_regressor() - Regression using random forests.

CLASSIFICATION

ml_linear_svc() - Classification using linear support vector machines

ml_logistic_regression() - Logistic regression

ml_multilayer_perceptron_classifier() - Classification model based on the Multilayer Perceptron.

ml_naive_bayes() - It supports Multinomial NB which can handle finitely supported discrete data

ml_one_vs_rest() - Reduction of Multiclass Classification to Binary Classification. Performs reduction using one against all strategy.

TREE

ml_decision_tree_classifier()|**ml_decision_tree()** |**ml_decision_tree_regressor()** - Classification and regression using decision trees

ml_gbt_classifier()|**ml_gradient_boosted_trees()** | **ml_gbt_regressor()** - Binary classification and regression using gradient boosted trees

ml_random_forest_classifier() - Classification and regression using random forests.

ml_feature_importances() | **ml_tree_feature_importance()** - Feature Importance for Tree Models

CLUSTERING

ml_bisecting_kmeans() - A bisecting k-means algorithm based on the paper

ml_lda() | **ml_describe_topics()** | **ml_log_likelihood()** | **ml_log_perplexity()** | **ml_topics_matrix()** - LDA topic model designed for text documents.

ml_gaussian_mixture() - Expectation maximization for multivariate Gaussian Mixture Models (GMMs)

ml_kmeans() | **ml_compute_cost()** |**ml_compute_silhouette_measure()** - Clustering with support for k-means

ml_power_iteration() - For clustering vertices of a graph given pairwise similarities as edge properties.

FEATURE

ml_chisquare_test(x,features,label) - Pearson's independence test for every feature against the label

ml_default_stop_words() - Loads the default stop words for the given language

STATS

ml_summary() - Extracts a metric from the summary object of a Spark ML model

ml_corr() - Compute correlation matrix

RECOMMENDATION

ml_als() | **ml_recommend()** - Recommendation using Alternating Least Squares matrix factorization

EVALUATION

ml_clustering_evaluator() - Evaluator for clustering

ml_evaluate() - Compute performance metrics

ml_binary_classification_evaluator() | **ml_binary_classification_eval()** | **ml_classification_eval()** - A set of functions to calculate performance metrics for prediction models.

FREQUENT PATTERN

ml_fpgrowth() | **ml_association_rules()** | **ml_freq_itemsets()** - A parallel FP-growth algorithm to mine frequent itemsets.

ml_freq_seq_patterns() | **ml_prefixspan()** - PrefixSpan algorithm for mining frequent itemsets.

UTILITIES

ml_call_constructor() - Identifies the associated sparklyr ML constructor for the JVM

ml_model_data() - Extracts data associated with a Spark ML model

ml_standardize_formula() - Generates a formula string from user inputs, to be used in `ml_model` constructor

ml_uid() - Extracts the UID of an ML object.

ML Pipelines

*Easily create a formal Spark Pipeline models using R. Save the Pipeline in native Sacala. The saved model will have **no dependencies** on R.*

INITIALIZE AND TRAIN

ml_pipeline() - Initializes a new Spark Pipeline

ml_fit() - Trains the model, outputs a Spark Pipeline Model.

SAVE AND RETRIEVE

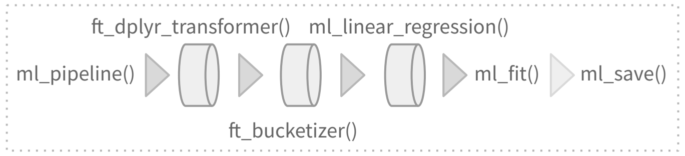
ml_save() - Saves into a format that can be read by Scala and PySpark .

ml_read() - Reads Spark object into sparklyr.

SQL AND DPLYR

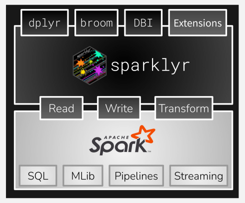
ft_sql_transformer() - Creates a Pipeline step based on the SQL statement passed to the command.

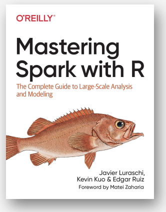
ft_dplyr_transformer() - Creates a Pipeline step based on one or several dplyr commands.



spark.rstudio.com/guides/pipelines

More Info




O'REILLY
Mastering Spark with R
The Complete Guide to Large-Scale Analysis and Modeling
Javier Luraschi, Kevin Kuo & Edgar Ruiz
Foreword by Matei Zaharia

spark.rstudio.com   therinspark.com

Sessions

YARN CLIENT

1. Install RStudio Server on an edge node
2. Locate path to the cluster's Spark Home Directory, it normally is **"/usr/lib/spark"**
3. Basic configuration example

```
conf <- spark_config()
conf$spark.executor.memory <- "300M"
conf$spark.executor.cores <- 2
conf$spark.executor.instances <- 3
conf$spark.dynamicAllocation.enabled<-"false"
```

4. Open a connection

```
sc <- spark_connect(master = "yarn",
          spark_home = "/usr/lib/spark/",
          version = "2.1.0", config = conf)
```

YARN CLUSTER

1. Make sure to have copies of the **yarn-site.xml** and **hive-site.xml** files in the RStudio Server
2. Point environment variables to the correct paths

```
Sys.setenv(JAVA_HOME="[Path]")
Sys.setenv(SPARK_HOME ="[Path]")
Sys.setenv(YARN_CONF_DIR ="[Path]")
```

3. Open a connection

```
sc <- spark_connect(master = "yarn-cluster")
```

STANDALONE CLUSTER

1. Install RStudio Server on one of the existing nodes or a server in the same LAN
2. Open a connection

```
spark_connect(master="spark://host:port",
   version = "2.0.1",
   spark_home = [path to Spark])
```

LOCAL MODE

No cluster required. Use for learning purposes only

1. Install a local version of Spark: `spark_install()`
2. Open a connection

```
sc <- spark_connect(master="local")
```

KUBERNETES

1. Use the following to obtain the Host and Port

```
system2("kubectl", "cluster-info")
```

2. Open a connection

```
sc <- spark_connect(config =
     spark_config_kubernetes(
      "k8s://https://[HOST]>:[PORT]",
       account = "default",
       image = "docker.io/owner/repo:version"))
```

CLOUD

Databricks - `spark_connect(method = "databricks")`

Qubole- `spark_connect(method = "qubole")`

RStudio