

SMOKE TEST DOCUMENT

Spark Smoke Test Cases

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1 NOTE

Set SPARK_HOME to \$PATH or “cd /usr/lib/spark/spark-2.4.0-bin-hadoop2.7” and run the following examples

All the Spark examples related to Python, Scala, Java & R are under this location
“/usr/lib/spark/spark-2.4.0-bin-hadoop2.7/examples/src/main”

Replace <Spark-master-IP> with the actual IP address

Reference link: <https://spark.apache.org/docs/latest/submitting-applications.html>

2 SAMPLE TEST CASE FOR SPARK-SUBMIT

- Run application locally on 8 cores

```
./bin/spark-submit \  
--class org.apache.spark.examples.SparkPi \  
--master local[8] \  
/usr/lib/spark/spark-2.4.0-bin-hadoop2.7/examples/jars/spark-examples_2.11-2.4.0.jar \  
100
```

- Run below command on a Spark standalone cluster in client deploy mode

```
./bin/spark-submit \  
--class org.apache.spark.examples.SparkPi \  
--master spark://<Spark-master-IP>:7077 \  
--executor-memory 20G \  
--total-executor-cores 100 \  
/usr/lib/spark/spark-2.4.0-bin-hadoop2.7/examples/jars/spark-examples_2.11-2.4.0.jar \  
1000
```

- Run below command on a Spark standalone cluster in cluster deploy mode with supervise

```
./bin/spark-submit \  
--class org.apache.spark.examples.SparkPi \  
--master spark://<Spark-master-IP>:7077 \  
--deploy-mode cluster \  
--supervise \  

```

- Run a python application on a Spark standalone cluster

```
./bin/spark-submit \  
--class org.apache.spark.examples.SparkPi \  
--master spark://<Spark-master-IP>:7077 \  
--deploy-mode cluster \  
--supervise \  

```

3 SAMPLE TEST CASE FOR SPARK-SHELL

We will first introduce the API through Spark's interactive shell (in Python or Scala) and then show how to write applications in Java, Scala and Python.

3.1 Interactive analysis with the Spark Shell

Spark's shell provides a simple way to learn the API, as well as a powerful tool to analyze data interactively. It is available in either Scala (which runs on the Java VM and is thus a good way to use existing Java libraries) or Python.

Start it by running the following in the Spark directory

```
./bin/spark-shell
```

- Make a new Dataset from the text of the README file in the Spark source directory:

```
scala> val textFile = spark.read.textFile("README.md")
```

```
textFile: org.apache.spark.sql.Dataset[String] = [value: string]
```

- Get values from Dataset directly, by calling some actions, or transform the Dataset to get new one

```
scala> textFile.count() // Number of items in this Dataset
```

```
res0: Long = 126 // May be different from yours as README.md
```

```
scala> textFile.first() // First item in this Dataset
```

```
res1: String = # Apache Spark
```

- Transform this Dataset into a new one

```
scala> val linesWithSpark = textFile.filter(line => line.contains("Spark"))
```

```
linesWithSpark: org.apache.spark.sql.Dataset[String] = [value: string]
```

- Chain together transformations and actions

```
scala> textFile.filter(line => line.contains("Spark")).count() // How many lines contain  
"Spark"?
```

```
res3: Long = 15
```

3.2 Caching operations on Spark Shell

Spark also supports pulling data sets into a cluster-wide in-memory cache. This is very useful when data is accessed repeatedly.

```
scala> linesWithSpark.cache()
```

```
res7: linesWithSpark.type = [value: string]
```

```
scala> linesWithSpark.count()
```

```
res8: Long = 15
```

```
scala> linesWithSpark.count()
```

```
res9: Long = 15
```

Reference Link: <https://spark.apache.org/docs/latest/quick-start.html#basics>

3.3 Example for Scala Word Count program:

Following are the commands that we shall use for Word Count Example in Spark Shell

- Using Spark context variable, sc to read a text file

```
scala> sc.textFile("usr/lib/spark/spark-2.4.0-bin-hadoop2.7/word.txt")
```

- Split each line using space " " as separator

```
scala> flatMap(line => line.split(" "))
```

- Map each word to a tuple (word, 1), 1 being the number of occurrences of word

```
scala> map(word => (word,1))
```

- Reduce all the words based on Key

```
scala> var counts = map.reduceByKey(_ + _);
```

- Save counts to local file

```
scala> counts.saveAsTextFile("usr/lib/spark/spark-2.4.0-bin-hadoop2.7/result.txt")
```

4 TEST CASES FOR JUPYTERHUB

- Create a Linux user on the master controller node or login as AD user.
- Login to Jupyterhub.

Note: All the Spark examples related to Python, Scala, Java & R are under this location
“/usr/lib/spark/spark-2.4.0-bin-hadoop2.7/examples/src/main/”

4.1 Spark Scala testing

Start a toree scala kernel -> Wait till kernel creates a spark shell. Run following Pearson's correlation. You can run up to 4 Spark shells with current configurations. If your shell doesn't start, you may have used up all the cores. Kill unused Kernels to release resources

Code: Running Pearson's correlation using mllib

Note: You can copy the sample code below from this link: <https://spark.apache.org/docs/latest/mllib-statistics.html>

```
import org.apache.spark.mllib.linalg._
```

```
import org.apache.spark.mllib.stat.Statistics
```

```
import org.apache.spark.rdd.RDD
```

```
val seriesX: RDD[Double] = sc.parallelize(Array(1, 2, 3, 3, 5)) // a series
```

```
// must have the same number of partitions and cardinality as seriesX
```

```
val seriesY: RDD[Double] = sc.parallelize(Array(11, 22, 33, 33, 555))
```

```
// compute the correlation using Pearson's method. Enter "spearman" for Spearman's method. If a
```

```
// method is not specified, Pearson's method will be used by default.
```

```
val correlation: Double = Statistics.corr(seriesX, seriesY, "pearson")
```

```
println(s"Correlation is: $correlation")

val data: RDD[Vector] = sc.parallelize(

Seq(

  Vectors.dense(1.0, 10.0, 100.0),

  Vectors.dense(2.0, 20.0, 200.0),

  Vectors.dense(5.0, 33.0, 366.0))

) // note that each Vector is a row and not a column

// calculate the correlation matrix using Pearson's method. Use "spearman" for Spearman's method

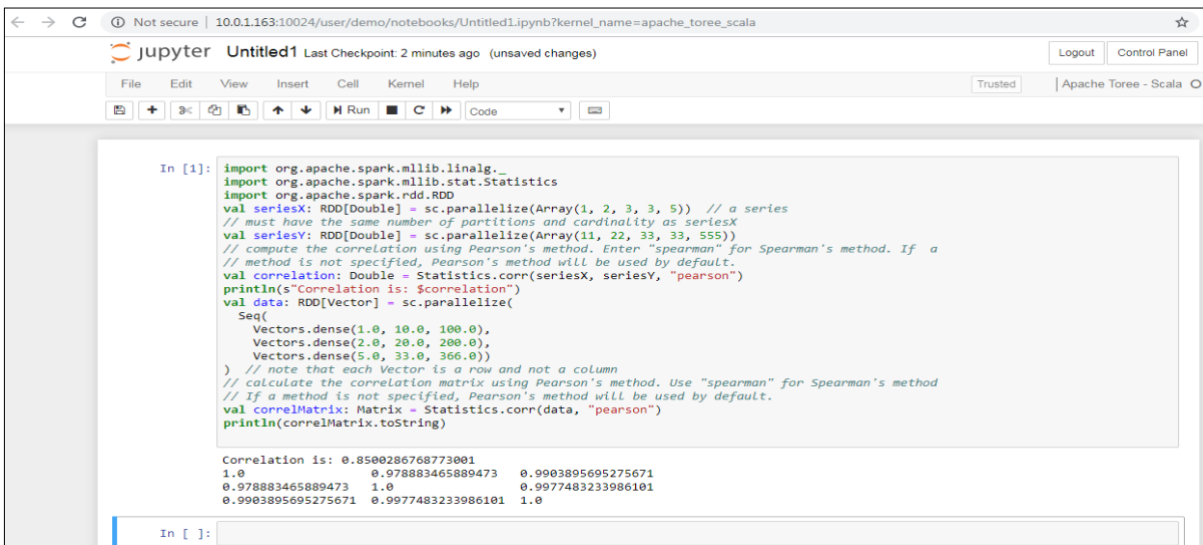
// If a method is not specified, Pearson's method will be used by default.

val correlMatrix: Matrix = Statistics.corr(data, "pearson")

println(correlMatrix.toString)
```

Input: Input is generated within the code. No external input is provided.

Output: Sample output is as given below.



```
In [1]: import org.apache.spark.mllib.linalg._
import org.apache.spark.mllib.stat.Statistics
import org.apache.spark.rdd.RDD
val seriesX: RDD[Double] = sc.parallelize(Array(1, 2, 3, 3, 5)) // a series
// must have the same number of partitions and cardinality as seriesX
val seriesY: RDD[Double] = sc.parallelize(Array(11, 22, 33, 33, 55))
// compute the correlation using Pearson's method. Enter "spearman" for Spearman's method. If a
// method is not specified, Pearson's method will be used by default.
val correlation: Double = Statistics.corr(seriesX, seriesY, "pearson")
println(s"Correlation is: $correlation")
val data: RDD[Vector] = sc.parallelize(
  Seq(
    Vectors.dense(1.0, 10.0, 100.0),
    Vectors.dense(2.0, 20.0, 200.0),
    Vectors.dense(5.0, 33.0, 366.0))
) // note that each Vector is a row and not a column
// calculate the correlation matrix using Pearson's method. Use "spearman" for Spearman's method
// If a method is not specified, Pearson's method will be used by default.
val correlMatrix: Matrix = Statistics.corr(data, "pearson")
println(correlMatrix.toString)
```

Correlation is: 0.8500286768773001

1.0	0.978883465889473	0.9903895695275671
0.978883465889473	1.0	0.9977483233986101
0.9903895695275671	0.9977483233986101	1.0

4.2 PySpark testing

Start a toree pySpark kernel -> Wait till kernel creates a spark shell. You can run up to 4 Spark shells with current configurations. If your shell doesn't start, you may have used up all the cores. Kill unused Kernels to release resources.

Code:

```
from pyspark.mllib.linalg import Matrices, Vectors
```

```
from pyspark.mllib.regression import LabeledPoint
```

```
from pyspark.mllib.stat import Statistics
```

```
vec = Vectors.dense(0.1, 0.15, 0.2, 0.3, 0.25) # a vector composed of the frequencies of events
```

```
# compute the goodness of fit. If a second vector to test against
```

```
# is not supplied as a parameter, the test runs against a uniform distribution.
```

```
goodnessOfFitTestResult = Statistics.chiSqTest(vec)
```

```
# summary of the test including the p-value, degrees of freedom,
```

```
# test statistic, the method used, and the null hypothesis.
```

```
print("%s\n" % goodnessOfFitTestResult)
```

```
mat = Matrices.dense(3, 2, [1.0, 3.0, 5.0, 2.0, 4.0, 6.0]) # a contingency matrix
```

```
# conduct Pearson's independence test on the input contingency matrix
```

```
independenceTestResult = Statistics.chiSqTest(mat)
```

```
# summary of the test including the p-value, degrees of freedom,
```

```
# test statistic, the method used, and the null hypothesis.
```

```
print("%s\n" % independenceTestResult)
```

```
obs = sc.parallelize(
```

```
    [LabeledPoint(1.0, [1.0, 0.0, 3.0]),
```

```
    LabeledPoint(1.0, [1.0, 2.0, 0.0]),
```

```
    LabeledPoint(1.0, [-1.0, 0.0, -0.5])]
```

```
) # LabeledPoint(label, feature)
```

```
# The contingency table is constructed from an RDD of LabeledPoint and used to conduct
```

```
# the independence test. Returns an array containing the ChiSquaredTestResult for every feature
```

```
# against the label.
```

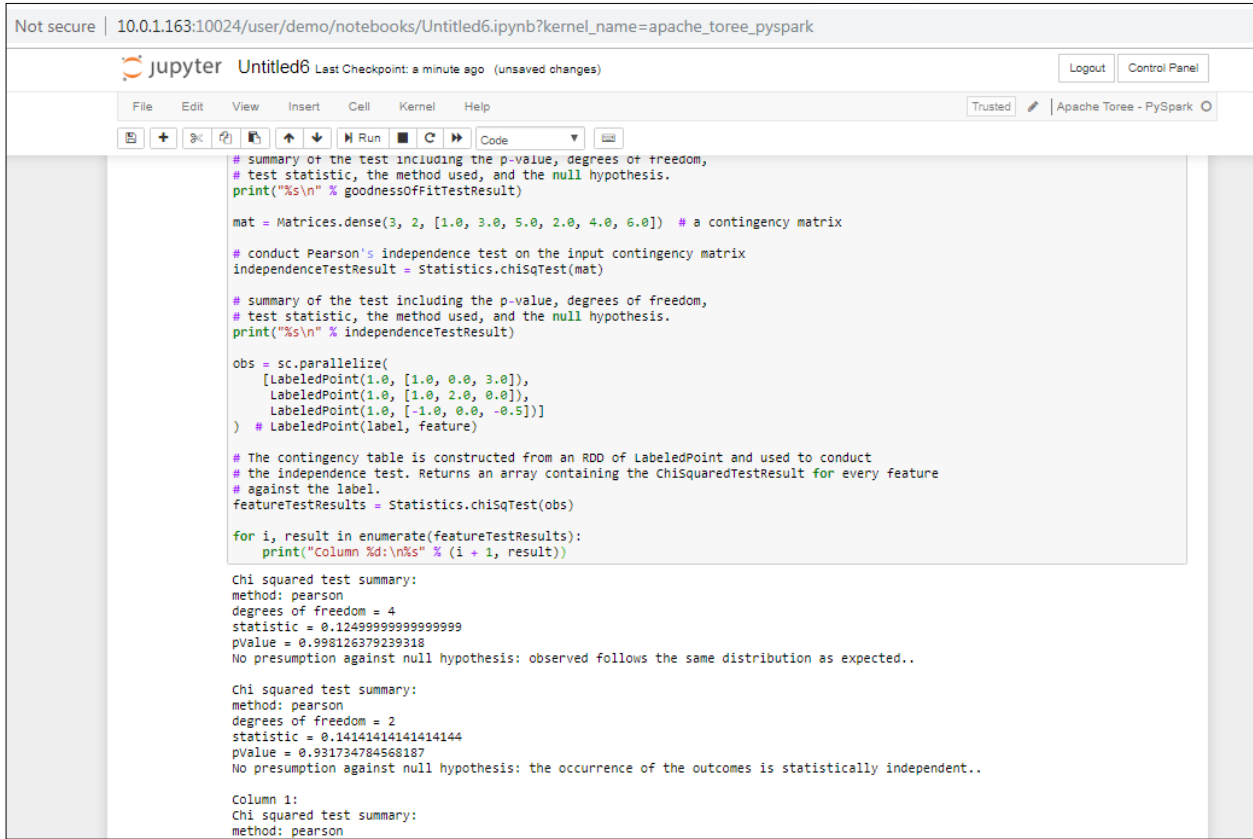
```
featureTestResults = Statistics.chiSqTest(obs)
```

```
for i, result in enumerate(featureTestResults):
```

```
    print("Column %d:\n%s" % (i + 1, result))
```

Input: No input files used. Data is generated in the code.

Output: Sample output is as given below.



```

# summary of the test including the p-value, degrees of freedom,
# test statistic, the method used, and the null hypothesis.
print("%s\n" % goodnessOfFitTestResult)

mat = Matrices.dense(3, 2, [1.0, 3.0, 5.0, 2.0, 4.0, 6.0]) # a contingency matrix

# conduct Pearson's independence test on the input contingency matrix
independenceTestResult = Statistics.chiSqTest(mat)

# summary of the test including the p-value, degrees of freedom,
# test statistic, the method used, and the null hypothesis.
print("%s\n" % independenceTestResult)

obs = sc.parallelize(
    [LabeledPoint(1.0, [1.0, 0.0, 3.0]),
     LabeledPoint(1.0, [1.0, 2.0, 0.0]),
     LabeledPoint(1.0, [-1.0, 0.0, -0.5])]
) # LabeledPoint(label, feature)

# The contingency table is constructed from an RDD of LabeledPoint and used to conduct
# the independence test. Returns an array containing the ChiSquaredTestResult for every feature
# against the label.
featureTestResults = Statistics.chiSqTest(obs)

for i, result in enumerate(featureTestResults):
    print("Column %d:\n%s" % (i + 1, result))

Chi squared test summary:
method: pearson
degrees of freedom = 4
statistic = 0.12499999999999999
pvalue = 0.998126379239318
No presumption against null hypothesis: observed follows the same distribution as expected..

Chi squared test summary:
method: pearson
degrees of freedom = 2
statistic = 0.14141414141414144
pvalue = 0.931734784568187
No presumption against null hypothesis: the occurrence of the outcomes is statistically independent..

Column 1:
Chi squared test summary:
method: pearson

```

4.3 Execute Spark Submit job on JupyterHub

Start a toree pySpark kernel -> Wait till kernel creates a spark shell. You can run up to 4 Spark shells with current configurations. If your shell doesn't start, you may have used up all the cores. Kill unused Kernels to release resources.

Code:

```
##sh
```

```
./bin/spark-submit \
--class org.apache.spark.examples.SparkPi \
--master local[8] \
```

```
/usr/lib/spark/spark-2.4.0-bin-hadoop2.7/examples/jars/spark-examples_2.11-2.4.0.jar \  
100
```

Output:

Check Spark master GUI that job is running under Running Applications section

5 SAMPLE TEST CASES FOR SPARK WITH NOTEBOOKS

5.1 PySpark testing

Start a toree pySpark kernel -> Wait till kernel creates a spark shell. You can run up to 4 Spark shells with current configurations. If your shell doesn't start, you may have used up all the cores. Kill unused Kernels to release resources.

Code:

```
from pyspark import SparkConf, SparkContext

from sklearn.datasets import make_classification

from sklearn.ensemble import ExtraTreesClassifier

import pandas as pd

import numpy as np

# Build a classification task using 3 informative features

X, y = make_classification(n_samples=12000,

                           n_features=10,

                           n_informative=3,

                           n_redundant=0,

                           n_repeated=0,

                           n_classes=2,

                           random_state=0,

                           shuffle=False)
```



```
# Partition data
```

```
def dataPart(X, y, start, stop): return dict(X=X[start:stop, :], y=y[start:stop])
```

```
def train(data):
```

```
    X = data['X']
```

```
    y = data['y']
```

```
    return ExtraTreesClassifier(n_estimators=100,random_state=0).fit(X,y)
```

```
# Merge 2 Models
```

```
from sklearn.base import copy
```

```
def merge(left,right):
```

```
    new = copy.deepcopy(left)
```

```
    new.estimators_ += right.estimators_
```

```
    new.n_estimators = len(new.estimators_)
```

```
    return new
```

```
data = [dataPart(X, y, 0, 4000), dataPart(X,y,4000,8000), dataPart(X,y,8000,12000)]
```

```
forest = sc.parallelize(data).map(train).reduce(merge)
```

```
importances = forest.feature_importances_
```

```
std = np.std([tree.feature_importances_ for tree in forest.estimators_],
```

```
             axis=0)
```

```
indices = np.argsort(importances)[::-1]
```

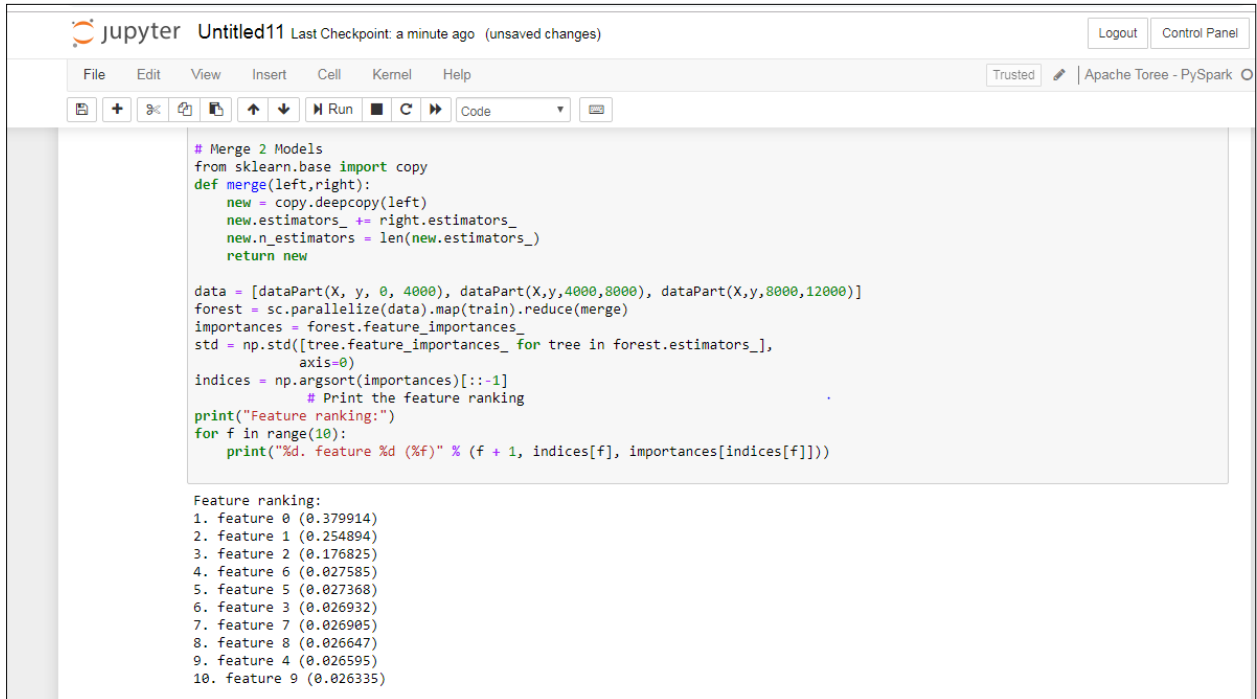
```
    # Print the feature ranking
```

```
print("Feature ranking:")
```

```
for f in range(10):
```

```
    print("%d. feature %d (%f)" % (f + 1, indices[f], importances[indices[f]]))
```

Output: Sample output is as given below.



```

# Merge 2 Models
from sklearn.base import copy
def merge(left,right):
    new = copy.deepcopy(left)
    new.estimators_ += right.estimators_
    new.n_estimators = len(new.estimators_)
    return new

data = [dataPart(X, y, 0, 4000), dataPart(X,y,4000,8000), dataPart(X,y,8000,12000)]
forest = sc.parallelize(data).map(train).reduce(merge)
importances = forest.feature_importances_
std = np.std([tree.feature_importances_ for tree in forest.estimators_],
              axis=0)
indices = np.argsort(importances)[::-1]
# Print the feature ranking
print("Feature ranking:")
for f in range(10):
    print("%d. feature %d (%f)" % (f + 1, indices[f], importances[indices[f]]))

Feature ranking:
1. feature 0 (0.379914)
2. feature 1 (0.254894)
3. feature 2 (0.176825)
4. feature 6 (0.027585)
5. feature 5 (0.027368)
6. feature 3 (0.026932)
7. feature 7 (0.026905)
8. feature 8 (0.026647)
9. feature 4 (0.026595)
10. feature 9 (0.026335)

```

5.2 Spark Scala testing

Start a toree scala kernel -> Wait till kernel creates a spark shell. Run following Pearson's correlation. You can run up to 4 Spark shells with current configurations. If your shell doesn't start, you may have used up all the cores. Kill unused Kernels to release resources

Code:

```
import org.apache.spark.mllib.linalg._
```

```
import org.apache.spark.mllib.stat.Statistics
```

```
import org.apache.spark.rdd.RDD

val seriesX: RDD[Double] = sc.parallelize(Array(1, 2, 3, 3, 5)) // a series

// must have the same number of partitions and cardinality as seriesX

val seriesY: RDD[Double] = sc.parallelize(Array(11, 22, 33, 33, 55))

// compute the correlation using Pearson's method. Enter "spearman" for Spearman's method. If a

// method is not specified, Pearson's method will be used by default.

val correlation: Double = Statistics.corr(seriesX, seriesY, "pearson")

println(s"Correlation is: $correlation")

    val data: RDD[Vector] = sc.parallelize(

Seq(

    Vectors.dense(1.0, 10.0, 100.0),

    Vectors.dense(2.0, 20.0, 200.0),

    Vectors.dense(5.0, 33.0, 366.0))

) // note that each Vector is a row and not a column

    // calculate the correlation matrix using Pearson's method. Use "spearman" for Spearman's

method

// If a method is not specified, Pearson's method will be used by default.

val correlMatrix: Matrix = Statistics.corr(data, "pearson")

println(correlMatrix.toString)

Spark R-studio test on jupyter notebook

Open R-studio GUI and execute the following scrip
```

```
library(data.table)
```

```
dt <- data.table(1:3)
```

```
print(dt)
```

```
for (i in 1:5) {
```

```
  print(i*2)
```

```
}
```

```
print(1:50)
```

Output: Sample output is as given below.

```
> library(data.table)
Error in library(data.table) : there is no package called 'data.table'
> dt <- data.table(1:3)
Error in data.table(1:3) : could not find function "data.table"
> print(dt)
function (x, df, ncp, log = FALSE)
{
  if (missing(ncp))
    .Call(C_dt, x, df, log)
  else .Call(C_dnt, x, df, ncp, log)
}
<bytecode: 0x5dad150>
<environment: namespace:stats>
> for (i in 1:5) {
+   print(i*2)
+ }
[1] 2
[1] 4
[1] 6
[1] 8
[1] 10
> print(1:50)
[1]  1  2  3  4  5  6  7  8  9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28
[29] 29 30 31 32 33 34 35 36 37 38 39 40 41 42 43 44 45 46 47 48 49 50
>
> |
```

6 TEST CASES FOR SPARK WITH RSTUDIO

- Create a Linux user on the master controller node or login as AD user.
- Login to R-studio

6.1 Base-R testing on RStudio GUI

```
data(iris) # Load the dataset iris
```

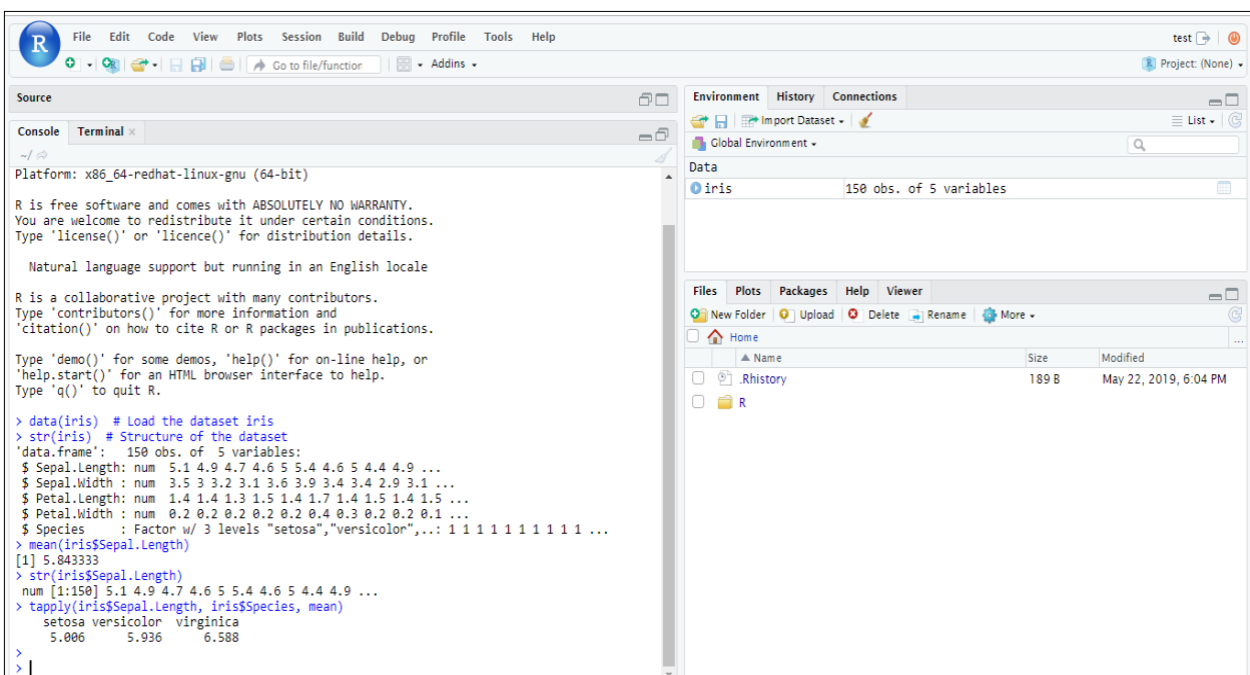
```
str(iris) # Structure of the dataset
```

```
mean(iris$Sepal.Length)
```

```
str(iris$Sepal.Length)
```

```
tapply(iris$Sepal.Length, iris$Species, mean)
```

Output: Sample output is as given below.



The screenshot shows the RStudio interface with the following components:

- Source Editor:** Contains the R code being executed.
- Console:** Displays the output of the R code, including the R startup message and the results of the commands.
- Environment:** Shows the global environment with the 'iris' dataset loaded, containing 150 observations of 5 variables.
- Files:** Shows the file explorer with the current project directory.

```
> data(iris) # Load the dataset iris
> str(iris) # Structure of the dataset
'data.frame': 150 obs. of 5 variables:
 $ Sepal.Length: num 5.1 4.9 4.7 4.6 5 5.4 4.6 5 4.4 4.9 ...
 $ Sepal.Width : num 3.5 3 3.2 3.1 3.6 3.9 3.4 3.4 2.9 3.1 ...
 $ Petal.Length: num 1.4 1.4 1.3 1.5 1.4 1.7 1.4 1.5 1.4 1.5 ...
 $ Petal.Width : num 0.2 0.2 0.2 0.2 0.2 0.4 0.3 0.2 0.2 0.1 ...
 $ Species : Factor w/ 3 levels "setosa","versicolor",...: 1 1 1 1 1 1 1 1 1 1 ...
> mean(iris$Sepal.Length)
[1] 5.843333
> str(iris$Sepal.Length)
num [1:150] 5.1 4.9 4.7 4.6 5 5.4 4.6 5 4.4 4.9 ...
> tapply(iris$Sepal.Length, iris$Species, mean)
      setosa versicolor  virginica 
      5.006      5.936      6.588
```

6.2 Sparklyr testing on RStudio GUI

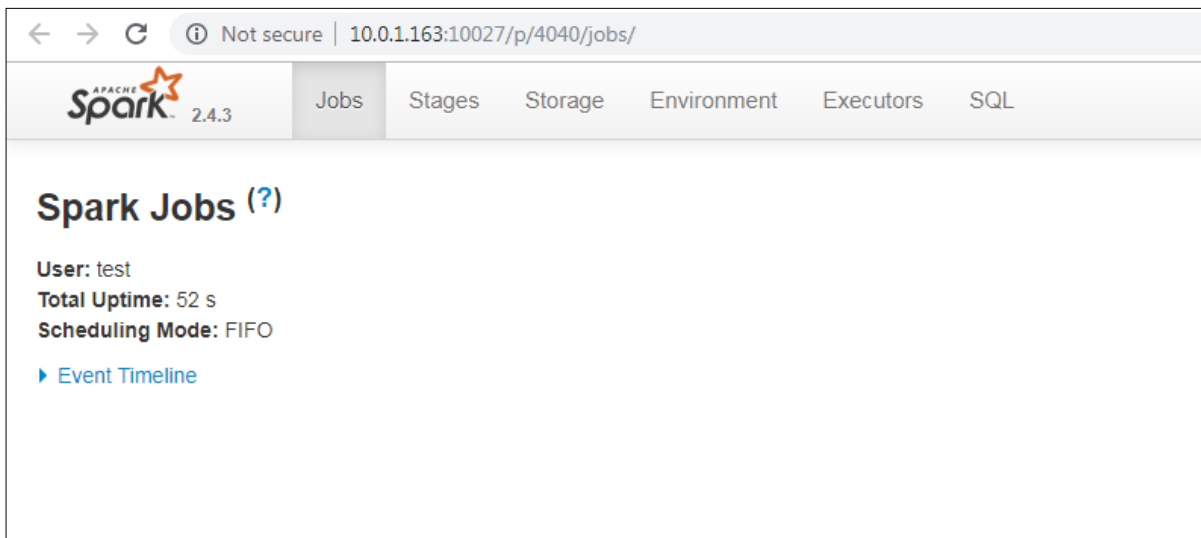
```
>install.packages("sparklyr")
```

```
>sparklyr::spark_install()
```

```
>library(sparklyr)
```

```
>sc <- spark_connect(master = 'local')
```

Output: Sample output is as given below.



6.3 Simple test on RStudio GUI

```
data(iris) # Load the dataset iris
```

```
str(iris) # Structure of the dataset
```

```
mean(iris$Sepal.Length)
```

```
str(iris$Sepal.Length)
```

```
tapply(iris$Sepal.Length, iris$Species, mean)
```

Output: Sample output is as given below

```
> data(iris) # Load the dataset iris
> str(iris) # Structure of the dataset
'data.frame': 150 obs. of 5 variables:
 $ Sepal.Length: num 5.1 4.9 4.7 4.6 5 5.4 4.6 5 4.4 4.9 ...
 $ Sepal.Width : num 3.5 3 3.2 3.1 3.6 3.9 3.4 3.4 2.9 3.1 ...
 $ Petal.Length: num 1.4 1.4 1.3 1.5 1.4 1.7 1.4 1.5 1.4 1.5 ...
 $ Petal.Width : num 0.2 0.2 0.2 0.2 0.2 0.4 0.3 0.2 0.2 0.1 ...
 $ Species : Factor w/ 3 levels "setosa","versicolor",...: 1 1 1 1 1 1 1 1 1 1 ...
> mean(iris$Sepal.Length)
[1] 5.843333
> str(iris$Sepal.Length)
num [1:150] 5.1 4.9 4.7 4.6 5 5.4 4.6 5 4.4 4.9 ...
> tapply(iris$Sepal.Length, iris$Species, mean)
      setosa versicolor virginica 
      5.006      5.936      6.588 
>
> |
```

6.4 MLLib usage test on RStudio GUI

```
>install.packages("sparklyr")
```

```
>sparklyr::spark_install()
```

```
>library(sparklyr)
```

```
>sc <- spark_connect(master = 'local')

> library(dplyr)

# copy mtcars into spark
> mtcars_tbl <- copy_to(sc, mtcars)

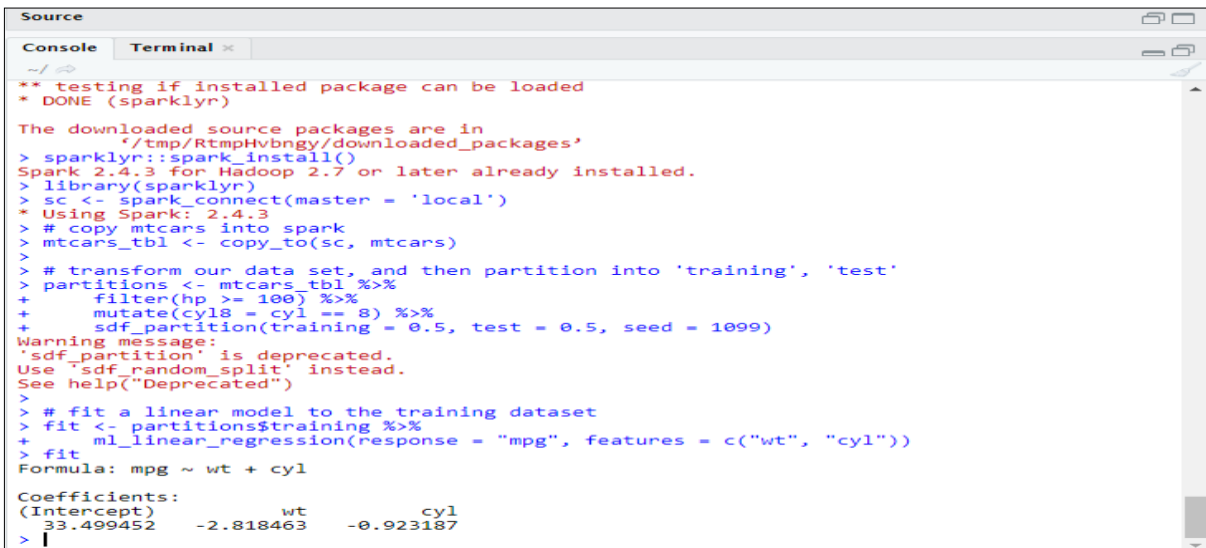
# ** May show an error regarding problem with database. Seems to work OK after that
>src_tbls(sc)

# transform our data set, and then partition into 'training', 'test'
> partitions <- mtcars_tbl %>%
  filter(hp >= 100) %>%
  mutate(cyl8 = cyl == 8) %>%
  sdf_partition(training = 0.5, test = 0.5, seed = 1099)

# fit a linear model to the training dataset
> fit <- partitions$training %>%
  ml_linear_regression(response = "mpg", features = c("wt", "cyl"))

> summary(fit)
```

Output: Sample output is as given below



```
Source
Console Terminal x
~/
** testing if installed package can be loaded
* DONE (sparklyr)

The downloaded source packages are in
  '/tmp/RtmpHvbngy/downloaded_packages'
> sparklyr::spark_install()
Spark 2.4.3 for Hadoop 2.7 or later already installed.
> library(sparklyr)
> sc <- spark_connect(master = 'local')
* Using Spark: 2.4.3
> # copy mtcars into spark
> mtcars_tbl <- copy_to(sc, mtcars)
>
> # transform our data set, and then partition into 'training', 'test'
> partitions <- mtcars_tbl %>%
+   filter(hp >= 100) %>%
+   mutate(cyl8 = cyl == 8) %>%
+   sdf_partition(training = 0.5, test = 0.5, seed = 1099)
Warning message:
'sdf_partition' is deprecated.
Use 'sdf_random_split' instead.
See help("Deprecated")
>
> # fit a linear model to the training dataset
> fit <- partitions$training %>%
+   ml_linear_regression(response = "mpg", features = c("wt", "cyl"))
> fit
Formula: mpg ~ wt + cyl

Coefficients:
(Intercept)      wt      cyl
 33.499452   -2.818463   -0.923187
> |
```


7 TEST CASES FOR SPARK WITH SQL

Spark SQL allows relational queries expressed in SQL or Scala to be executed using Spark. At the core of this component is a new type of RDD, SchemaRDD. SchemaRDDs are composed of Row objects, along with a schema that describes the data types of each column in the row. A SchemaRDD is similar to a table in a traditional relational database. The SchemaRDD can be created from an existing RDD, a Parquet file, a JSON dataset.

Follow these tests for testing spark-sql for your cluster:

7.1 Testing with user defined functions

- Creating a dataset “hello world”

```
val dataset = Seq((0, "hello"),(1, "world")).toDF("id","text")
```

- Defining a function ‘upper’ which converts a string into upper case

```
val upper: String => String =_.toUpperCase
```

- We now import the ‘udf’ package into Spark

```
import org.apache.spark.sql.functions.udf
```

- Defining our UDF, ‘upperUDF’ and importing our function ‘upper’

```
val upperUDF = udf(upper)
```

- Displaying the results of our User Defined Function in a new column ‘upper’

```
dataset.withColumn("upper", upperUDF('text')).show
```

7.2 Starting a Spark Session and displaying DataFrame of people.json

For the querying examples, we will be using files, 'people.txt' and 'people.json'. These files are stored at '/usr/lib/spark/spark-2.4.0-bin-hadoop2.7/examples/src/main/resources/'

- We first import a Spark Session into Apache Spark

```
import org.apache.spark.sql.SparkSession
```

- Creating a Spark Session 'spark' using the 'builder()' function

```
val spark = SparkSession.builder().appName("Spark SQL basic  
example").config("spark.some.config.option", "some-value").getOrCreate()
```

- Importing the Implicits class into our 'spark' Session.

```
import spark.implicits._
```

- We now create a DataFrame 'df' and import data from the 'employee.json' file.

```
val df = spark.read.json("examples/src/main/resources/people.json ")
```

- Displaying the DataFrame 'df'. The result is a table of 5 rows of ages and names from our 'employee.json' file.

```
df.show()
```

7.3 Creating a Dataset

- Creating a class 'Employee' to store name and age of an employee

```
case class Employee(name: String, age: Long)
```

- Assigning a Dataset 'caseClassDS' to store the record of Andrew

```
val caseClassDS = Seq(Employee("Andrew", 55)).toDS()
```

- Displaying the Dataset 'caseClassDS'

```
caseClassDS.show()
```

- Creating a primitive Dataset to demonstrate mapping of DataFrames into Datasets

```
val primitiveDS = Seq(1, 2, 3).toDS
```

- Assigning the above sequence into an array

```
primitiveDS.map(_ + 1).collect()
```