

Mini 7

part 1: load data into HDFS

log onto server

ssh aisrexit@montana.dataapplab.com -p 49233

mkdir mushroom # create a file mushroom in server

cd mushroom

copy local csv file to file mushroom in server (from a new terminal window)

scp -P 49233 ./mushrooms.csv aisrexit@montana

dataapplab.com:/home/aisrexit/mushroom

upload the file into an existed file named 'data' in HDFS

hdfs dfs -put mushrooms.csv hdfs:///user/aisrexit/data

double check if data is successfully uploaded to 'data'

hdfs dfs -ls hdfs:///user/aisrexit/data

#####

part 2: Pyspark

SPARK_MAJOR_VERSION=2 pyspark

from pyspark.sql import SQLContext

from pyspark.sql.types import *

from pyspark.ml import Pipeline

from pyspark.ml.classification import DecisionTreeClassifier

from pyspark.ml.feature import StringIndexer, VectorIndexer, VectorAssembler

from pyspark.ml.evaluation import MulticlassClassificationEvaluator

Create a structure

struct = StructType([

 StructField('class',StringType(),True),

 StructField('cap-shape',StringType(),True),

 StructField('cap-surface',StringType(),True),

 StructField('cap-color',StringType(),True),

 StructField('bruises',StringType(),True),

 StructField('odor',StringType(),True),

 StructField('gill-attachment',StringType(),True),

 StructField('gill-spacing',StringType(),True),

 StructField('gill-size',StringType(),True),

 StructField('gill-color',StringType(),True),

 StructField('stalk-shape',StringType(),True),

 StructField('stalk-root',StringType(),True),

 StructField('stalk-surface-above-ring',StringType(),True),

 StructField('stalk-surface-below-ring',StringType(),True),

 StructField('stalk-color-above-ring',StringType(),True),

 StructField('stalk-color-below-ring',StringType(),True),

 StructField('veil-type',StringType(),True),

```
StructField('veil-color',StringType(),True),  
StructField('ring-number',StringType(),True),  
StructField('ring-type',StringType(),True),  
StructField('spore-print-color',StringType(),True),  
StructField('population',StringType(),True),  
StructField('habitat',StringType(),True),  
])
```

```
# find the path for mushroom.csv  
hdfs dfs -ls /user/aisrexit/data/
```

```
# load data as DF  
df_mush =  
spark.read.csv('hdfs:///user/aisrexit/data/mushrooms.csv',header=True,schema=s  
truct)
```

```
df_mush
```



The screenshot shows a terminal window with a title bar that reads "meierhabarexiti — aisrexit@m1:~ — ssh aisrexit@montana.dataapplab.com -p...". The terminal content includes a Java stack trace for a warning message: "17/11/03 03:35:33 WARN DataSource: Error while looking for metadata directory." Below this, a user enters the command "[>>> df)mush", which results in a "SyntaxError: invalid syntax" message. The user then enters "[>>> df_mush", which successfully loads a DataFrame with 11 columns: class, cap-shape, cap-surface, cap-color, bruises, odor, gill-attachment, gill-spacing, gill-size, gill-color, stalk-shape, stalk-surface-above-ring, stalk-surface-below-ring, stalk-color-above-ring, stalk-color-below-ring, veil-type, veil-color, ring-number, ring-type, spore-print-color, population, and habitat. The DataFrame is displayed as a single row of values for these columns.

```
#### Indexing  
## features  
# index features : good  
indexers = [StringIndexer(inputCol=column,  
outputCol=column+"_index").fit(df_mush) for column in list(set(df_mush.columns))]  
  
pipeline = Pipeline(stages=indexers)
```

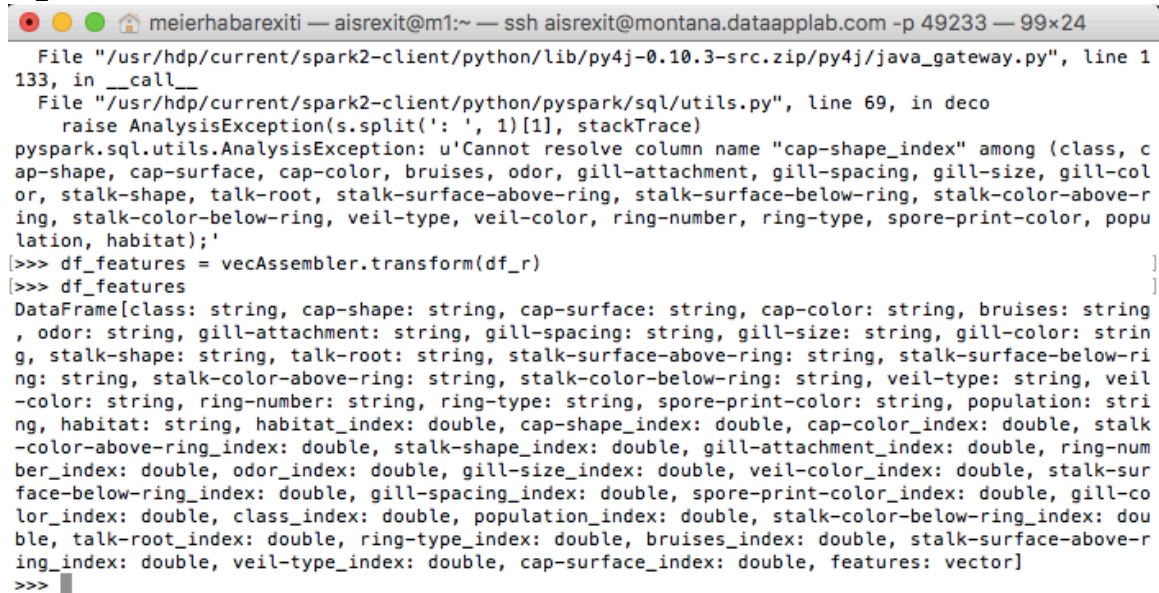
```
df_r = pipeline.fit(df_mush).transform(df_mush)
```

```
df_r.show() # the new dataframe with coloumn and columm+index
```

```
# Assemble all features: all features in VectorAssembler has to be numeric
vecAssembler = VectorAssembler(inputCols=["cap-shape_index","cap-
surface_index","cap-color_index","bruises_index","odor_index","gill-
attachment_index","gill-spacing_index","gill-size_index","gill-color_index","stalk-
shape_index","talk-root_index","stalk-surface-above-ring_index","stalk-surface-
below-ring_index","stalk-color-above-ring_index","stalk-color-below-
ring_index","veil-type_index","veil-color_index","ring-number_index","ring-
type_index","spore-print-
color_index","population_index","habitat_index"],outputCol="features")
vecAssembler
```

```
df_features = vecAssembler.transform(df_r)
```

```
df_features:
```



The screenshot shows a terminal window with a title bar indicating a remote SSH session. The terminal displays an error message from Spark SQL: 'Cannot resolve column name "cap-shape_index" among (class, cap-shape, cap-surface, cap-color, bruises, odor, gill-attachment, gill-spacing, gill-size, gill-color, stalk-shape, talk-root, stalk-surface-above-ring, stalk-surface-below-ring, stalk-color-above-ring, stalk-color-below-ring, veil-type, veil-color, ring-number, ring-type, spore-print-color, population, habitat);'. Below the error, the user enters a command to transform the DataFrame, and the terminal shows the resulting DataFrame schema, which includes columns for mushroom features and their corresponding indices, along with a 'features' vector column.

```
meierhabarexiti — aisrexit@m1:~ — ssh aisrexit@montana.dataapplab.com -p 49233 — 99x24
File "/usr/hdp/current/spark2-client/python/lib/py4j-0.10.3-src.zip/py4j/java_gateway.py", line 1
133, in __call__
File "/usr/hdp/current/spark2-client/python/pyspark/sql/utils.py", line 69, in deco
raise AnalysisException(s.split(':', 1)[1], stackTrace)
pyspark.sql.utils.AnalysisException: u'Cannot resolve column name "cap-shape_index" among (class, c
ap-shape, cap-surface, cap-color, bruises, odor, gill-attachment, gill-spacing, gill-size, gill-col
or, stalk-shape, talk-root, stalk-surface-above-ring, stalk-surface-below-ring, stalk-color-above-r
ing, stalk-color-below-ring, veil-type, veil-color, ring-number, ring-type, spore-print-color, popu
lation, habitat);'
[>>> df_features = vecAssembler.transform(df_r)
[>>> df_features
DataFrame[class: string, cap-shape: string, cap-surface: string, cap-color: string, bruises: string
, odor: string, gill-attachment: string, gill-spacing: string, gill-size: string, gill-color: strin
g, stalk-shape: string, talk-root: string, stalk-surface-above-ring: string, stalk-surface-below-ri
ng: string, stalk-color-above-ring: string, stalk-color-below-ring: string, veil-type: string, veil
-color: string, ring-number: string, ring-type: string, spore-print-color: string, population: stri
ng, habitat: string, habitat_index: double, cap-shape_index: double, cap-color_index: double, stalk
-color-above-ring_index: double, stalk-shape_index: double, gill-attachment_index: double, ring-num
ber_index: double, odor_index: double, gill-size_index: double, veil-color_index: double, stalk-sur
face-below-ring_index: double, gill-spacing_index: double, spore-print-color_index: double, gill-co
lor_index: double, class_index: double, population_index: double, stalk-color-below-ring_index: dou
ble, talk-root_index: double, ring-type_index: double, bruises_index: double, stalk-surface-above-r
ing_index: double, veil-type_index: double, cap-surface_index: double, features: vector]
>>>
```

```
df_features.head(1)
```

```

| e|      b|      s|      w|      t|      l|      f|      c|      b|      n|      e|      c|      n|
s|      s|      w|      p|      w|      o|      p|      n|      n|
m|      5.0|      3.0|      4.0|      0.0|      1.0|      0.0|      0.0|
5.0|      0.0|      0.0|      0.0|      0.0|      1.0|      3.0|      0.0|
3.0|      0.0|      3.0|      0.0|      1.0|      0.0|      0.0|
1.0|(22,[0,1,2,3,4,...])
+-----+-----+-----+-----+-----+-----+-----+-----+-----+-----+
+-----+-----+-----+-----+-----+-----+-----+-----+-----+-----+
+-----+-----+-----+-----+-----+-----+-----+-----+-----+-----+
+-----+-----+-----+-----+-----+-----+-----+-----+-----+-----+
+-----+-----+-----+-----+-----+-----+-----+-----+-----+-----+
only showing top 3 rows

>>> df_features.head(1)
[Row(class=u'p', cap-shape=u'x', cap-surface=u's', cap-color=u'n', bruises=u't', odor=u'p', gill-attachment=u'f', gill-spacing=u'c', gill-size=u'n', gill-color=u'k', stalk-shape=u'e', talk-root=u'e', stalk-surface-above-ring=u's', stalk-surface-below-ring=u's', stalk-color-above-ring=u'w', stalk-color-below-ring=u'w', veil-type=u'p', veil-color=u'w', ring-number=u'o', ring-type=u'p', spore-print-color=u'k', population=u's', habitat=u'u', habitat_index=4.0, cap-shape_index=0.0, cap-color_index=0.0, stalk-color-above-ring_index=0.0, stalk-shape_index=1.0, gill-attachment_index=0.0, ring-number_index=0.0, odor_index=6.0, gill-size_index=1.0, veil-color_index=0.0, stalk-surface-below-ring_index=0.0, gill-spacing_index=0.0, spore-print-color_index=2.0, gill-color_index=7.0, class_index=1.0, population_index=2.0, stalk-color-below-ring_index=0.0, talk-root_index=2.0, ring-type_index=0.0, bruises_index=1.0, stalk-surface-above-ring_index=0.0, veil-type_index=0.0, cap-surface_index=1.0, features=SparseVector(22, {1: 1.0, 3: 1.0, 4: 6.0, 7: 1.0, 8: 7.0, 9: 1.0, 10: 2.0, 19: 2.0, 20: 2.0, 21: 4.0})))]
>>>

```

Fitting model

train test split

```
(trainingData, testData) = df_features.randomSplit([0.7, 0.3])
```

fitting a decision tree model

```
dt = DecisionTreeClassifier(labelCol="class_index", featuresCol="features")
```

```
dt_model = dt.fit(trainingData)
```

```
meierhabarexiti — aisrexiti@ml1:~ — ssh aisrexiti@montana.dataaplab.com -p 49233 — 105x24
```

```
+-----+
+-----+
+-----+
only showing top 3 rows

>>> df_features.head(1)
[Row(class=u'p', cap-shape=u'x', cap-surface=u's', cap-color=u'n', bruises=u't', odor=u'p', gill-attachme
nt=u'f', gill-spacing=u'c', gill-size=u'n', gill-color=u'k', stalk-shape=u'e', talk-root=u'e', stalk-surf
ace-above-ring=u's', stalk-surface-below-ring=u's', stalk-color-above-ring=u'w', stalk-color-below-ring=u
'w', veil-type=u'p', veil-color=u'w', ring-number=u'o', ring-type=u'p', spore-print-color=u'k', populatio
n=u's', habitat=u'u', habitat_index=4.0, cap-shape_index=0.0, cap-color_index=0.0, stalk-color-above-ring
_index=0.0, stalk-shape_index=1.0, gill-attachment_index=0.0, ring-number_index=0.0, odor_index=6.0, gill
-size_index=1.0, veil-color_index=0.0, stalk-surface-below-ring_index=0.0, gill-spacing_index=0.0, spore-
print-color_index=2.0, gill-color_index=7.0, class_index=1.0, population_index=2.0, stalk-color-below-rin
g_index=0.0, talk-root_index=2.0, ring-type_index=0.0, bruises_index=1.0, stalk-surface-above-ring_index=
0.0, veil-type_index=0.0, cap-surface_index=1.0, features=SparseVector(22, {1: 1.0, 3: 1.0, 4: 6.0, 7: 1.
0, 8: 7.0, 9: 1.0, 10: 2.0, 19: 2.0, 20: 2.0, 21: 4.0})]]

>>> (trainingData, testData) = df_features.randomSplit([0.7, 0.3])
>>> dt = DecisionTreeClassifier(labelCol="class_index", featuresCol="features")
>>> dt_model = dt.fit(trainingData)
>>> dt_model
DecisionTreeClassificationModel (uid=DecisionTreeClassifier_496a858acea9fa1e8f0d) of depth 5 with 11 node
s
>>>
```

make predictions on testdata

```
df_predicted = dt_model.transform(testData.select('features','class_index'))
```

```
df_predicted.show(2)
```

```
meierhabarexiti — aisrexit@m1:~ — ssh aisrexit@montana.dataaplab.com -p 49233 — 105x24
```

```
n=u's', habitat=u'u', habitat_index=4.0, cap-shape_index=0.0, cap-color_index=0.0, stalk-color-above-ring_index=0.0, stalk-shape_index=1.0, gill-attachment_index=0.0, ring-number_index=0.0, odor_index=6.0, gill-size_index=1.0, veil-color_index=0.0, stalk-surface-below-ring_index=0.0, gill-spacing_index=0.0, spore-print-color_index=2.0, gill-color_index=7.0, class_index=1.0, population_index=2.0, stalk-color-below-ring_index=0.0, stalk-root_index=2.0, ring-type_index=0.0, bruises_index=1.0, stalk-surface-above-ring_index=0.0, veil-type_index=0.0, cap-surface_index=1.0, features=SparseVector(22, {1: 1.0, 3: 1.0, 4: 6.0, 7: 1.0, 8: 7.0, 9: 1.0, 10: 2.0, 19: 2.0, 20: 2.0, 21: 4.0}))]
```

```
>>> (trainingData, testData) = df_features.randomSplit([0.7, 0.3])
>>> dt = DecisionTreeClassifier(labelCol="class_index", featuresCol="features")
>>> dt_model = dt.fit(trainingData)
>>> dt_model
DecisionTreeClassificationModel (uid=DecisionTreeClassifier_496a858acea9fa1e8f0d) of depth 5 with 11 nodes
>>> df_predicted = dt_model.transform(testData.select('features','class_index'))
>>> df_predicted.show(2)
```

features	class_index	rawPrediction	probability	prediction
(22,[0,1,2,6,8,9,...]	0.0	[2888.0,6.0]	[0.99792674498963...	0.0
(22,[0,1,2,6,8,9,...]	0.0	[2888.0,6.0]	[0.99792674498963...	0.0

```
only showing top 2 rows
>>>
```

```
df_predicted.select("features","class_index","prediction").show(5)
```

```
meierhabarexiti — aisrexit@m1:~ — ssh aisrexit@montana.dataaplab.com -p 49233 — 105x24
```

```
DataFrame[features: vector, class_index: double, rawPrediction: vector, probability: vector, prediction: double]
>>> df_predicted.show(2)
```

features	class_index	rawPrediction	probability	prediction
(22,[0,1,2,6,8,9,...]	0.0	[2888.0,6.0]	[0.99792674498963...	0.0
(22,[0,1,2,6,8,9,...]	0.0	[2888.0,6.0]	[0.99792674498963...	0.0

```
only showing top 2 rows

>>> df_predicted.select("features","class_index","prediction").show(5)
```

features	class_index	prediction
(22,[0,1,2,6,8,9,...]	0.0	0.0
(22,[0,1,2,6,8,9,...]	0.0	0.0
(22,[0,1,2,6,8,9,...]	0.0	0.0
(22,[0,1,2,6,8,9,...]	0.0	0.0
(22,[0,1,2,6,8,9,...]	0.0	0.0

```
only showing top 5 rows
>>>
```

Evaluation

Select (prediction, true label) and compute test error

```
evaluator = MulticlassClassificationEvaluator(labelCol="class_index",
predictionCol="prediction")
```



```
evaluator.evaluate(df_predicted)
```

```
accuracy = evaluator.evaluate(df_predicted)  
print("Test Error = %g " % (1.0 - accuracy))
```

```
meierhabarexiti — aisrexit@m1:~ — ssh aisrexit@montana.dataapplab.com -p 49233 — 105x24  
[>>> evaluator  
MulticlassClassificationEvaluator_4bb59c11ffc3e3144804  
[>>> accuracy = evaluator.evaluate(df_predicted)  
Traceback (most recent call last):  
  File "<stdin>", line 1, in <module>  
  File "/usr/hdp/current/spark2-client/python/pyspark/ml/evaluation.py", line 68, in evaluate  
    return self._evaluate(dataset)  
  File "/usr/hdp/current/spark2-client/python/pyspark/ml/evaluation.py", line 98, in _evaluate  
    return self._java_obj.evaluate(dataset._jdf)  
  File "/usr/hdp/current/spark2-client/python/lib/py4j-0.10.3-src.zip/py4j/java_gateway.py", line 1133, i  
n __call__  
  File "/usr/hdp/current/spark2-client/python/pyspark/sql/utils.py", line 79, in deco  
    raise IllegalArgumentException(s.split(': ', 1)[1], stackTrace)  
pyspark.sql.utils.IllegalArgumentException: u'Field "label" does not exist.'  
[>>> evaluator = MulticlassClassificationEvaluator(labelCol="class_index", predictionCol="prediction")  
[>>> evaluator  
MulticlassClassificationEvaluator_4ac082c1ae54d61a0569  
[>>> evaluator.evaluate(df_predicted)  
0.997561712613218  
>>> accuracy = evaluator.evaluate(df_predicted)  
  
>>> print("Test Error = %g " % (1.0 - accuracy))  
Test Error = 0.00243829  
>>> ]
```