In [2]: %reload\_ext autoreload
%autoreload 2

# MIDS - w261 Machine Learning At Scale

Course Lead: Dr James G. Shanahan (email Jimi via James.Shanahan AT gmail.com)

## **Assignment - HW10**

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**Week:** 10

# **Table of Contents**

- 1. HW Introduction
- 2. HW References
- 3. HW Problems
  - 10.0. Short Answer Questions
  - 10.1. Word Count plus sorting
  - 10.2. MLlib-centric Kmeans
  - 10.3. Homegrown KMeans in Spark
  - 10.4. Making Homegrown KMeans more efficient
  - 10.5. OPTIONAL Weighted KMeans
  - 10.6. OPTIONAL Linear Regression
  - 10.7. OPTIONAL Error surfaces

## 1 Instructions

#### Back to Table of Contents

- Homework submissions are due by Tueday, 07/28/2016 at 11AM (West Coast Time).
- Prepare a single Jupyter note, please include questions, and question numbers in the questions and in the responses. Submit your homework notebook via the following form:
  - <u>Submission Link Google Form</u>
     (<a href="https://docs.google.com/forms/d/1ZOr9Rnle\_A06AcZDB6K1mJN4vrLeSmS2PD6Xm3eOiis/viewform\_usp=send\_form">https://docs.google.com/forms/d/1ZOr9Rnle\_A06AcZDB6K1mJN4vrLeSmS2PD6Xm3eOiis/viewform\_usp=send\_form</a>)

#### **Documents:**

- IPython Notebook, published and viewable online.
- PDF export of IPython Notebook.

## 2 Useful References

#### Back to Table of Contents

- Karau, Holden, Konwinski, Andy, Wendell, Patrick, & Zaharia, Matei. (2015). Learning Spark: Lightning-fast big data analysis. Sebastopol, CA: O'Reilly Publishers.
- Hastie, Trevor, Tibshirani, Robert, & Friedman, Jerome. (2009). The elements of statistical learning:
   Data mining, inference, and prediction (2nd ed.). Stanford, CA: Springer Science+Business Media.
   (Download for free <a href="here">here</a>
  - (http://statweb.stanford.edu/~tibs/ElemStatLearn/printings/ESLII print10.pdf))

## 3 HW Problems

**Back to Table of Contents** 

# **HW10.0: Short answer questions**

Back to Table of Contents

What is Apache Spark and how is it different to Apache Hadoop?

Spark is an optimized engine that supports general execution graphs over an RDD. Spark allows increased parallelism and higher-level data management. As a result, it is much faster tahn Hadoop, and has specialized libarries for machine learning, graph processing, and database management.

Fill in the blanks: Spark API consists of interfaces to develop applications based on it in Java, Scala, Python and R languages.

Using Spark, resource management can be done either in a single server instance or using a framework such as Mesos or YARN in a distributed manner.

What is an RDD and show a fun example of creating one and bringing the first element back to the driver program.

RDD (Resilient Distributed Datasets) is a fundamental data structure of Spark. It is an immutable distributed collection of objects. Each dataset in RDD is divided into logical partitions, which may be computed on different nodes of the cluster.

```
In [3]: #Fun Example:
    rdd = sc.parallelize('file_to_be_processed.txt') #distributes the str
    ing
    rdd.first()
Out[3]: 'f'
```

## **HW10.1 WordCount plus sorting**

Back to Table of Contents

The following notebooks will be useful to jumpstart this collection of Homework exercises:

- <u>Example Notebook with Debugging tactics in Spark</u>
   (<a href="http://nbviewer.jupyter.org/urls/dl.dropbox.com/s/jqjllp8kmf1eolk/WordCountDebugging-Example.ipynb">http://nbviewer.jupyter.org/urls/dl.dropbox.com/s/jqjllp8kmf1eolk/WordCountDebugging-Example.ipynb</a>)
- Word Count Quiz (http://nbviewer.jupyter.org/urls/dl.dropbox.com/s/vgmpivsi4rvqz0s/WordCountQuiz.ipynb)
- Work Count Solution
   (http://nbviewer.jupyter.org/urls/dl.dropbox.com/s/dxv3dmp1vluuo8i/WordCountQuiz-Solution.ipvnb)

In Spark write the code to count how often each word appears in a text document (or set of documents). Please use this homework document (with no solutions in it) as a the example document to run an experiment. Report the following:

• provide a sorted list of tokens in decreasing order of frequency of occurrence limited to [top 20 most frequent only] and [bottom 10 least frequent].

**OPTIONAL** Feel free to do a secondary sort where words with the same frequncy are sorted alphanumerically increasing. Plseas refer to the <u>following notebook</u> (<a href="http://nbviewer.jupyter.org/urls/dl.dropbox.com/s/uu5afr3ufpm9fy8/SecondarySort.ipynb">http://nbviewer.jupyter.org/urls/dl.dropbox.com/s/uu5afr3ufpm9fy8/SecondarySort.ipynb</a>) for examples of secondary sorts in Spark. Please provide the following [top 20 most frequent terms only] and [bottom 10 least frequent terms]

**NOTE** [Please incorporate all referenced notebooks directly into this master notebook as cells for HW submission. I.e., HW submissions should comprise of just one notebook]\_\_

# In [2]: ## Set up Spark

```
import os
import sys
import pyspark
from pyspark.sql import SQLContext

# We can give a name to our app (to find it in Spark WebUI) and config
ure execution mode
# In this case, it is local multicore execution with "local[*]"
app_name = "example-logs"
master = "local[*]"
conf = pyspark.SparkConf().setAppName(app_name).setMaster(master)
sc = pyspark.SparkContext(conf=conf)
sqlContext = SQLContext(sc)
print sc
print sqlContext
```

<pyspark.context.SparkContext object at 0x7f5ba0397e50>
<pyspark.sql.context.SQLContext object at 0x7f5b77217390>

```
In [86]: ## Drivers & Runners

# complete word count
#Count words in file/directory
logFileNAME = 'enronemail_lh.txt'
text_file = sc.textFile(logFileNAME)

counts = text_file.flatMap(lambda line: line.split(" ")) \
    .map(lambda word: (word, 1)) \
    .reduceByKey(lambda a, b: a + b, 1) \
    .sortBy(lambda x: -x[1])

wordCounts = counts.collect()

print ("20 most popular terms:")
print wordCounts[:20]

print ("10 least popular terms:")
print wordCounts[-10:]
```

```
20 most popular terms:
[(u'', 3504), (u'the', 1240), (u'to', 908), (u'and', 646), (u'of', 5
55), (u'a', 514), (u'in', 412), (u'your', 389), (u'you', 376), (u'fo
r', 368), (u'@', 361), (u'on', 253), (u'this', 243), (u'is', 243), (
u'""', 239), (u'will', 234), (u'i', 232), (u'|', 228), (u'be', 216),
(u'that', 213)]
10 least popular terms:
[(u'kinds', 1), (u'pinion/hou/ect', 1), (u'webpage,', 1), (u'inheren
tly', 1), (u'my,', 1), (u'my.', 1), (u'coffee,', 1), (u'"">>', 1), (u'commencement', 1), (u'volumes', 1)]
```

## HW10.1.1

#### **Back to Table of Contents**

Modify the above word count code to count words that begin with lower case letters (a-z) and report your findings. Again sort the output words in decreasing order of frequency.

```
20 most popular terms:
[(u'the', 1240), (u'to', 908), (u'and', 646), (u'of', 555), (u'a', 5
14), (u'in', 412), (u'your', 389), (u'you', 376), (u'for', 368), (u'
on', 253), (u'this', 243), (u'is', 243), (u'will', 234), (u'i', 232)
, (u'be', 216), (u'that', 213), (u'with', 197), (u'have', 168), (u'a
re', 168), (u'we', 161)]
10 least popular terms:
[(u'249.', 0), (u'11)', 0), (u'12/15', 0), (u'8,703', 0), (u'*', 0),
(u'877.', 0), (u'07:47', 0), (u'-->', 0), (u'9,908', 0), (u'"">>', 0
)]
```

## **HW10.2: MLlib-centric KMeans**

Back to Table of Contents

Using the following MLlib-centric KMeans code snippet:

```
from pyspark.mllib.clustering import KMeans, KMeansModel
from numpy import array
from math import sqrt
# Load and parse the data
# NOTE kmeans data.txt is available here
           https://www.dropbox.com/s/q85t0ytb9apggnh/kmeans_data.txt?dl=0
data = sc.textFile("kmeans data.txt")
parsedData = data.map(lambda line: array([float(x) for x in line.split(' '
)]))
# Build the model (cluster the data)
clusters = KMeans.train(parsedData, 2, maxIterations=10,
        runs=10, initializationMode="random")
# Evaluate clustering by computing Within Set Sum of Squared Errors
def error(point):
    center = clusters.centers[clusters.predict(point)]
    return sqrt(sum([x**2 for x in (point - center)]))
WSSSE = parsedData.map(lambda point: error(point)).reduce(lambda x, y: x +
y)
print("Within Set Sum of Squared Error = " + str(WSSSE))
# Save and load model
clusters.save(sc, "myModelPath")
sameModel = KMeansModel.load(sc, "myModelPath")
```

#### NOTE

The **kmeans\_data.txt** is available here <a href="https://www.dropbox.com/s/q85t0ytb9apggnh/kmeans\_data.txt?">https://www.dropbox.com/s/q85t0ytb9apggnh/kmeans\_data.txt?</a> dl=0 (https://www.dropbox.com/s/q85t0ytb9apggnh/kmeans\_data.txt?dl=0)

#### **TASKS**

- Run this code snippet and list the clusters that your find.
- compute the Within Set Sum of Squared Errors for the found clusters. Comment on your findings.

```
In [5]: %%writefile kmeans_data.txt
0.0 0.0 0.0
0.1 0.1 0.1
0.2 0.2 0.2
9.0 9.0 9.0
9.1 9.1 9.1
9.2 9.2 9.2
```

Overwriting kmeans data.txt

```
In [6]: ## Code goes here
        from pyspark.mllib.clustering import KMeans, KMeansModel
        from numpy import array
        from math import sqrt
        # Load and parse the data
        data = sc.textFile("kmeans data.txt")
        parsedData = data.map(lambda line: array([float(x) for x in line.split
        ('')]))
        # Build the model (cluster the data)
        clusters = KMeans.train(parsedData, 2, maxIterations=10, runs=10, init
        ializationMode="random")
        # Evaluate clustering by computing Within Set Sum of Squared Errors
        def error(point):
            center = clusters.centers[clusters.predict(point)]
            return sqrt(sum([x**2 for x in (point - center)]))
        WSSSE = parsedData.map(lambda point: error(point)).reduce(lambda x, y:
        print("Within Set Sum of Squared Error = " + str(WSSSE))
        # Save and load model
        clusters.save(sc, "myModelPath")
        sameModel = KMeansModel.load(sc, "myModelPath")
```

/usr/local/spark/python/pyspark/mllib/clustering.py:176: UserWarning : Support for runs is deprecated in 1.6.0. This param will have no e ffect in 1.7.0.

"Support for runs is deprecated in 1.6.0. This param will have no effect in 1.7.0.")

Within Set Sum of Squared Error = 0.692820323028

```
Py4JJavaErrorTraceback (most recent call last)
<ipython-input-6-c0273e617df9> in <module>()
```

21

```
22 # Save and load model
---> 23 clusters.save(sc, "myModelPath")
     24 sameModel = KMeansModel.load(sc, "myModelPath")
/usr/local/spark/python/pyspark/mllib/clustering.py in save(self, sc
, path)
    150
                java centers = py2java(sc, [ convert to vector(c) f
or c in self.centers])
                java model = sc. jvm.org.apache.spark.mllib.clusteri
ng.KMeansModel(java centers)
--> 152
                java_model.save(sc._jsc.sc(), path)
    153
    154
            @classmethod
/usr/local/spark/python/lib/py4j-0.9-src.zip/py4j/java_gateway.py in
call (self, *args)
                answer = self.gateway client.send command(command)
    811
    812
                return value = get return value(
--> 813
                    answer, self.gateway client, self.target id, sel
f.name)
    814
    815
                for temp arg in temp args:
/usr/local/spark/python/pyspark/sql/utils.py in deco(*a, **kw)
     43
            def deco(*a, **kw):
     44
                try:
---> 45
                    return f(*a, **kw)
                except py4j.protocol.Py4JJavaError as e:
     46
     47
                    s = e.java exception.toString()
/usr/local/spark/python/lib/py4j-0.9-src.zip/py4j/protocol.py in get
_return_value(answer, gateway_client, target_id, name)
    306
                        raise Py4JJavaError(
    307
                            "An error occurred while calling {0}{1}{
2}.\n".
--> 308
                            format(target id, ".", name), value)
    309
                    else:
    310
                        raise Py4JError(
```

Py4JJavaError: An error occurred while calling o87.save.

: org.apache.hadoop.mapred.FileAlreadyExistsException: Output direct ory file:/home/jovyan/work/root/Documents/MIDS/W 261 Machine Learnin g at Scale/myModelPath/metadata already exists

at org.apache.hadoop.mapred.FileOutputFormat.checkOutputSpecs(FileOutputFormat.java:132)

at org.apache.spark.rdd.PairRDDFunctions\$\$anonfun\$saveAsHado opDataset\$1.apply\$mcV\$sp(PairRDDFunctions.scala:1179)

at org.apache.spark.rdd.PairRDDFunctions\$\$anonfun\$saveAsHado opDataset\$1.apply(PairRDDFunctions.scala:1156)

at org.apache.spark.rdd.PairRDDFunctions\$\$anonfun\$saveAsHado

```
opDataset$1.apply(PairRDDFunctions.scala:1156)
        at org.apache.spark.rdd.RDDOperationScope$.withScope(RDDOper
ationScope.scala:150)
        at org.apache.spark.rdd.RDDOperationScope$.withScope(RDDOper
ationScope.scala:111)
        at org.apache.spark.rdd.RDD.withScope(RDD.scala:316)
        at org.apache.spark.rdd.PairRDDFunctions.saveAsHadoopDataset
(PairRDDFunctions.scala:1156)
        at org.apache.spark.rdd.PairRDDFunctions$$anonfun$saveAsHado
opFile$4.apply$mcV$sp(PairRDDFunctions.scala:1060)
        at org.apache.spark.rdd.PairRDDFunctions$$anonfun$saveAsHado
opFile$4.apply(PairRDDFunctions.scala:1026)
        at org.apache.spark.rdd.PairRDDFunctions$$anonfun$saveAsHado
opFile$4.apply(PairRDDFunctions.scala:1026)
        at org.apache.spark.rdd.RDDOperationScope$.withScope(RDDOper
ationScope.scala:150)
        at org.apache.spark.rdd.RDDOperationScope$.withScope(RDDOper
ationScope.scala:111)
        at org.apache.spark.rdd.RDD.withScope(RDD.scala:316)
        at org.apache.spark.rdd.PairRDDFunctions.saveAsHadoopFile(Pa
irRDDFunctions.scala:1026)
        at org.apache.spark.rdd.PairRDDFunctions$$anonfun$saveAsHado
opFile$1.apply$mcV$sp(PairRDDFunctions.scala:952)
        at org.apache.spark.rdd.PairRDDFunctions$$anonfun$saveAsHado
opFile$1.apply(PairRDDFunctions.scala:952)
        at org.apache.spark.rdd.PairRDDFunctions$$anonfun$saveAsHado
opFile$1.apply(PairRDDFunctions.scala:952)
        at org.apache.spark.rdd.RDDOperationScope$.withScope(RDDOper
ationScope.scala:150)
        at org.apache.spark.rdd.RDDOperationScope$.withScope(RDDOper
ationScope.scala:111)
        at org.apache.spark.rdd.RDD.withScope(RDD.scala:316)
        at org.apache.spark.rdd.PairRDDFunctions.saveAsHadoopFile(Pa
irRDDFunctions.scala:951)
        at org.apache.spark.rdd.RDD$$anonfun$saveAsTextFile$1.apply$
mcV$sp(RDD.scala:1457)
        at org.apache.spark.rdd.RDD$$anonfun$saveAsTextFile$1.apply(
RDD.scala:1436)
```

at org.apache.spark.rdd.RDD\$\$anonfun\$saveAsTextFile\$1.apply( RDD.scala:1436)

at org.apache.spark.rdd.RDDOperationScope\$.withScope(RDDOper ationScope.scala:150)

at org.apache.spark.rdd.RDDOperationScope\$.withScope(RDDOper ationScope.scala:111)

at org.apache.spark.rdd.RDD.withScope(RDD.scala:316)

at org.apache.spark.rdd.RDD.saveAsTextFile(RDD.scala:1436)

at org.apache.spark.mllib.clustering.KMeansModel\$SaveLoadV1 0\$.save(KMeansModel.scala:131)

at org.apache.spark.mllib.clustering.KMeansModel.save(KMeans Model.scala:96)

```
at sun.reflect.NativeMethodAccessorImpl.invoke0(Native Metho
d)
        at sun.reflect.NativeMethodAccessorImpl.invoke(NativeMethodA
ccessorImpl.java:57)
        at sun.reflect.DelegatingMethodAccessorImpl.invoke(Delegatin
gMethodAccessorImpl.java:43)
        at java.lang.reflect.Method.invoke(Method.java:606)
        at py4j.reflection.MethodInvoker.invoke(MethodInvoker.java:2
31)
        at py4j.reflection.ReflectionEngine.invoke(ReflectionEngine.
java:381)
        at py4j.Gateway.invoke(Gateway.java:259)
        at py4j.commands.AbstractCommand.invokeMethod(AbstractComman
d.java:133)
        at py4j.commands.CallCommand.execute(CallCommand.java:79)
        at py4j.GatewayConnection.run(GatewayConnection.java:209)
        at java.lang.Thread.run(Thread.java:745)
```

The clusters found are:

```
In [6]: for centroid in clusters.centers:
    print centroid

[ 0.1 0.1 0.1]
    [ 9.1 9.1 9.1]
```

And the Within Set Sum of Squared Errors for the found clusters is:

This means that by running the code above, we identified clusters centered at (0.1, 0.1, 0.1) and (9.1, 9.1, 9.1), with a Within Set Sum of Squared Errors = 0.6928.

## **HW10.3: Homegrown KMeans in Spark**

Back to Table of Contents

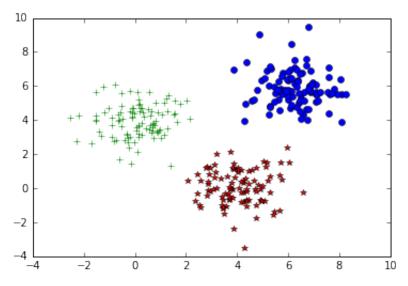
Download the following KMeans <u>notebook</u> (<a href="http://nbviewer.jupyter.org/urls/dl.dropbox.com/s/3nsthvp8g2rrrdh/EM-Kmeans.ipynb">http://nbviewer.jupyter.org/urls/dl.dropbox.com/s/3nsthvp8g2rrrdh/EM-Kmeans.ipynb</a>).

Generate 3 clusters with 100 (one hundred) data points per cluster (using the code provided). Plot the data. Then run MLlib's Kmean implementation on this data and report your results as follows:

- plot the resulting clusters after 1 iteration, 10 iterations, after 20 iterations, after 100 iterations.
- in each plot please report the Within Set Sum of Squared Errors for the found clusters (as part of the title WSSSE). Comment on the progress of this measure as the KMEans algorithms runs for more iterations. Then plot the WSSSE as a function of the iteration (1, 10, 20, 30, 40, 50, 100).

```
In [71]:
         %matplotlib inline
          import numpy as np
         import pylab
         import json
         size1 = size2 = size3 = 100
         samples1 = np.random.multivariate normal([4, 0], [[1, 0], [0, 1]], size
         1)
         data = samples1
         samples2 = np.random.multivariate normal([6, 6], [[1, 0],[0, 1]], size
         2)
         data = np.append(data,samples2, axis=0)
         samples 3 = \text{np.random.multivariate normal}([0, 4], [[1, 0], [0, 1]], \text{size}
         3)
         data = np.append(data,samples3, axis=0)
         # Randomlize data
         data = data[np.random.permutation(size1+size2+size3),]
         np.savetxt('data.csv',data,delimiter = ',')
```

```
In [72]: pylab.plot(samples1[:, 0], samples1[:, 1],'*', color = 'red')
    pylab.plot(samples2[:, 0], samples2[:, 1],'o',color = 'blue')
    pylab.plot(samples3[:, 0], samples3[:, 1],'+',color = 'green')
    pylab.show()
```



```
[ 0.05657819 3.97671718]
[ 6.26427572 5.86742717]
[ 4.02084087 0.08266008]
```

```
In [74]:
         import os
         import sys
         import numpy as np
         #Calculate which class each data point belongs to
         def nearest centroid(line):
             x = np.array([float(f) for f in line.split(',')])
             closest centroid idx = np.sum((x - centroids)**2, axis=1).argmin()
             return (closest centroid idx_{i}(x,1))
         #plot centroids and data points for each iteration
         def plot iteration(means):
             pylab.plot(samples1[:, 0], samples1[:, 1], '.', color = 'blue')
             pylab.plot(samples2[:, 0], samples2[:, 1], '.', color = 'blue')
             pylab.plot(samples3[:, 0], samples3[:, 1],'.', color = 'blue')
             pylab.plot(means[0][0], means[0][1],'*',markersize =10,color = 're
         d')
             pylab.plot(means[1][0], means[1][1],'*',markersize =10,color = 're
         d')
             pylab.plot(means[2][0], means[2][1],'*',markersize =10,color = 're
         d')
             pylab.show()
         def error(point):
             center = clusters.centers[clusters.predict(point)]
             return sqrt(sum([x**2 for x in (point - center)]))
```

```
In [75]: \#K = 3
         # Initialization: initialization of parameter is fixed to show an exam
         centroids = np.array([[0.0,0.0],[2.0,2.0],[0.0,7.0]])
         D = sc.textFile("./data.csv")
         parsedData = D.map(lambda line: array([float(x) for x in line.split(',
         ')]))
         wse = []
         iter_num = -1
         for i in range(101):
             res = D.map(nearest centroid).reduceByKey(lambda x,y : (x[0]+y[0],
         x[1]+y[1]).collect()
             res = sorted(res, key = lambda x : x[0]) #sort based on clusted ID
             centroids_new = np.array([x[1][0]/x[1][1]  for x in res])
                                                                         #divide
         by cluster size
             if np.sum(np.absolute(centroids new-centroids))<0.000001:</pre>
                 break
             iter num +=1
             centroids = centroids new
             if i in [1, 10, 20, 100]:
                 print "Iteration" + str(iter num)
                 print centroids
                 plot iteration(centroids)
                 WSSSE = parsedData.map(lambda point: error(point)).reduce(lamb
         da x, y: x + y
                 print("Within Set Sum of Squared Error = " + str(WSSSE))
         print "Final Results:"
         print centroids
```

```
Iteration1
[[ 1.70557734  0.8494642 ]
 [ 5.54041084 3.19625608]
 [ 1.2730003
               4.91916871]]
 8
 6
  4
  2
 0
-2
                                            10
Within Set Sum of Squared Error = 358.527731123
Final Results:
[[ 4.02084087  0.08266008]
 [ 6.26427572 5.86742717]
 [ 0.05657819  3.97671718]]
```

This process converges in less than 10 iterations, so we can't compare its WSE after multiple 20, 50 or 100 iterations.

# **HW10.4: KMeans Experiments**

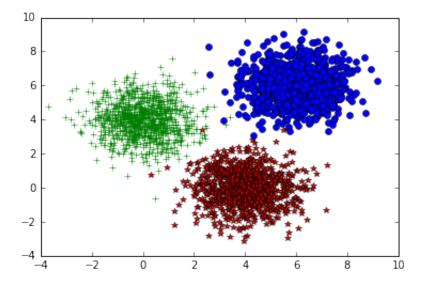
**Back to Table of Contents** 

Using this provided <u>homegrown Kmeans code</u> (<a href="http://nbviewer.jupyter.org/urls/dl.dropbox.com/s/3nsthvp8g2rrrdh/EM-Kmeans.ipynb">http://nbviewer.jupyter.org/urls/dl.dropbox.com/s/3nsthvp8g2rrrdh/EM-Kmeans.ipynb</a>) repeat the experiments in HW10.3. Explain any differences between the results in HW10.3 and HW10.4.

#### **Data Generation**

```
In [65]:
          %matplotlib inline
         import numpy as np
         import pylab
         import json
         size1 = size2 = size3 = 1000
         samples1 = np.random.multivariate normal([4, 0], [[1, 0], [0, 1]], size
         1)
         data = samples1
         samples2 = np.random.multivariate normal([6, 6], [[1, 0],[0, 1]], size
         data = np.append(data,samples2, axis=0)
         samples3 = np.random.multivariate_normal([0, 4], [[1, 0],[0, 1]], size
         3)
         data = np.append(data,samples3, axis=0)
         # Randomlize data
         data = data[np.random.permutation(size1+size2+size3),]
         np.savetxt('data.csv',data,delimiter = ',')
```

#### **Data Visualiazation**

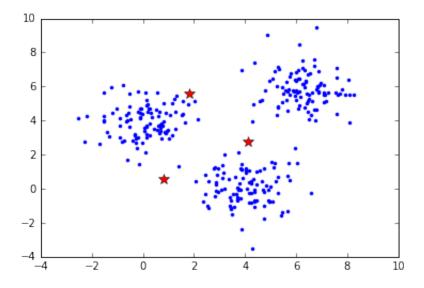


```
In [67]:
         import numpy as np
         #Calculate which class each data point belongs to
         def nearest centroid(line):
             x = np.array([float(f) for f in line.split(',')])
             closest centroid idx = np.sum((x - centroids)**2, axis=1).argmin()
             return (closest centroid idx,(x,1))
         #plot centroids and data points for each iteration
         def plot iteration(means):
             pylab.plot(samples1[:, 0], samples1[:, 1], '.', color = 'blue')
             pylab.plot(samples2[:, 0], samples2[:, 1], '.', color = 'blue')
             pylab.plot(samples3[:, 0], samples3[:, 1],'.', color = 'blue')
             pylab.plot(means[0][0], means[0][1],'*',markersize =10,color = 're
         d')
             pylab.plot(means[1][0], means[1][1],'*',markersize =10,color = 're
         d')
             pylab.plot(means[2][0], means[2][1],'*',markersize =10,color = 're
         d')
             pylab.show()
```

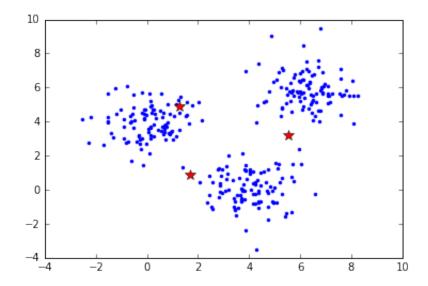
### **Distributed KMeans in Spark**

```
In [77]: | K = 3
         # Initialization: initialization of parameter is fixed to show an exam
         centroids = np.array([[0.0,0.0],[2.0,2.0],[0.0,7.0]])
         D = sc.textFile("./data.csv") #.cache()
         parsedData = D.map(lambda line: array([float(x) for x in line.split(',
         ')1))
         iter num = 0
         for i in range(10):
             res = D.map(nearest centroid).reduceByKey(lambda x,y : (x[0]+y[0],
         x[1]+y[1]).collect()
             res = sorted(res, key = lambda x : x[0]) #sort based on clusted ID
             centroids new = np.array([x[1][0]/x[1][1]  for x in res]) #divide
         by cluster size
             if np.sum(np.absolute(centroids new-centroids))<0.01:</pre>
                 break
             print "Iteration" + str(iter num)
             iter num = iter num + 1
             centroids = centroids new
             print centroids
             plot iteration(centroids)
             WSSSE = parsedData.map(lambda point: error(point)).reduce(lambda x
         , y: x + y
             print("Within Set Sum of Squared Error = " + str(WSSSE))
         print "Final Results:"
         print centroids
         Iteration0
         [[ 0.82299188  0.56796758]
```

```
[ 4.12230006 2.76406673]
[ 1.79893782 5.57176309]]
```

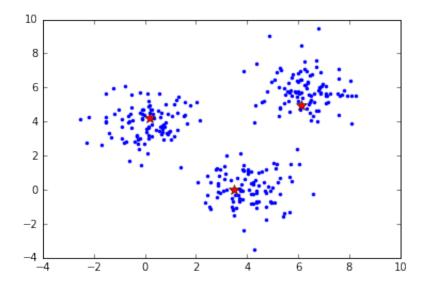


Within Set Sum of Squared Error = 358.527731123 Iteration1



Within Set Sum of Squared Error = 358.527731123 Iteration2

[ 0.20215824 4.19267863]]

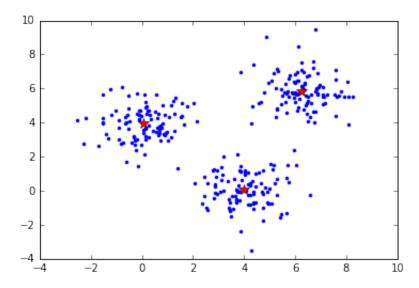


Within Set Sum of Squared Error = 358.527731123 Iteration3

[[ 4.00163563 0.05975173]

[ 6.26107859 5.83283378]

[ 0.05657819 3.97671718]]

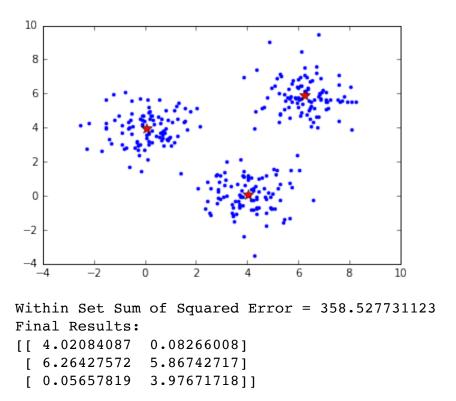


Within Set Sum of Squared Error = 358.527731123 Iteration4

[[ 4.02084087 0.08266008]

[ 6.26427572 5.86742717]

[ 0.05657819 3.97671718]]



#### **MLlib KMeans**

```
In [69]:
         from pyspark.mllib.clustering import KMeans, KMeansModel
         from numpy import array
         from math import sqrt
         # Load and parse the data
         data = sc.textFile("data.csv")
         parsedData = data.map(lambda line: array([float(x) for x in line.split
         (',')]))
         # Build the model (cluster the data)
         clusters = KMeans.train(parsedData, 3, maxIterations=20,
                 runs=10, initializationMode="random")
         for centroid in clusters.centers:
             print centroid
         [ 6.02635913  6.01593624]
         [ 0.00545017 3.97702197]
         [ 3.97598146 -0.02083839]
```

There doesn't seem to be much difference between the two codes, besides slightly faster convergence of the first one.

## **HW10.4.1: Making Homegrown KMeans more efficient**

**Back to Table of Contents** 

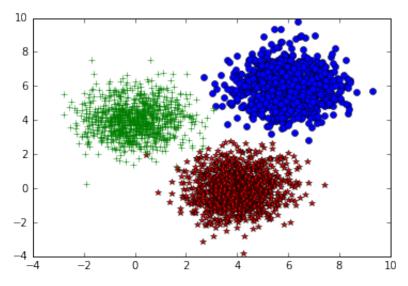
The above provided homegrown KMeans implentation in not the most efficient. How can you make it more efficient? Make this change in the code and show it work and comment on the gains you achieve.

# HINT: have a look at this linear regression notebook (<a href="http://nbviewer.jupyter.org/urls/dl.dropbox.com/s/atzqkc0p1eajuz6/LinearRegreNotebook-Challenge.ipynb">http://nbviewer.jupyter.org/urls/dl.dropbox.com/s/atzqkc0p1eajuz6/LinearRegreNotebook-Challenge.ipynb</a>)

This notebook hints to use broadcasting.

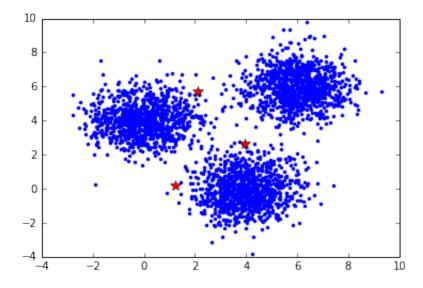
```
In [87]:
          %matplotlib inline
          import numpy as np
          import pylab
         import json
         size1 = size2 = size3 = 1000
         samples1 = np.random.multivariate normal([4, 0], [[1, 0],[0, 1]], size
         1)
         data = samples1
         samples2 = np.random.multivariate normal([6, 6], [[1, 0],[0, 1]], size
         2)
         data = np.append(data,samples2, axis=0)
         samples 3 = \text{np.random.multivariate normal}([0, 4], [[1, 0], [0, 1]], \text{size}
         3)
         data = np.append(data,samples3, axis=0)
         # Randomlize data
         data = data[np.random.permutation(size1+size2+size3),]
         np.savetxt('data.csv',data,delimiter = ',')
```

```
In [88]: pylab.plot(samples1[:, 0], samples1[:, 1],'*', color = 'red')
    pylab.plot(samples2[:, 0], samples2[:, 1],'o',color = 'blue')
    pylab.plot(samples3[:, 0], samples3[:, 1],'+',color = 'green')
    pylab.show()
```



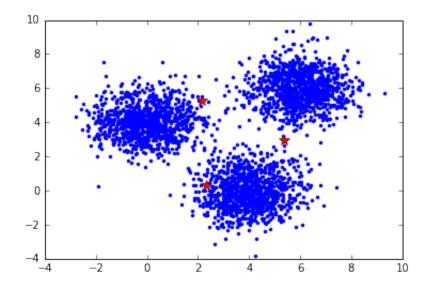
```
In [89]:
         import numpy as np
         #Calculate which class each data point belongs to
         def nearest centroid(line):
             x = np.array([float(f) for f in line.split(',')])
             closest centroid idx = np.sum((x - centroids)**2, axis=1).argmin()
             return (closest centroid idx_{i}(x,1))
         #plot centroids and data points for each iteration
         def plot iteration(means):
             pylab.plot(samples1[:, 0], samples1[:, 1], '.', color = 'blue')
             pylab.plot(samples2[:, 0], samples2[:, 1], '.', color = 'blue')
             pylab.plot(samples3[:, 0], samples3[:, 1],'.', color = 'blue')
             pylab.plot(means[0][0], means[0][1],'*',markersize =10,color = 're
         d')
             pylab.plot(means[1][0], means[1][1],'*',markersize =10,color = 're
         d')
             pylab.plot(means[2][0], means[2][1],'*',markersize =10,color = 're
         d')
             pylab.show()
```

```
In [102]:
          K = 3
          # Initialization: initialization of parameter is fixed to show an exam
          centroids = np.array([[0.0,0.0],[2.0,2.0],[0.0,7.0]])
          D = sc.textFile("./data.csv") #.cache()
          parsedData = D.map(lambda line: array([float(x) for x in line.split(',
          ')1))
          featureLen = len(parsedData.take(1)[0])-1
          n = parsedData.count()
          learningRate=0.05
          w = np.random.normal(size=featureLen) # w should be broadcasted if it
          is large
          wBroadcast = sc.broadcast(w) #make available in memory as read-only
          to the executors (for mappers and reducers)
          iter num = 0
          for i in range(10):
              wBroadcast = sc.broadcast(w) #make available in memory as read-o
          nly to the executors (for mappers and reducers)
              res = D.map(nearest centroid).reduceByKey(lambda x,y : (x[0]+y[0],
          x[1]+y[1]).collect()
              res = sorted(res, key = lambda x : x[0]) #sort based on clusted ID
              centroids new = np.array([x[1][0]/x[1][1]  for x in res]) #divide
          by cluster size
              if np.sum(np.absolute(centroids new-centroids))<0.01:</pre>
                  break
              print "Iteration" + str(iter num)
              iter num = iter num + 1
              centroids = centroids new
              print centroids
              plot_iteration(centroids)
              w = w - learningRate/n
              WSSSE = parsedData.map(lambda point: error(point)).reduce(lambda x
          , y: x + y
              #print("Within Set Sum of Squared Error = " + str(WSSSE))
          print "Final Results:"
          print centroids
          Iteration0
          [[ 1.22669683  0.16077825]
           [ 3.94327877 2.65087401]
           [ 2.10937398 5.70463155]]
```



#### Iteration1

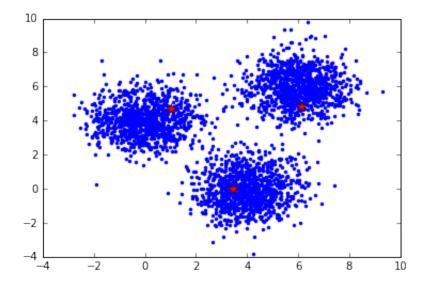
[[ 2.33581065 0.31989059] [ 5.38171434 2.92142592] [ 2.14111435 5.25234035]]



#### Iteration2

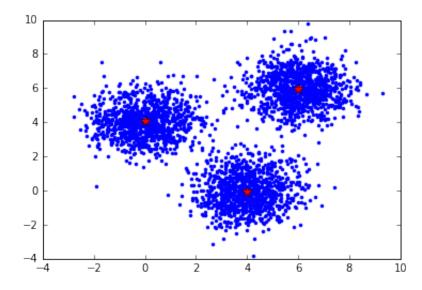
[[ 3.45392296 -0.02907791] [ 6.10580771 4.83576349]

[ 1.03525368 4.71383842]]



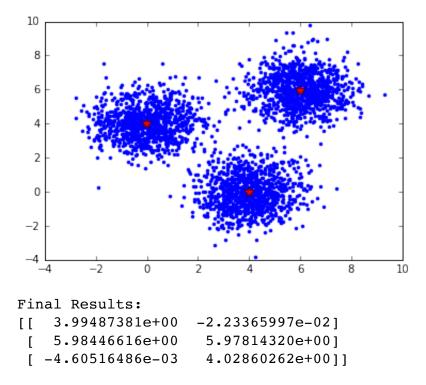
#### Iteration3

[[ 3.97168289 -0.0378151 ] [ 6.0099452 5.95150522] [ 0.02883925 4.05006856]]



#### Iteration4

[[ 3.99487381e+00 -2.23365997e-02] [ 5.98446616e+00 5.97814320e+00] [ -4.60516486e-03 4.02860262e+00]]



With broadcasting, we can make the serial tasks run in parallel. As a result, we made this process run faster.

# **HW10.5: OPTIONAL Weighted KMeans**

Back to Table of Contents

Using this provided <u>homegrown Kmeans code</u>

(http://nbviewer.jupyter.org/urls/dl.dropbox.com/s/3nsthvp8g2rrrdh/EM-Kmeans.ipynb), modify it to do a weighted KMeans and repeat the experiements in HW10.3. Explain any differences between the results in HW10.3 and HW10.5.

NOTE: Weight each example as follows using the inverse vector length (Euclidean norm):

```
weight(X)= 1/||X||,
```

where  $||X|| = SQRT(X.X) = SQRT(X1^2 + X2^2)$ 

Here X is vector made up of two values X1 and X2.

[Please incorporate all referenced notebooks directly into this master notebook as cells for HW submission. I.e., HW submissions should comprise of just one notebook]

```
In [71]: ## Code goes here
In [72]: ## Drivers & Runners
In [73]: ## Run Scripts, S3 Sync
```

## **HW10.6 OPTIONAL Linear Regression**

**Back to Table of Contents** 

## **HW10.6.1 OPTIONAL Linear Regression**

**Back to Table of Contents** 

Using this linear regression notebook

(http://nbviewer.jupyter.org/urls/dl.dropbox.com/s/atzqkc0p1eajuz6/LinearRegression-Notebook-Challenge.ipynb):

- Generate 2 sets of data with 100 data points using the data generation code provided and plot each in separate plots. Call one the training set and the other the testing set.
- Using MLLib's LinearRegressionWithSGD train up a linear regression model with the training dataset and evaluate with the testing set. What a good number of iterations for training the linear regression model? Justify with plots (e.g., plot MSE as a function of the number of iterations) and words.

## **HW10.6.2 OPTIONAL Linear Regression**

**Back to Table of Contents** 

In the notebook provided above, in the cell labeled "Gradient descent (regularization)".

- Fill in the blanks and get this code to work for LASSO and RIDGE linear regression.
- Using the data from HW10.6.1 tune the hyper parameters of your LASS0 and RIDGE regression.
   Report your findings with words and plots.

```
In [74]: ## Code goes here
In [75]: ## Drivers & Runners
In [76]: ## Run Scripts, S3 Sync
```

#### **HW10.7 OPTIONAL Error surfaces**

**Back to Table of Contents** 

Here is a link to R code with 1 test drivers that plots the linear regression model in model space and in the domain space:

https://www.dropbox.com/s/3xc3kwda6d254l5/PlotModelAndDomainSpaces.R?dl=0 (https://www.dropbox.com/s/3xc3kwda6d254l5/PlotModelAndDomainSpaces.R?dl=0)

Here is a sample output from this script:

https://www.dropbox.com/s/my3tnhxx7fr5qs0/image%20%281%29.png?dl=0 (https://www.dropbox.com/s/my3tnhxx7fr5qs0/image%20%281%29.png?dl=0)

Please use this as inspiration and code a equivalent error surface and heatmap (with isolines) in Spark and show the trajectory of learning taken during gradient descent(after each n-iterations of Gradient Descent):

Using Spark and Python (using the above R Script as inspiration), plot the error surface for the linear regression model using a heatmap and contour plot. Also plot the current model in the original domain space for every 10th iteration. Plot them side by side if possible for each iteration: lefthand side plot is the model space(w0 and w01) and the righthand side plot is domain space (plot the corresponding model and training data in the problem domain space) with a final pair of graphs showing the entire trajectory in the model and domain space. Make sure to label your plots with iteration numbers, function, model space versus original domain space, MSE on the training data etc.

Also plot the MSE as a function of each iteration (possibly every 10th iteration). Dont forget to label both axis and the graph also. [Please incorporate all referenced notebooks directly into this master notebook as cells for HW submission. I.e., HW submissions should comprise of just one notebook]

```
In [77]: ## Code goes here
In [78]: ## Drivers & Runners
In [79]: ## Run Scripts, S3 Sync
```

#### **Back to Table of Contents**

# ----- END OF HOWEWORK ------

In [ ]:	]:	