HW11, DATSCI W261

Team: Kuan Lin, Alejandro J. Rojas, Ricardo Barrera

Emails: kuanlin@ischool.berkeley.edu, ale@ischool.berkeley.edu,

ricardofrank@ischool.berkeley.edu

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W261-1, Spring 2016 Week 11 Homework

Took 0 seconds. (outdated)

HW11.0 Broadcast versus Caching in Spark

What is the difference between broadcasting and caching data in Spark? Give an example (in the context of machine learning) of each mechanism (at a highlevel). Feel free to cut and paste code examples from the lectures to support your answer.

Review the following Spark-notebook-based implementation of KMeans and use the broadcast pattern to make this implementation more efficient. Please describe your changes in English first, implement, comment your code and highlight your changes:

Notebook

https://www.dropbox.com/s/41q9lgyqhy8ed5g/EM-Kmeans.ipynb?dl=0

Notebook via NBViewer

http://nbviewer.ipython.org/urls/dl.dropbox.com/s/41q9lgyqhy8ed5g/EM-Kmeans.ipynb Took 0 seconds. (outdated)

```
%pyspark
# zeppelin interop to matplotlib
import StringIO
import matplotlib.pyplot as plt
def show(p):
    img = StringIO.StringIO()
    p.savefig(img, format='svg')
    img.seek(0)
    print "%html <div style='width:600px'>" + img.buf + "</div>"
```

Took 99 seconds.

Caching vs. Broadcasting

Caching data in Sapark is useful to keep in memory data that you need to process multiple times. An example of it would be cacg=hing a training dataset. On the other hand boradcasting is implemented whem you need to let worker nodes the status of a specific variable. For example when running logistic regression you need to broadcast the value of the weights after each iteration so that worker nodes can process gradient descent using the most current weights.

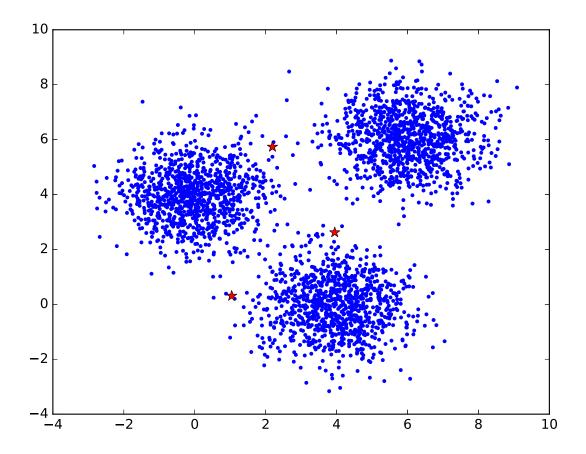
To make the K-Means code more efficient we will boradcast the values of the centroids of each cluster so that the worker nodes get that info on their own memory Took 0 seconds. (outdated)

Broadcasted K-Mean

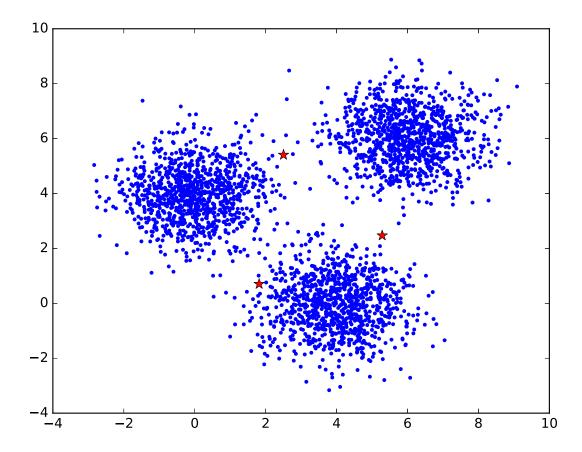
Took 0 seconds. (outdated)

```
%pyspark
# generate data
import numpy as np
size1 = size2 = size3 = 1000
samples1 = np.random.multivariate_normal([4, 0], [[1, 0],[0, 1]], size1)
data = samples1
samples2 = np.random.multivariate_normal([6, 6], [[1, 0],[0, 1]], size2)
data = np.append(data,samples2, axis=0)
samples3 = np.random.multivariate_normal([0, 4], [[1, 0],[0, 1]], size3)
data = np.append(data,samples3, axis=0)
# Randomlize data
data = data[np.random.permutation(size1+size2+size3),]
np.savetxt('/data/data.csv',data,delimiter = ',')
import pylab
# 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\_broadcasted.value)**2, axis=1).argmin() # use
    return (closest centroid idx,(x,1))
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 = 'red')
pylab.plot(means[1][0], means[1][1],'*',markersize =10,color = 'red')
pylab.plot(means[2][0], means[2][1],'*',markersize =10,color = 'red')
    #pylab.show()
    show(pylab.plt)
# Initialization: initialization of parameter is fixed to show an example
centroids = np.array([[0.0,0.0],[2.0,2.0],[0.0,7.0]])
```

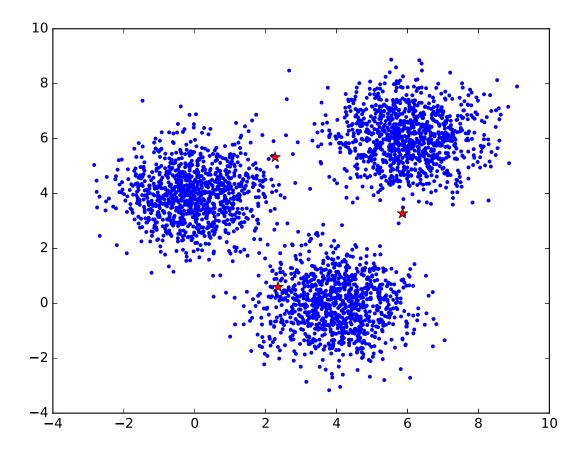
```
D = sc.textFile("file:///data/data.csv").cache()
iter_num = 0
for i in range(10):
    centroids broadcasted = sc.broadcast(centroids) # broadcast centroids before any distribute
    res = D.map(nearest\_centroid).reduceByKey(lambda x,y : (x[0]+y[0],x[1]+y[1])).collect()
    res = sorted(res, key = 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>
    print "Iteration" + str(iter_num)
    iter_num = iter_num + 1
    centroids = centroids_new
    print centroids
    plot_iteration(centroids)
print "Final Results:"
print centroids
```



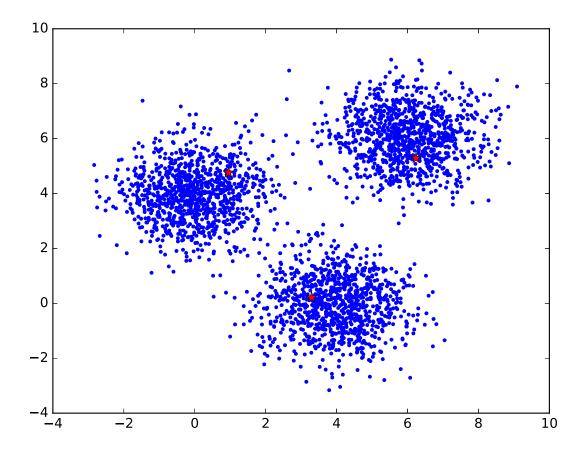
Iteration1 [[1.81718093 0.70375309] [5.28602178 2.46978436] [2.50336223 5.407373]] %html



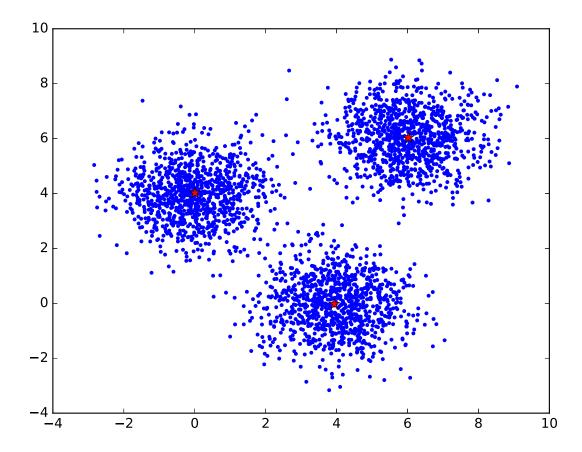
 $Iteration 2 \hbox{ [[2.34995127 \ 0.59701223] [5.8580947 \ 3.27051641] [2.26729464 \ 5.31927826]] } \% html. \\$



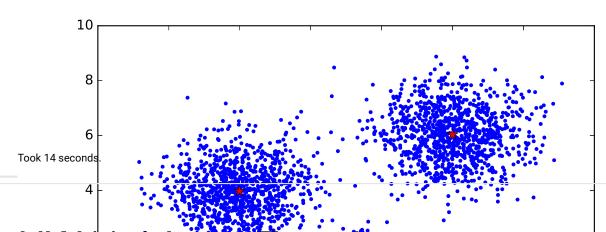
 $Iteration 3 \; [[\; 3.29638299\; 0.21308633]\; [\; 6.23980831\; 5.28776201]\; [\; 0.95424176\; 4.7580034\;]]\; \% html$



 $Iteration 4 \hbox{ [[3.93649784 -0.02283234] [6.027216 \ 6.03508886] [0.0076037 \ 4.02187443]] \ \% html} \\$



 $Iteration 5 \ [[\ 3.966843\ -0.0419858\]\ [\ 6.00188246\ 6.0371927\]\ [-0.00836659\ 3.97631336]]\ \% html$



HW41. unctions

In the context of binary classification problems, tipes the linear SVM learning algorithm yield the same result as a L2 penalized logistic regession learning alog thm?

In your reponse, please discuss the loss functions, and the learnt models, and separating surfaces between_the two classes.

In the context of binary classification problems, does the linear SVM learning algorithm yield the same result as a perceptron learning algorithm?

[OPTIONAL] generate an artifical binary classification dataset with 2 input features and plot the learnt separating surface for both a linear SVM and for logistic regression. Comment on the learnt surfaces. Please feel free to do this in Python (no need to use Spark). Final Results: [[3.966843 -0.0419858] [6.00188246 6.0371927] [-0.00836659 3.97631336]]

In the context of binary clasification problems, linear SVM should yield similar result as a L2penalized logistic regression, as both algorithm pushes to have some positive margin. However, the SVM loss function does not attempt to further distinguish non support vectors, that is, data points with margins greater than 1. This is different from logistic regression which will preferentially push for more margin if possible.

Linear SVM will not likely yield the same result as a perceptron learning algorithm. The perceptron algorithm does not provide further possibility for gradient descent as long as the data point is classified correctly, even with only a very small margin.

Took 6 seconds. (outdated)

HW11.2 Gradient descent

In the context of logistic regression describe and define three flavors of penalized loss functions. Are these all supported in Spark MLLib (include online references to support your answers)? Descibe probabilitic interpretations of the L1 and L2 priors for penalized logistic regression (HINT: see synchronous slides for week 11 for details)

Took 2 seconds. (outdated)

In the context of logistic regression, the three flavors of penalized terms are:

· L1 Reg, which penalizes for sum of absolute weights:

$$l_{reg}(w) = \lambda \sum ||w_i||$$

L2 Reg, penalizes sum of squared weights:

$$l_{reg}(w) = \lambda \sum w_i^2$$

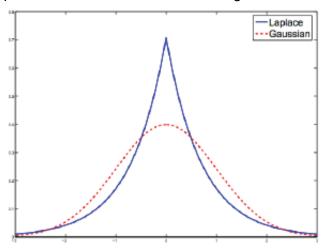
• Elastic Net, penalizes a linear combination of L1 and L2 norms:

$$\alpha \cdot ||w|| + (1 - \alpha) \cdot 1/2 \cdot ||w||^2$$

All of the above three regularization methods are supported by spark.mllib: http://spark.apache.org/docs/latest/mllib-linear-methods.html#regularizers

Probablisitic interpretation of L1 and L2 priors:

L1 regularization can be interpreted as using Laplace distribution as the prior distribution for the model weights, where as L2 regularization can be interpreted as using gaussian distribution as the prior distribution for the model weights.



The Laplace distribution has more density closer to mean (usually zero in most settings) in comparison to the gaussian distribution, and therefore L1 regularization will tend to push the model weights toward zero.

Took 12 seconds. (outdated)

HW11.3 Logistic Regression

Generate 2 sets of linearly separable data with 100 data points each using the data generation code provided below and plot each in separate plots. Call one the training set and the other the testing set.

def generateData(n):

generates a 2D linearly separable dataset with n samples.

The third element of the sample is the label

xb = (rand(n)2-1)/2-0.5

yb = (rand(n)2-1)/2+0.5

xr = (rand(n)2-1)/2+0.5

yr = (rand(n)2-1)/2-0.5

inputs = []

for i in range(len(xb)):

inputs.append([xb[i],yb[i],1])

inputs.append([xr[i],yr[i],-1])

return inputs

Modify this data generation code to generating non-linearly separable training and testing datasets (with approximately 10% of the data falling on the wrong side of the separating hyperplane. Plot the resulting datasets.

NOTE: For the remainder of this problem please use the non-linearly separable training and testing datasets.

Using MLLib train up a LASSO logistic regression model with the training dataset and evaluate with the testing set. What a good number of iterations for training the logistic regression model? Justify with plots and words.

Derive and implement in Spark a weighted LASSO logistic regression. Implement a convergence test of your choice to check for termination within your training algorithm.

Weight the above training dataset as follows: Weight each example 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 X1 and X2.

Evaluate your homegrown weighted LASSO logistic regression on the test dataset. Report misclassification error (1 - Accuracy) and how many iterations does it took to converge.

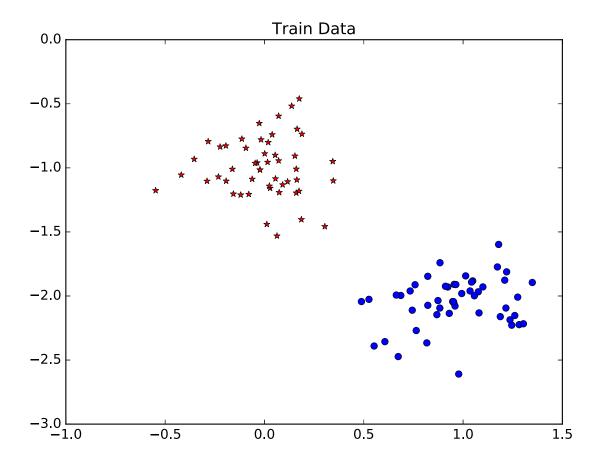
Does Spark MLLib have a weighted LASSO logistic regression implementation. If so use it and report your findings on the weighted training set and test set.

Took 7 seconds. (outdated)

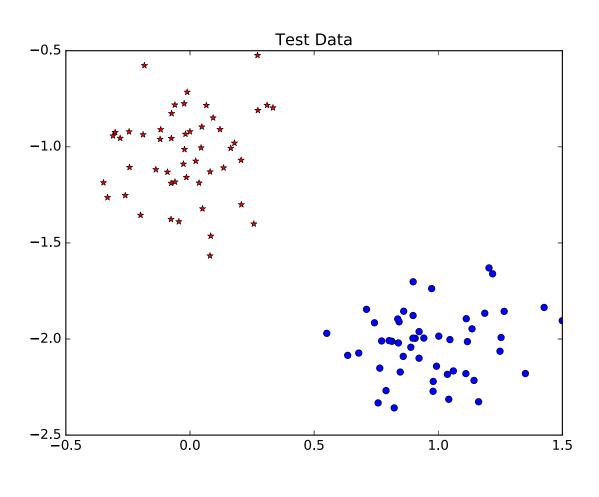
```
%pyspark
import numpy as np
np.random.seed(0)
def generateData(n):
    generates a 2D linearly separable dataset with n samples.
    The third element of the sample is the label
    xb = (np.random.normal(1,0.2,n)*2-1)/2-0.5
    yb = (np.random.normal(-1,0.2,n)*2-1)/2+0.5
    xr = (np.random.normal(1,0.2,n)*2-1)/2+0.5
    yr = (np.random.normal(-1,0.2,n)*2-1)/2-0.5
    inputs = []
    for i in range(len(xb)):
        inputs.append([xb[i],yb[i],1])
        inputs.append([xr[i],yr[i],-1])
    return inputs
```

Took 2 seconds.

```
%pyspark
import pylab
# generate 100 linearly seperatable data and plot them
data_lin_seperable_train = generateData(50) # train data
data lin seperable test = generateData(50) # test data
pylab.plot(\lceil d\lceil \theta \rceil for d in data lin seperable train if d\lceil 2 \rceil = 1 \rceil, \lceil d\lceil 1 \rceil for d in data lin seperak
pylab.plot(\lceil d\lceil \theta \rceil for d in data lin seperable train if d\lceil 2 \rceil = -1 \rceil, \lceil d\lceil 1 \rceil for d in data lin sepera
pylab.plt.title("Train Data")
#pylab.show()
show(pylab.plt)
pylab.plot([d[0] for d in data_lin_seperable_test if d[2]==1], [d[1] for data_lin_seperable_test if d[2]==1], [d[1] for data_lin_seperable_test if d[2]==1], [d[1] for data_
pylab.plot([d[0] for d in data_lin_seperable_test if d[2]==-1], [d[1] for d in data_lin_seperable_test if d[2]=-1], [d[1] for d in data_lin_seperable_test if d[2]=-1
pylab.plt.title("Test Data")
#pylab.show()
show(pylab.plt)
```



%html



Modify this data generation code to generating non-linearly separable training and testing datasets (with approximately 10% of the data falling on the wrong side of the separating hyperplane. Plot the resulting datasets

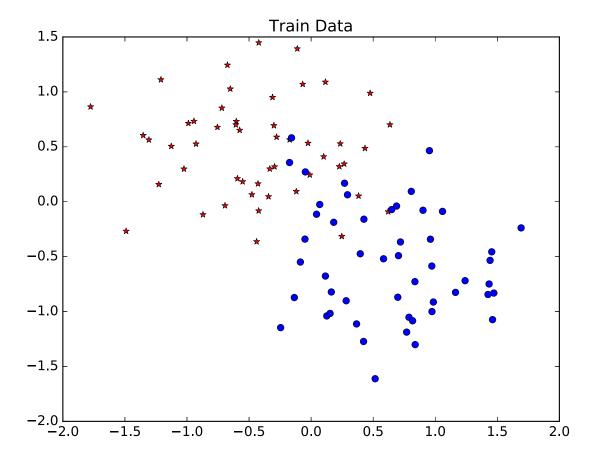
Took 6 seconds. (outdated)

Took 12 seconds.

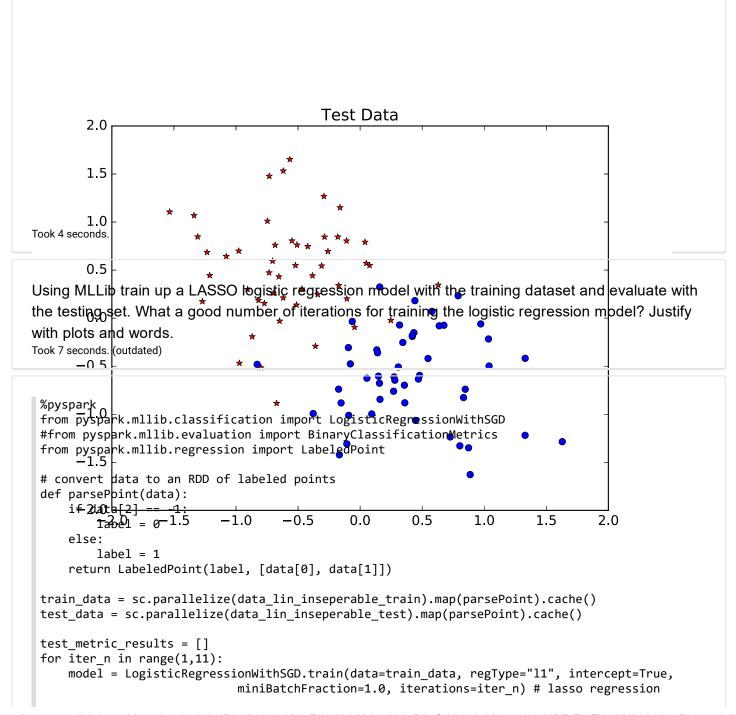
Took 3 seconds.

```
%pyspark
import numpy as np
np.random.seed(0)
def generateData2(n):
    non-linearly seperable data
    xb = np.random.normal(0,0.5,n)-0.5
    yb = np.random.normal(0,0.5,n)+0.5
    xr = np.random.normal(0,0.5,n)+0.5
    yr = np.random.normal(0,0.5,n)-0.5
    inputs = []
    for i in range(len(xb)):
        inputs.append([xb[i],yb[i],1])
        inputs.append([xr[i],yr[i],-1])
    return inputs
```

```
%pyspark
import pylab
# generate 100 linearly seperatable data and plot them
data_lin_inseperable_train = generateData2(50) # train data
data_lin_inseperable_test = generateData2(50) # test data
pylab.plot([d[0] for d in data_lin_inseperable_train if d[2]==1], [d[1] for data_lin_inseperable_train if d[2]==1], [d[1] for data_lin_inseperable_train if d[2]==1], [d[1] fo
pylab.plot(\lceil d \lceil 0 \rceil for d in data lin inseperable train if d \lceil 2 \rceil = -1 \rceil, \lceil d \lceil 1 \rceil for d in data lin inse
pylab.plt.title("Train Data")
#pylab.show()
show(pylab.plt)
pylab.plot([d[0] \ for \ d \ in \ data\_lin\_inseperable\_test \ if \ d[2]==1], \ [d[1] \ for \ d \ in \ data\_lin\_inseperable\_test \ if \ d[2]==1], \ [d[1] \ for \ d \ in \ data\_lin\_inseperable\_test \ if \ d[2]==1], \ [d[1] \ for \ d \ in \ data\_lin\_inseperable\_test \ if \ d[2]==1], \ [d[1] \ for \ d \ in \ data\_lin\_inseperable\_test \ if \ d[2]==1], \ [d[1] \ for \ d \ in \ data\_lin\_inseperable\_test \ if \ d[2]==1], \ [d[1] \ for \ d \ in \ data\_lin\_inseperable\_test \ if \ d[2]==1], \ [d[1] \ for \ d \ in \ data\_lin\_inseperable\_test \ if \ d[2]==1], \ [d[1] \ for \ d \ in \ data\_lin\_inseperable\_test \ if \ d[2]==1], \ [d[1] \ for \ d \ in \ data\_lin\_inseperable\_test \ if \ d[2]==1], \ [d[1] \ for \ d \ in \ data\_lin\_inseperable\_test \ if \ d[2]==1], \ [d[1] \ for \ d \ in \ data\_lin\_inseperable\_test \ if \ d[2]==1], \ [d[1] \ for \ d \ in \ data\_lin\_inseperable\_test \ if \ d[2]==1], \ [d[1] \ for \ d \ in \ data\_lin\_inseperable\_test \ if \ d[2]==1], \ [d[1] \ for \ d \ in \ data\_lin\_inseperable\_test \ if \ d[2]==1], \ [d[1] \ for \ d \ in \ data\_lin\_inseperable\_test \ if \ d[2]==1], \ [d[1] \ for \ d \ in \ data\_lin\_inseperable\_test \ if \ d[2]==1], \ [d[1] \ for \ d \ in \ data\_lin\_inseperable\_test \ if \ d[2]==1], \ [d[1] \ for \ d \ in \ data\_lin\_inseperable\_test \ if \ d[2]==1], \ [d[1] \ for \ d \ in \ data\_lin\_inseperable\_test \ if \ d[2]==1], \ [d[1] \ for \ d \ in \ data\_lin\_inseperable\_test \ if \ d[2]==1], \ [d[1] \ for \ d \ in \ data\_lin\_inseperable\_test \ if \ d[2]==1], \ [d[1] \ for \ d \ in \ data\_lin\_inseperable\_test \ if \ d[2]==1], \ [d[1] \ for \ d \ in \ data\_lin\_inseperable\_test \ in \ d[2]==1], \ [d[1] \ for \ d \ in \ d[2]==1], \ [d[1] \ for \ d \ in \ d[2]==1], \ [d[1] \ for \ d \ in \ d[2]==1], \ [d[1] \ for \ d \ in \ d[2]==1], \ [d[1] \ for \ d \ in \ d[2]==1], \ [d[1] \ for \ d \ in \ d[2]==1], \ [d[1] \ for \ d \ in \ d[2]==1], \ [d[1] \ for \ d \ in \ d[2]==1], \ [d[1] \ for \ d \ in \ d[2]==1], \ [d[1] \ for \ d \ in \ d[2]==1], \ [d[1] \ for \ d \ in \ d[2]==1], \ [d[1] \ for \ d 
pylab.plot([d[0] for d in data_lin_inseperable_test if d[2]==-1], [d[1] for d in data_lin_inser
pylab.plt.title("Test Data")
#pylab.show()
show(pylab.plt)
```

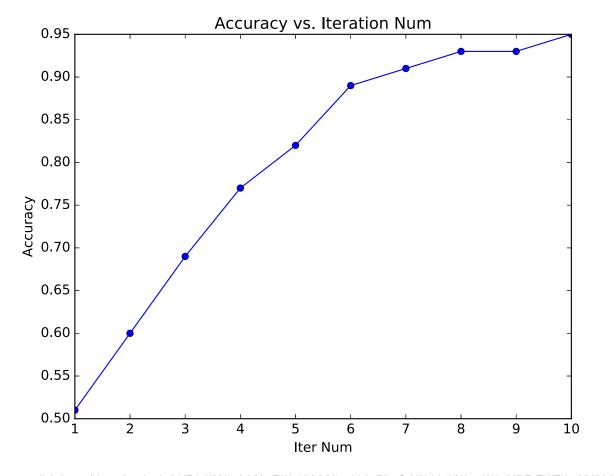


%html



```
accuracy = test_data.map(lambda lp: ((float(model.predict(lp.features))==lp.label), 1)).rec
     accuracy = 1.0*accuracy[0]/accuracy[1]
     test_metric_results.append((iter_n,accuracy))
     print "Iter %s | Accuracy:%.4f" %(iter_n, accuracy)
Iter 1 | Accuracy:0.5100
Iter 2 | Accuracy:0.6000
Iter 3 | Accuracy:0.6900
Iter 4 | Accuracy:0.7700
Iter 5 | Accuracy:0.8200
Iter 6 | Accuracy:0.8900
Iter 7 | Accuracy:0.9100
Iter 8 | Accuracy:0.9300
Iter 9 | Accuracy:0.9300
Iter 10 | Accuracy:0.9500
Took 46 seconds.
```

```
%pyspark
import matplotlib.pyplot as plt
plt.plot([d[0] for d in test_metric_results], [d[1] for d in test_metric_results], '-o')
plt.title("Accuracy vs. Iteration Num")
plt.xlabel("Iter Num")
plt.ylabel("Accuracy")
show(plt)
```



Took 5 seconds.

Based on accuracy, the SGD implementation of LogisticRegression with L1 regularization seems to begin to level-out when iteration is about 8 or more.

Took 7 seconds. (outdated)

Derive and implement in Spark a weighted LASSO logistic regression. Implement a convergence test of your choice to check for termination within your training algorithm.

Weight the above training dataset as follows: Weight each example using the inverse vector length (Euclidean norm):

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

```
where ||X|| = SQRT(X.X) = SQRT(X1^2 + X^22)
```

Here X is vector made up of X1 and X2.

Evaluate your homegrown weighted LASSO logistic regression on the test dataset. Report misclassification error (1 - Accuracy) and how many iterations does it took to converge.

Does Spark MLLib have a weighted LASSO logistic regression implementation. If so use it and report your findings on the weighted training set and test set.

Took 7 seconds. (outdated)

```
%pyspark
import numpy as np
def readPoint(data):
    label = data[2]
    x = [data[0], data[1], 1.0] #add bias term
    return (x, label)
def vectorWeight(v1, v2):
    weight = 1.0/((v1**2+v2**2)**0.5)
    if weight < 0.1:
       weight = 0.1
    elif weight > 10:
       weight = 10
    return weight
def WeightedlogisticRegressionGD(data, wInitial=None, learningRate=0.05, iterations=10, regPara
    featureLen = len(data.take(1)[0][0])
    #total_weight = data.count()
    total_weight = data.map(lambda p: vectorWeight(p[0][0], p[0][1])).reduce(lambda a,b: a+b)
    if wInitial is None:
        w = np.random.normal(size=featureLen) # w should be broadcasted if it is large
    else:
        w = wInitial
    for i in range(iterations):
        #print "Iteration %s"%(i+1)
        wBroadcast = sc.broadcast(w)
        gradient = data.map(lambda p: vectorWeight(p[0][0], p[0][1])*(1 / (1 + np.exp(-p[1]*np.
                    .reduce(lambda a, b: a + b)
        #gradient = data.map(lambda p: (1 / (1 + np.exp(-p[1]*np.dot(wBroadcast.value, p[0]))).
                     .reduce(lambda a, b: a + b)
        if regType == "Ridge":
            wReg = w * 1
            wReg[-1] = 0 #last value of weight vector is bias term, ignored in regularization
        elif regType == "Lasso":
```

```
wReg = w * 1
       wReg[-1] = 0 #last value of weight vector is bias term, ignored in regularization
       wReg = (wReg>0).astype(int) * 2-1
    else:
       wReg = np.zeros(w.shape[0])
    gradient = gradient + regParam * wReg #gradient: GD of Sqaured Error+ GD of regularia
    wdelta = learningRate * gradient / total_weight # scale by total weight
    if sum(abs(wdelta))<=stopCriteria*sum(abs(w)): #Convergence condition
        print "convergenced reached at iteration %"%(i+1)
        break
    #w = w - learningRate * gradient / n
    w = w - wdelta
    #print w
print "total iterations: %s"%(i+1)
return w
```

Took 5 seconds.

```
%pyspark
 train_data = sc.parallelize(data_lin_inseperable_train).map(readPoint).cache()
 test data = sc.parallelize(data lin inseperable test).map(readPoint).cache()
 w = WeightedlogisticRegressionGD(train data, regType="Lasso", stopCriteria=0.001, iterations=56
 print "final weight:%s"%str(w)
total iterations: 50
final weight: [ 0.48698643    1.39042164 -0.08542408]
Took 14 seconds.
```

```
%pyspark
# evaluate on test set
def isAccurate(x, w, label):
    score = 1.0/(1.0 + np.exp(-1.0*np.dot(w,x)))
    print "label|score: %s | %.4f"%(label, score)
    if (label == 1 and score >= 0.5) or (label == -1 and score < 0.5):
       return 1
    else:
       return 0
def LogisticRegAccuracy(data, w):
    wBroadcast = sc.broadcast(w)
    return result[0]*1.0/result[1]
accuracy = LogisticRegAccuracy(test data, w)
print "Misclassification Error Rate: %s"%(str(100-round(accuracy*100, 4))+"%")
Misclassification Error Rate: 23.0%
```

Took 5 seconds.

HW11.4 SVMs

Use the non-linearly separable training and testing datasets from HW11.3 in this problem. Using MLLib train up a soft SVM model with the training dataset and evaluate with the testing set. What is a good number of iterations for training the SVM model? Justify with plots and words. Derive and Implement in Spark a weighted soft linear sym classification learning algorithm. Evaluate your homegrown weighted soft linear sym classification learning algorithm on the weighted training dataset and test dataset from HW11.3. Report misclassification error (1 - Accuracy) and how many iterations does it took to converge? How many support vectors do you end up?

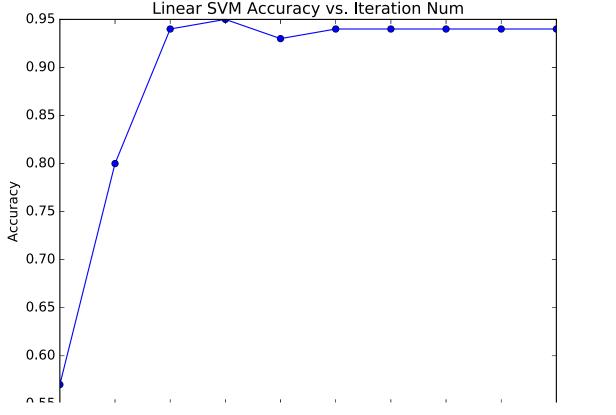
Does Spark MLLib have a weighted soft SVM learner. If so use it and report your findings on the weighted training set and test set.

Took 6 seconds. (outdated)

```
%pyspark
# using mllib
from pyspark.mllib.classification import SVMWithSGD
#from pyspark.mllib.evaluation import BinaryClassificationMetrics
from pyspark.mllib.regression import LabeledPoint
# convert data to an RDD of labeled points
def parsePoint(data):
    if data[2] == -1:
        label = 0
    else:
        label = 1
    return LabeledPoint(label, [data[0], data[1]])
train data = sc.parallelize(data lin inseperable train).map(parsePoint).cache()
test_data = sc.parallelize(data_lin_inseperable_test).map(parsePoint).cache()
test_metric_results = []
for iter n in range(1,11):
    model = SVMWithSGD.train(data=train_data, regType="11", intercept=True,
                            miniBatchFraction=1.0, iterations=iter_n) # lasso, same parameters
    accuracy = test_data.map(lambda lp: ((float(model.predict(lp.features))==lp.label), 1)).rec
    accuracy = 1.0*accuracy[0]/accuracy[1]
    test metric_results.append((iter_n,accuracy))
    print "Iter %s | Accuracy:%.4f" %(iter_n, accuracy)
```

```
Iter 1 | Accuracy:0.5700
Iter 2 | Accuracy:0.8000
Iter 3 | Accuracy:0.9400
Iter 4 | Accuracy: 0.9500
Iter 5 | Accuracy:0.9300
Iter 6 | Accuracy:0.9400
Iter 7 | Accuracy:0.9400
Iter 8 | Accuracy:0.9400
Iter 9 | Accuracy:0.9400
Iter 10 | Accuracy:0.9400
Took 24 seconds.
```

```
%pyspark
import matplotlib.pyplot as plt
plt.plot([d[0] for d in test_metric_results], [d[1] for d in test_metric_results], '-o')
plt.title("Linear SVM Accuracy vs. Iteration Num")
plt.xlabel("Iter Num")
plt.ylabel("Accuracy")
show(plt)
```



Zeppelin**Hub** (/)Viewer (/viewer)

MfDS-W261-2016_HWK11-Week11-Team5.ipynb

 \ominus

For this particular dataset LinearSVM seems to converge very fast, plateauing at 94% accuracy after 3 iterations. This is much faster than logistic regression.

Took 8 seconds. (outdated)

```
%pyspark
# home-grown SVM code
def vectorWeight(v1, v2):
    weight = 1.0/((v1**2+v2**2)**0.5)
    if weight < 0.1:
        weight = 0.1
    elif weight > 10:
       weight = 10
    return weight
def WeightedSVMRegressionGD(data, wInitial=None, learningRate=0.05, iterations=100, regParam=0.
    featureLen = len(data.take(1)[0][0])
    total_weight = data.map(lambda p: vectorWeight(p[0][0], p[0][1])).reduce(lambda a,b: a+b)
    if wInitial is None:
        w = np.random.normal(size=featureLen) # w should be broadcasted if it is large
    else:
        w = wInitial
    for i in range(iterations):
        wBroadcast = sc.broadcast(w)
        sv = data.filter(lambda p: p[1]*np.dot(wBroadcast.value, p[0]) < 1).cache() # find the
        if sv.isEmpty(): # no more support vectors
            break
        # calculate weighted gradients (hinge loss)
        gradient = -1.0*sv.map(lambda p: vectorWeight(p[0][0], p[0][1])*p[1]*np.array(p[0])).re
        if regType == "Ridge":
            wReg = w * 1
            wReg[-1] = 0 #last value of weight vector is bias term, ignored in regularization
        elif regType == "Lasso":
            wReg = w * 1
            wReg[-1] = 0 #last value of weight vector is bias term, ignored in regularization
            wReg = (wReg>0).astype(int) * 2-1
        else:
            wReg = np.zeros(w.shape[0])
        gradient = gradient + regParam * wReg #gradient: GD of Sqaured Error+ GD of regulariz
        wdelta = learningRate * gradient
        if sum(abs(wdelta))<=stopCriteria*sum(abs(w)): #Convergence condition
            print "convergenced reached at iteration %"%(i+1)
            break
        w = w - wdelta
        #print "iteration %s"%i
        #print w
    return (w, i+1, sv.count()) # learned weights, total iters, support vector counts
```

Took 6 seconds.

```
%pyspark
 train_data = sc.parallelize(data_lin_inseperable_train).map(readPoint).cache()
 test_data = sc.parallelize(data_lin_inseperable_test).map(readPoint).cache()
 w = WeightedSVMRegressionGD(train_data, regType="Lasso", stopCriteria=0.001, iterations=50, reg
 print "final weight:%s | total iters:%s | SV counts: %s"%(str(w[0]), str(w[1]), str(w[2]))
 # eval with test data
 def isAccurateSVM(x, w, label):
    score = label*np.dot(w, x)
    print "label|score: %s | %.4f"%(label, score)
    if score >= 0:
        return 1
    else:
        return 0
 def SVMRegAccuracy(data, w):
    wBroadcast = sc.broadcast(w)
    result = data.map(lambda p: (isAccurateSVM(p[0], wBroadcast.value, p[1]), 1)).reduce(lambda
    return result[0]*1.0/result[1]
 accuracy = SVMRegAccuracy(test_data, w[0])
 print "Misclassification Error Rate: %s"%(str(100-round(accuracy*100, 4))+"%")
Misclassification Error Rate: 5.0%
Took 20 seconds.
```

11.5 [OPTIONAL] Distributed Perceptron algorithm

Using the following papers as background:

http://static.googleusercontent.com/external content/untrusted dlcp/research.google.com/en//pubs/

https://www.dropbox.com/s/a5pdcp0r8ptudgj/gesmundo-tomeh-eacl-2012.pdf?dl=0

http://www.slideshare.net/matsubaray/distributed-perceptron

Implement each of the following flavors of perceptron learning algorithm:

- Serial (All Data): This is the classifier returned if trained serially on all the available data. On a single computer for example (Mistake driven)
- Serial (Sub Sampling): Shard the data, select one shard randomly and train serially.
- Parallel (Parameter Mix): Learn a perceptron locally on each shard: Once learning is

complete combine each learnt percepton using a uniform weighting

• Parallel (Iterative Parameter Mix) as described in the above papers.

```
Took 6 seconds. (outdated)
```

```
%pyspark

# perceptron gradient descent calculation

def perceptronSubGradientCalc(data, w, regParam, feature_size):
    #gradient = data.map(lambda p: -p[1]*np.array(p[0]) if (p[1]*np.dot(w.value,p[0]))<0 else r
    incorrect = data.filter(lambda p: np.dot(w.value,p[0])*p[1] < 0)
    if incorrect.isEmpty():
        gradient = np.zeros(feature_size)
    else:
        gradient = incorrect.map(lambda p: -p[1]*np.array(p[0])).reduce(lambda a,b: a+b)
    wreg = np.array(w.value)  # L2 reg
    wreg[-1] = 0 # don't count the bias term in reg
    return gradient/data.count() + regParam*wreg

def perceptronAccuracy(data, w):
    correct_count = data.map(lambda p: 1 if np.dot(w.value, p[0])*p[1] > 0 else 0).reduce(lambc)
    return 1.0*correct_count/data.count()
```

Took 6 seconds.

```
%pyspark
 # train serially with all data
 def perceptronGDSerialAll(data, wInitial=None, nShards=1, nIter=10, stopCriteria=0.001,
                  learningRate=0.05, regParam=0.01):
     feature size = len(data.take(1)[0][0])
     if wInitial is None:
         w = np.random.normal(size=feature_size)
     else:
         w = wInitial
     #model data = data.randomSplit([1.0/nShards]*nShards) # split data into shards
     for i in range(nIter):
         wb = sc.broadcast(w) # broadcast weight
         wdelta = learningRate*perceptronSubGradientCalc(data, wb, regParam, feature size)
         if sum(abs(wdelta))<=stopCriteria*sum(abs(w)):</pre>
             break
         w = w - wdelta
     return w
 w = perceptronGDSerialAll(train_data, nIter=50)
 print "Train serially on all data"
 print "learned weight: %s" % str(w)
 print "Accuracy: %s" % perceptronAccuracy(test_data, sc.broadcast(w))
Train serially on all data
learned weight: [-0.36552607 0.00169171 0.05655527]
Accuracy: 0.83
```

Took 21 seconds.

```
%pyspark
 # train serially on random shards
 def perceptronGDSerialShards(data, wInitial=None, nShards=1, nIter=10, stopCriteria=0.001,
                  learningRate=0.05, regParam=0.01):
     feature_size = len(data.take(1)[0][0])
     if wInitial is None:
         w = np.random.normal(size=feature size)
     else:
         w = wInitial
     for shared data in data.randomSplit([1.0/nShards]*nShards): # split data into shards
         for i in range(nIter):
             wb = sc.broadcast(w) # broadcast weight
             wdelta = learningRate*perceptronSubGradientCalc(shared_data, wb, regParam, feature_
             if sum(abs(wdelta))<=stopCriteria*sum(abs(w)):</pre>
                 break
             w = w - wdelta
     return w
 w = perceptronGDSerialShards(train data, nShards=4, nIter=10)
 print "Train serially on random shards"
 print "learned weight: %s" % str(w)
 print "Accuracy: %s" % perceptronAccuracy(test data, sc.broadcast(w))
Train serially on random shards
learned weight: [ 0.01615768  0.14727661 -0.00092179]
Accuracy: 0.87
Took 19 seconds.
```

```
%pyspark
# non-iterative parameter mixing
def percetronTrainLocal(data, w, regParam, feature_size, learningRate, nIter):
    # train with local data instead of RDD
    weight = np.array(w.value)
    for i in range(nIter):
        gradient = sum(map(lambda p: -p[1]*np.array(p[0]) if (p[1]*np.dot(weight,p[0]))<0 else
        wreg = weight*1 # L2 reg
        wreg[-1] = 0 # don't count the bias term in reg
        weight -= learningRate*(gradient/len(data) + regParam*wreg)
    return weight
def perceptronGDSerialParaMix(data, wInitial=None, nShards=1, nIter=10, learningRate=0.05, regf
    feature size = len(data.take(1)[0][0])
    if wInitial is None:
        w = np.random.normal(size=feature_size)
    else:
        w = wInitial
    wb = sc.broadcast(w)
    model_data = sc.parallelize([d.collect() for d in data.randomSplit([1.0/nShards]*nShards)])
    learned w = model data.map(lambda data: percetronTrainLocal(data, wb, regParam, feature siz
    return sum(learned_w)/len(learned_w) # take uniform average of the learned weights
```

```
w = perceptronGDSerialParaMix(train data, nShards=4, nIter=50)
 print "non-iterative parameter mixing"
 print "learned weight: %s" % str(w)
 print "Accuracy: %s" % perceptronAccuracy(test_data, sc.broadcast(w))
non-iterative parameter mixing
learned weight: [ 0.64234314  0.12919467  0.64654108]
Accuracy: 0.43
Took 8 seconds.
```

```
%pyspark
 # Iterative Param Mixing
 def perceptronGDSerialIterParaMix(data, wInitial=None, nShards=1, nIter=10, learningRate=0.05,
     feature_size = len(data.take(1)[0][0])
     if wInitial is None:
         w = np.random.normal(size=feature_size)
     else:
         w = wInitial
     model_data = sc.parallelize([d.collect() for d in data.randomSplit([1.0/nShards]*nShards)])
     for i in range(nIter):
         wb = sc.broadcast(w)
         # train only 1 epoch and then do param mixing
         learned_w = model_data.map(lambda data: percetronTrainLocal(data, wb, regParam, feature
         w = sum(learned_w)/len(learned_w) # take uniform average of the learned weights
     return w
 w = perceptronGDSerialIterParaMix(train data, nShards=4, nIter=50)
 print "Iterative Param Mixing"
 print "learned weight: %s" % str(w)
 print "Accuracy: %s" % perceptronAccuracy(test data, sc.broadcast(w))
Iterative Param Mixing
learned weight: [-0.72533034 0.15420804 0.05242582]
Accuracy: 0.91
Took 13 seconds.
```

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