return (x[0], val)

```
iteratorVal = {0:2, 1:2} #number of columns of second matrix, number of rows for first
db = [(0, ((0, 1), -1)), (0, ((0, 2), 2)), (0, ((1, 0), 4)), (0, ((1, 1), 11)), (0, ((1, 2), 2)), (1, ((0, 0), 3)),
(1, ((0, 1), -1)), (1, ((1, 0), 1)), (1, ((1, 1), 2)), (1, ((2, 0), 6)), (1, ((2, 1), 1))] # Example matrix
matrix = sc.parallelize(db) # Can also read from the disk/ db/ table.
# Checking for matrix 0/1 and creating (i, k) intermediate key value pairs accordingly
matrix = matrix.flatMap(lambda x: [((x[1][0][0], iter), (x[0], x[1][0][1], x[1][1])) if (x[0]==0) else
((iter, x[1][0][1]), (x[0], x[1][0][0], x[1][1])) for iter in range(iteratorVal[x[0]]))
matrix = matrix.groupByKey()
matrix = matrix.map(reducer).filter(lambda x: x[1]!=0).collect() #Filter to produce a sparse
matrix at the end
# Technically, GroupByKey + Map = Reduce operation. Hence the single pass is not violated!
def reducer(x):
       x1 = list(x[1])
       m1, m2 = [], []
       for i in x1:
               if(list(i)[0] == 0):
                       m1.append(list(i))
               else:
                       m2.append(list(i))
       m1 = sorted(m1, key=lambda x: x[1])
       m2 = sorted(m2, key=lambda x: x[1])
       maxIndex = max(m1[-1][1], m2[-1][1])
       for i in range(maxIndex):
       # Handling sparse matrix resulting in different intermediate key value pairs.
               if(m1[i][1] != i):
                       m1.insert(i, None)
               if(m2[i][1]!=i):
                       m2.insert(i, None)
       val = 0
       for i in range(maxIndex+1):
               if(m1[i] == None or m2[i] == None ):
                       val = val
               elif(m1[i][1] == i and m2[i][1] == i):
                                                     #Multiplying the j-th value alone
                      val += (m1[i][2]*m2[i][2])
```

```
#Certain variables like input features are assumed to be defined already.
import tensorflow as tf
X = tf.constant(features, dtype=tf.float32, name="X")
y = tf.constant(y.reshape(-1,1), dtype=tf.float32, name="y")
n = 10
learning rate = 0.0001
penalty = tf.constant(1.0, dtype=tf.float32, name="penalty")
beta = tf.Variable(tf.random_uniform([features.shape[1], 1], -1., 1.), name = "beta")
m = tf.Variable(tf.zeros([features.shape[1], 1]), name="momentum_weights")
momentum = 0.9
y_pred = tf.matmul(X, beta, name="predictions")
penalizedCost = tf.reduce_sum(tf.square(y - y_pred)) + penalty * tf.reduce_sum(tf.square(beta))
grads = tf.gradients(penalizedCost, [beta])[0]
momentum_op = tf.assign(m, momentum*m + learning_rate*grads)
training_op = tf.assign(beta, beta-momentum_op)
init = tf.global variables initializer()
with tf.Session() as sess:
sess.run(init)
for epoch in range(n epochs):
 if epoch %1 == 0: #print debugging output
 print("Epoch", epoch, "; penalizedCost =", int(penalizedCost.eval()))
 sess.run(training_op)
best beta = beta.eval()
                                     RESULT ANALYSIS
Momentum: 0.9
('Epoch', 0, '; penalizedCost =', 248912)
('Epoch', 1, '; penalizedCost =', 208998)
('Epoch', 2, '; penalizedCost =', 146552)
('Epoch', 3, '; penalizedCost =', 83685)
('Epoch', 4, '; penalizedCost =', 38788)
('Epoch', 5, '; penalizedCost =', 20954)
('Epoch', 6, '; penalizedCost =', 28629)
('Epoch', 7, '; penalizedCost =', 52131)
('Epoch', 8, '; penalizedCost =', 78465)
('Epoch', 9, '; penalizedCost =', 96344)
('Mean Absolute Error:', 13.02788132525499)
('Pearson Correlation:', (0.6373451939992985, 4.927043496055146e-07))
```

Momentum: 0.0 ('Epoch', 0, '; penalizedCost =', 289662) ('Epoch', 1, '; penalizedCost =', 242613) ('Epoch', 2, '; penalizedCost =', 203794) ('Epoch', 3, '; penalizedCost =', 171763) ('Epoch', 4, '; penalizedCost =', 145335) ('Epoch', 5, '; penalizedCost =', 123528) ('Epoch', 6, '; penalizedCost =', 105535) ('Epoch', 7, '; penalizedCost =', 90689) ('Epoch', 8, '; penalizedCost =', 78439) ('Epoch', 9, '; penalizedCost =', 68331) ('Mean Absolute Error:', 9.376570549976392) ('Pearson Correlation:', (0.6373451939992985, 4.927043496055146e-07))

We can clearly see that the momentum helps in stable reduction of loss and better performance sooner.

Link to code

A. If the Jaccard similarity between two sets (say A, B) = 0, then there is no element that lies in common between them.

This means that no element (∈ union(A, B)) would have result in a tuple of [1, 1] in the characteristic table together.

This implies no hash function can produce the same value [of first row with a 1] for both these sets. Hence the minhashing estimate would also be 0.

```
B.
import numpy as np
import random
rdd = sc.textFile(hdfs://*.json)
                                   #Reading all the ison files
rdd = rdd.map(json.loads)
#To club all the elements across different partition belonging to the same set
rdd = rdd.reduceByKey( lambda x,y: x.union(y))
# Op: {'set Name 1': set(element1, element2, ...), 'set Name 2': set(element1, element2, ...)
...}
#Collects all the elements from all the sets
rows = rdd.reduce( lambda x, y: x.union(y))
numSets = 0
                     # Can also be computed as count of keys
def getCharacteristicMatrix(x):
# Returns the characteristic Matrix for a given set
       numSets += 1
       row = list(rows[0])
                            # converting set to list
       elements = list(x[1]) # converting set to list
       element flag = []
       for i in row:
              element_flag.append(elements.count(i)>0)
       return (x[0], np.array(element_flag))
charMatrix = rdd.map(getCharacteristicMatrix).reduce(lambda x,y:
np.concatenate((x.reshape(-1,1), y.reshape(-1,1)), axis=1))
#Turns it to (num_elements, numsets): Characteristic matrix. Easier to fetch a row now.
hashes = [getHfunc(randint(1,10000)) for i in range(500)] #500 hash functions, r seeded
```

```
signatureMat = np.zeros(size=(500, numSets))
                                                   # Signature Matrix init
signatureMat[:, :] = np.inf
def minHashing(x):
       rowNum = getRowNum(charMatrix, x)
                                                   #Function that searches the row num
       row = list(rows[0])
       h = [hashes[i](row[rowNum]) for i in range(500)]
                                                          #Hash results for that row
       for s in range(numSets):
              If x[s] == 1:
                      for i in range(500):
                             if (h[i] < signatureMat[i][s]):</pre>
                                    signatureMat[i][s] = h[i]
                             else:
                                    pass
#foreach on 2d np matrix reads line by line
```

charMatrix = charMatrix.foreach(minHashing)

print (signatureMat)

A. Band size r=5; implies b = 100, jaccard = 0.75Probability that set1 and set2 matches in 1 band: $(0.75)^5 = 0.237$ Probability that set1 and set2 didn't match in 1 band: $1 - (0.75)^5 = 0.763$

Probability that set1 and set2 matches in at least 1 band

- = 1 Probability that set1 and set2 matched in no band
- $= 1 (0.763)^{100} = 0.9999999999982817 \sim 0.999$
- B. Probability that set1 and set2 matches in a band = s^r
 Probability that set1 and set2 don't match in a band = 1 s^r

Probability that they match in at least 1 band = $1 - (1 - s^r)^b$

$$1 - (1 - s^r)^b >= 0.99$$

$$0.01 >= (1 - s^r)^b$$

$$s = 0.9$$

$$0.01 >= (1 - 0.9^{r})^{b}$$

 $log(0.01) >= b log(1 - 0.9^r)$

b * r = n [Taking n to be 500 from q3(b)]

$$b = 27.9$$
, $r = 17.8$

Since they need to be whole numbers (27, 18), since higher b would imply higher false positive rate.

C. Since some false positive rate is allowed to be a bit more than the usual implementation, I would follow the following approach.

I would pick first 'p' rows from the characteristic matrix [Having p smaller than total number of rows, for large speedup]

And I would build the minHashing on this, so that it has fewer comparisons to make. If both sets have a minhash for a row it is dealt the normal way based on the equality/inequality.

If a set doesn't have its minHash value within the first p rows and the other set has, then it is counted as unequal minhash.

If both the sets being compared don't have a minHash from the first p rows, then it is not considered a valid example.

Due to the last condition the denominator tends to decrease, which would increase the false positive rate. However the speedup would be large.