# **MIDS Machine Learning at Scale**

# **End of Term exam**

Week 15 Summer, 2016

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# Exam Schedule (All times are in California Time)

4:00 PM - 6:00 PM

# Instructions for exam

1. Please submit your solutions and notebook via the following form:

Submission Form (http://goo.gl/forms/ggNYfRXz0t)

- 1. Please acknowledge receipt of exam by sending a quick reply to the instructors
- 2. Review the submission form first to scope it out (it will take a 5-10 minutes to input your answers and other information into this form)
- 3. Please keep all your work and responses in ONE (1) notebook only (and submit via the submission form)
- 4. Please make sure that the NBViewer link for your Submission notebook works
- 5. Please do NOT discuss this exam with anyone (including your class mates) until after Monday, Midday, of week 16
- 6. This is an open book exam meaning you can consult webpages and textbooks, class notes, slides etc. but you can not consult each other or any other person/group. Please complete this exam by yourself within the time limit.
- 7. For markdown help in iPython Notebooks please see: <a href="https://sourceforge.net/p/ipython/discussion/markdown\_syntax">https://sourceforge.net/p/ipython/discussion/markdown\_syntax</a>

# Exam questions begins here

Assume your tasked with modeling a REGRESSION problem. How do you determine which variables may be important?

- 1. If your data has unknown structure, start with: Tree-based methods
- 2. If statistical measures of importance are needed, start with: linear models (think Generalized linear models (GLMs)
- 3. If statistical measures of importance are not needed, start with: Regression with shrinkage (e.g., LASSO, Elastic net)
- 4. If statistical measures of importance are not needed, use: Stepwise regression

Select the single most correct response from the following:

- (a) 1
- (b) 2
- (c) 3
- (d) 1, 2, 3, 4

Answer: (d)

# **ET:2**

Using one-hot-encoding, a categorical feature with four distinct values would be represented by how many features?

- (a) 1 feature
- (b) 2 features
- (c) 4 features
- (d) none of the above

Answer: (c) 4 features

In the following (and also referring to HW12:

http://nbviewer.jupyter.org/urls/dl.dropbox.com/s/1wb2rdqbet54y1h/MIDS-MLS-Project-Criteo-CTR.ipynb (http://nbviewer.jupyter.org/urls/dl.dropbox.com/s/1wb2rdqbet54y1h/MIDS-MLS-Project-Criteo-CTR.ipynb)) we have hashed the three sample points using numBuckets=4 and numBuckets=100. Complete the three statements below about these hashed features summarized in the following table using each answer once.

Name	Raw Features	4 Buckets	100 Buckets
sampleOne	[(0, 'mouse'), (1, 'black')]	{2: 1.0, 3: 1.0}	{14: 1.0, 31: 1.0}
sampleTwo	[(0, 'cat'), (1, 'tabby'), (2, 'mouse')]	{0: 2.0, 2: 1.0}	{40: 1.0, 16: 1.0, 62: 1.0}
sampleThree	[(0, 'bear'), (1, 'black'), (2, 'salmon')	{0: 1.0, 1: 1.0, 2: 1.0}	{72: 1.0, 5: 1.0, 14: 1.0}

With 100 buckets, sampleOne and sampleThree both contain index 14 due to \_\_\_.

- (a) A hash collision
- (b) Underlying properties of the data
- (c) The fact that used 100 buckets
- (d) none of the above

Answer: (b) underlying properties of the data

In the following (and also referring to HW12:

http://nbviewer.jupyter.org/urls/dl.dropbox.com/s/1wb2rdqbet54y1h/MIDS-MLS-Project-Criteo-CTR.ipynb (http://nbviewer.jupyter.org/urls/dl.dropbox.com/s/1wb2rdqbet54y1h/MIDS-MLS-Project-Criteo-CTR.ipynb)) we have hashed the three sample points using numBuckets=4 and numBuckets=100. Complete the three statements below about these hashed features summarized in the following table using each answer once.

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sampleThree	[(0, 'bear'), (1, 'black'), (2, 'salmon')	{0: 1.0, 1: 1.0, 2: 1.0}	{72: 1.0, 5: 1.0, 14: 1.0}

It is likely that sample Two has two indices with 4 buckets, but three indices with 100 buckets due to \_\_.

- (a) A hash collision
- (b) Underlying properties of the data
- (c) The fact that we go from 4 to 100 buckets
- (d) none of the above

**Answer**: (c) The fact that we go from 4 to 100 buckets.

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sampleThree	[(0, 'bear'), (1, 'black'), (2, 'salmon')	{0: 1.0, 1: 1.0, 2: 1.0}	{72: 1.0, 5: 1.0, 14: 1.0}

With 4 buckets, sampleTwo and sampleThree both contain index 0 due to \_\_\_.

- (a) A hash collision
- (b) Underlying properties of the data
- (c) The fact that we use 4 buckets
- (d) none of the above

Answer: (a) a hash collision

# ET:6 When applying numerical machine learning approaches (and for non-numerical approaches if required) to big data problems which of the following steps are could be used during modeling and are recommended:

- (a) Convert categorical features to numerical features via one-hot-encoding and store in a dense representation
- (b) Transform categorical features using hashing regardless of how many unique categorical values exist in training and test data
- (c) Use matrix factorization to remap your input vectors to latent concepts
- (d) none of the above

**Answer**: (a), (b), (c)

When dealing with numercial data which of the following are ways to deal with missing data:

- (a) Delete records that have missing input values
- (b) Standardize the data and set all missing values to 1 (one)
- (c) Use K-nearest neighbours based on the test set to fill in missing values in the training set
- (d) none of the above

**Answer**: (a) would be the method that most people would use; Some people would also use and (c).

# **ET:8**

In the Criteo project, we're trying to predict what:

- (a) Revenue from click events
- (b) Click-through vs not click event
- (c) Probability of a purchase
- (d) none of the above

Answer: (b) Click-through vs not click event

# **ET:9**

Which of the following are true about the purpose of a loss function?

- (a) It's a way to penalize a model for incorrect predictions
- (b) It precisely defines the optimization problem to be solved for a particular learning model
- (c) Loss functions can be used for modeling both classification and regression problems
- (d) none of the above

**Answer**: (a), (b), (c)

When implementing Logistic Regression with Regularization in Spark which of the following apply?

- (a) When lambda equals one, it provides the same result as standard logistic regression
- (b) One only needs to modify the standard logistic regression by modifying the Mapper
- (c) Can be framed as minimizing a convex function
- (d) none of the above

Answer: (b), (c)

# ET:11

In the context of ecommerce you have just deployed a new conversion rate prediction model to production. This model (aka treatment model) will challenge the control nodel (i.e., the current model) in AB Test manner to see if it can be produce better revenue. Here is the data that was taken from this live AB Test.

CONTROL MODEL	(our new CTR model)
Impression ID	Revenue
1	\$0.50
2	\$0.50
3	\$3.00
• • • • •	
20000	\$3.00
20001	\$3.00
20002	\$3.00
20003	\$3.00
• • • • •	
50,001	\$3.00
• • • • •	
100,000	\$4.00

All other impressions in this 100,000 sample resulted in zero transactions and therefore zero revenue.

```
TREATMENT MODEL (our new CTR model)
Impression ID Revenue

1 $1.50
2 $0.50
3 $0.00
.....
50,001 $3.00
.....
100,000 $4.00
```

All other impressions in this 100,000 sample resulted in zero transactions and therefore zero revenue.

P-values are a common way to determine the statistical significance of a test. The smaller it is, the more confident you can be that the test results are due to something other than random chance. A common p-value of .05 is a 5% significance level. Similarly, a p-value of .01 is a 1% significance level. A p-value of .20 is a 20% significance level. For this problem set the p-value to 0.01

Which of the following are true:

512994))

- (a) Based on revenue there is no statistical significant difference between the Control and the Treatment at p-value of 0.05 for a one-sided t-test
- (b) Based on transaction rates (tranactions that generated revenue versus not) there is no statistical significant difference between the Control and the Treatment at p-value of 0.05 for a one-sided t-test
- (c) AB testing using differences in revenue for this problem is a useful means of determining if the Treatment conversion rate prediction model is better than the control model.
- (d) none of the above

```
import numpy as np
In [1]:
        from scipy.sparse import coo matrix
        import scipy.stats
        control=[.5, .5, 3, 3, 3, 3, 3, 4]
        control.extend([0]*(100000-len(control)))
        control=np.asarray(control)
        treatment=[ 0, 0, 0, 0, 0, 0, 0, 0]
        treatment.extend([0]*(100000-len(treatment)))
        treatment=np.asarray(treatment)
        #nb 2-sided test
        scipy.stats.ttest ind(control, treatment), scipy.stats.ttest ind(contr
        ol.astype(bool), treatment.astype(bool))
Out[1]: (Ttest indResult(statistic=2.7393492169656311, pvalue=0.006156639881
        4161729),
         Ttest indResult(statistic=3.000120008400637, pvalue=0.0026990648570
```

Answer: (c)

### ET:12

Given this graph expressed in the form of an adjacency list,

```
Node
      adjacentNode:weightAssociatedWithEdge
N1
      N6:10, N2:2
N2
      N3:1
И3
      N4:1
N4
      N5:1
N5
      N6:1
      N7:1
N6
Ν7
      N8:1
N8
      N9:1
```

Using the parallel breadth-first search algorithm for determining the shortest path from a single source, how many iterations are required to discover the shortest distances to all nodes from Node 1

A 7

B 8

C 13

D None of the above

Answer: B 8

# ET:13 Assume the Lagrangian for SOFT SVMs (unconstrained optimization) is as follows:

```
minimize [\lambda/2 * w'w + C\Sigma i=1:n \xi i + 1/n \Sigma i=:n max(0, 1 - \xi i - y i(w'x i - b)))]
```

- (a) When  $\lambda$  is super small (e.g., 0.000001), then the above Langrangian will yield a Hard SVM
- (b) In the context of support vector machines, linear kernels can be readily parallelized in map reduce frameworks such as Spark
- (c) Sequential learning via algorithms such perceptron can take advantage of map-reduce frameworks and yield exactly the same results as a single core implementation with significant reductions in training time (d) When  $\lambda$  is 1.0, then the above Langrangian will yield a Soft SVM

**Answer**: (a), (b), (d)

# ET:14 Given the following paired RDDs

```
RDD1 = \{(1, 2), (3, 4), (3, 6)\} RDD2 = \{(3, 9), (3, 6)\}
```

B: [(3, (4, 9)), (3, (4, 6)), (3, (6, 9)), (3, (6, 6))]

A: [(3, (4, 9)), (3, (6, 9))]

Using PySpark, write code to perform an inner join of these paired RDDs. What is the resulting RDD? Make your Spark available in your notebook:

```
C: [(3, (4, 9)), (3, (4, 6)), (3, (6, 9)), (3, (6, 9))]
D: None of the above
  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 0x7f2916f25790>
           <pyspark.sql.context.SQLContext object at 0x7f28f513a210>
  In [3]: RDD1=sc.parallelize([(1, 2), (3, 4), (3, 6)])
           RDD2=sc.parallelize([(3, 9), (3, 6)])
           RDD1.join(RDD2).collect()
  Out[3]: [(3, (4, 9)), (3, (4, 6)), (3, (6, 9)), (3, (6, 6))]
```

Answer: B

# ET:15 You have been tasked to build a predictive model to forecast beer sales for a chain of stores.

After doing basic exploratory analysis on the data, what is the first thing you do regarding modeling?

- (a) Construct a baseline model
- (b) Determine a metric to evaluate your machine learnt models
- (c) Split your data into training, validation and test subsets (or split using cross fold validatation)
- (d) All of the of the above

Answer: (d) All of the above.

## ET:16

Use Spark and the following notebook to answer this question:

- <a href="http://nbviewer.jupyter.org/urls/dl.dropbox.com/s/6s5ph41h74bggwi/Linear-Regression-on-Beer-Data.ipynb">http://nbviewer.jupyter.org/urls/dl.dropbox.com/s/6s5ph41h74bggwi/Linear-Regression-on-Beer-Data.ipynb</a>)
- <a href="https://www.dropbox.com/s/6s5ph41h74bggwi/Linear-Regression-on-Beer-Data.ipynb?dl=0">https://www.dropbox.com/s/6s5ph41h74bggwi/Linear-Regression-on-Beer-Data.ipynb?dl=0</a> (<a href="https://www.dropbox.com/s/6s5ph41h74bggwi/Linear-Regression-on-Beer-Data.ipynb?dl=0">https://www.dropbox.com/s/6s5ph41h74bggwi/Linear-Regression-on-Beer-Data.ipynb?dl=0</a>

The mean absolute percentage error (MAPE), also known as mean absolute percentage deviation (MAPD), is a measure of prediction accuracy of a model for say a forecasting method in statistics, for example in trend estimation. It usually expresses accuracy as a percentage, and is defined by the formula:

MAPE = average over all examples (100\*Abs(Actual - Predicted) / Actual))

Note when Actual is zero that test row is dropped from the evaluation.

Construct a mean model for target variable CASES18PK. Calculate the MAPE for the mean model over the training set. Select the closest answer.

- (a) 200%
- (b) 250%
- (c) 20%
- (d) 180%

Answer: (d)

Use Spark and the following notebook to answer this question:

- <a href="http://nbviewer.jupyter.org/urls/dl.dropbox.com/s/6s5ph41h74bggwi/Linear-Regression-on-Beer-Data.ipynb">http://nbviewer.jupyter.org/urls/dl.dropbox.com/s/6s5ph41h74bggwi/Linear-Data.ipynb</a> Regression-on-Beer-Data.ipynb)
- <a href="https://www.dropbox.com/s/6s5ph41h74bggwi/Linear-Regression-on-Beer-Data.ipynb?dl=0">https://www.dropbox.com/s/6s5ph41h74bggwi/Linear-Regression-on-Beer-Data.ipynb?dl=0</a> (<a href="https://www.dropbox.com/s/6s5ph41h74bggwi/Linear-Regression-on-Beer-Data.ipynb?dl=0">https://www.dropbox.com/s/6s5ph41h74bggwi/Linear-Regression-on-Beer-Data.ipynb?dl=0</a>

The target variable CASES18PK is skewed, so take the log of it (and make it more normally distributed) and compute the MAPE of the mean model for CASES18PK. Select the closest answer to your calculated MAPE.

- (a) 200%
- (b) 30%
- (c) 20%
- (d) 10%

Answer: (b)

# ET:18

Use Spark and the following notebook to answer this question:

- http://nbviewer.jupyter.org/urls/dl.dropbox.com/s/6s5ph41h74bggwi/Linear-Regression-on-Beer-Data.ipynb (http://nbviewer.jupyter.org/urls/dl.dropbox.com/s/6s5ph41h74bggwi/Linear-Regression-on-Beer-Data.ipynb)
- <a href="https://www.dropbox.com/s/6s5ph41h74bggwi/Linear-Regression-on-Beer-Data.ipynb?dl=0">https://www.dropbox.com/s/6s5ph41h74bggwi/Linear-Regression-on-Beer-Data.ipynb?dl=0</a> (<a href="https://www.dropbox.com/s/6s5ph41h74bggwi/Linear-Regression-on-Beer-Data.ipynb?dl=0">https://www.dropbox.com/s/6s5ph41h74bggwi/Linear-Regression-on-Beer-Data.ipynb?dl=0</a>

Build a linear regression model using the following variables:

Log(CASES18PK) ~ log(PRICE12PK), log(PRICE18PK), log(PRICE30PK)

Calculate MAPE over the test set and select the closest answer.

- (a) 4.3%
- (b) 4.6%
- (c) 3.5%
- (d) 3.9%

Recall that Spark automatically sends all variables referenced in your closures to the worker nodes. While this is convenient, it can also be inefficient because (1) the default task launching mechanism is optimized for small task sizes, and (2) you might, in fact, use the same variable in multiple parallel operations, but Spark will send it separately for each operation. As an example, say that we wanted to write a Spark program that looks up countries by their call signs (e.g., the call sign for Ireland is EJZ) by prefix matching in an table. In the following the "signPrefixes" variable is essentially a table with two columns "Sign" and "Country Name". The goal is to join the following tables:

signPrefixes table with columns "Sign" and "Country Name"
contactCounts table with columns "Sign" and "count"

to yield a new table:

countryContactCounts with the following columns "Country Name" and "count"

Use Spark and the following notebook to answer this question:

- <a href="http://nbviewer.jupyter.org/urls/dl.dropbox.com/s/6s5ph41h74bggwi/Linear-Regression-on-Beer-Data.ipynb">http://nbviewer.jupyter.org/urls/dl.dropbox.com/s/6s5ph41h74bggwi/Linear-Data.ipynb</a> (http://nbviewer.jupyter.org/urls/dl.dropbox.com/s/6s5ph41h74bggwi/Linear-Regression-on-Beer-Data.ipynb)
- <a href="https://www.dropbox.com/s/6s5ph41h74bggwi/Linear-Regression-on-Beer-Data.ipynb?dl=0">https://www.dropbox.com/s/6s5ph41h74bggwi/Linear-Regression-on-Beer-Data.ipynb?dl=0</a> (<a href="https://www.dropbox.com/s/6s5ph41h74bggwi/Linear-Regression-on-Beer-Data.ipynb?dl=0">https://www.dropbox.com/s/6s5ph41h74bggwi/Linear-Regression-on-Beer-Data.ipynb?dl=0</a>

How can we modfify this code to make it more efficient? Choose one response only

- (a) modify line 18 with sc.broadcast(loadCallSignTable())
- (b) Use accumulators to store the counts for each country
- (c) The code is already optimal
- (d) none of the above

Answer:			
(a)			
In [ ]:			