DATASCI W261: Machine Learning at Scale

W261-1 Fall 2015 Week 12: Criteo CTR Project November 14, 2015

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Click-Through Rate Prediction Lab

This lab covers the steps for creating a click-through rate (CTR) prediction pipeline. You will work with the <u>Criteo Labs (http://labs.criteo.com/)</u> dataset that was used for a recent <u>Kaggle competition (https://www.kaggle.com/c/criteo-display-ad-challenge)</u>.

This lab will cover:

- ####Part 1: Featurize categorical data using one-hot-encoding (OHE)
- ####Part 2: Construct an OHE dictionary
- ####Part 3: Parse CTR data and generate OHE features
 - Visualization 1: Feature frequency
- ####Part 4: CTR prediction and logloss evaluation
 - Visualization 2: ROC curve
- ####Part 5: Reduce feature dimension via feature hashing
 - Visualization 3: Hyperparameter heat map

Note that, for reference, you can look up the details of the relevant Spark methods in <u>Spark's Python API (https://spark.apache.org/docs/latest/api/python/pyspark.html#pyspark.RDD)</u> and the relevant NumPy methods in the <u>NumPy Reference</u> (http://docs.scipy.org/doc/numpy/reference/index.html)

```
In [82]: import os
    import sys

spark_home = os.environ['SPARK_HOME']
    if not spark_home:
        raise ValueError('SPARK_HOME enviroment variable is not set')
    sys.path.insert(0,os.path.join(spark_home,'python'))
    sys.path.insert(0,os.path.join(spark_home,'python/lib/py4j-0.9-src.zip'))
    from pyspark import SparkContext, SparkConf,SparkFiles

    try:
        sc.stop()
    except:
        pass
    conf = SparkConf().setAppName("my app").setMaster("spark://master:7077")
    sc = SparkContext(conf=conf)
```

```
In [83]: labVersion = 'MIDS_MLS_week12_v_0_9'
```

Part 1: Featurize categorical data using one-hot-encoding

(1a) One-hot-encoding

We would like to develop code to convert categorical features to numerical ones, and to build intuition, we will work with a sample unlabeled dataset with three data points, with each data point representing an animal. The first feature indicates the type of animal (bear, cat, mouse); the second feature describes the animal's color (black, tabby); and the third (optional) feature describes what the animal eats (mouse, salmon).

In a one-hot-encoding (OHE) scheme, we want to represent each tuple of (featureID, category) via its own binary feature. We can do this in Python by creating a dictionary that maps each tuple to a distinct integer, where the integer corresponds to a binary feature. To start, manually enter the entries in the OHE dictionary associated with the sample dataset by mapping the tuples to consecutive integers starting from zero, ordering the tuples first by featureID and next by category.

Later in this lab, we'll use OHE dictionaries to transform data points into compact lists of features that can be used in machine learning algorithms.

```
In [84]: # Data for manual OHE
    # Note: the first data point does not include any value for the optional third fea
    sampleOne = [(0, 'mouse'), (1, 'black')]
    sampleTwo = [(0, 'cat'), (1, 'tabby'), (2, 'mouse')]
    sampleThree = [(0, 'bear'), (1, 'black'), (2, 'salmon')]
    sampleDataRDD = sc.parallelize([sampleOne, sampleTwo, sampleThree])
```

```
In [85]: # TODO: Replace <FILL IN> with appropriate code
    sampleOHEDictManual = {}
    sampleOHEDictManual[(0,'bear')] = 0
    sampleOHEDictManual[(0,'cat')] = 1
    sampleOHEDictManual[(0,'mouse')] = 2
    sampleOHEDictManual[(1, 'black')] = 3
    sampleOHEDictManual[(1, 'tabby')] = 4
    sampleOHEDictManual[(2, 'mouse')] = 5
    sampleOHEDictManual[(2, 'salmon')] = 6
```

```
In [86]: # A testing helper
         #https://pypi.python.org/pypi/test_helper/0.2
         import hashlib
         class TestFailure(Exception):
         class PrivateTestFailure(Exception):
         class Test(object):
           passed = 0
           numTests = 0
           failFast = False
           private = False
           @classmethod
           def setFailFast(cls):
             cls.failFast = True
           @classmethod
           def setPrivateMode(cls):
             cls.private = True
           @{\it classmethod}
           def assertTrue(cls, result, msg=""):
             cls.numTests += 1
             if result == True:
               cls.passed += 1
               print "1 test passed."
                print "1 test failed. " + msg
                if cls.failFast:
                  if cls.private:
                    raise PrivateTestFailure(msg)
                  else:
                    raise TestFailure(msg)
           @classmethod
           def assertEquals(cls, var, val, msg=""):
             cls.assertTrue(var == val, msg)
           @classmethod
           def assertEqualsHashed(cls, var, hashed_val, msg=""):
             cls.assertEquals(cls._hash(var), hashed_val, msg)
           @classmethod
           def printStats(cls):
             print "{0} / {1} test(s) passed.".format(cls.passed, cls.numTests)
           @classmethod
           def _hash(cls, x):
             return hashlib.sha1(str(x)).hexdigest()
```

```
In [87]:
         # TEST One-hot-encoding (1a)
         from test helper import Test
         Test.assertEqualsHashed(sampleOHEDictManual[(0, 'bear')],
                                  'b6589fc6ab0dc82cf12099d1c2d40ab994e8410c',
                                  "incorrect value for sampleOHEDictManual[(0,'bear')]")
         Test.assertEqualsHashed(sampleOHEDictManual[(0,'cat')],
                                  '356a192b7913b04c54574d18c28d46e6395428ab',
                                  "incorrect value for sampleOHEDictManual[(0,'cat')]")
         Test.assertEqualsHashed(sampleOHEDictManual[(0, 'mouse')],
                                  'da4b9237bacccdf19c0760cab7aec4a8359010b0',
                                  "incorrect value for sampleOHEDictManual[(0,'mouse')]")
         Test.assertEqualsHashed(sampleOHEDictManual[(1,'black')],
                                  '77de68daecd823babbb58edb1c8e14d7106e83bb',
                                  "incorrect value for sampleOHEDictManual[(1,'black')]")
         Test.assertEqualsHashed(sampleOHEDictManual[(1, 'tabby')],
                                  '1b6453892473a467d07372d45eb05abc2031647a',
                                  "incorrect value for sampleOHEDictManual[(1,'tabby')]")
         Test.assertEqualsHashed(sampleOHEDictManual[(2, 'mouse')],
                                  'ac3478d69a3c81fa62e60f5c3696165a4e5e6ac4'.
                                  "incorrect value for sampleOHEDictManual[(2,'mouse')]")
         Test.assertEqualsHashed(sampleOHEDictManual[(2, 'salmon')],
                                  'c1dfd96eea8cc2b62785275bca38ac261256e278',
                                  "incorrect value for sampleOHEDictManual[(2,'salmon')]")
         Test.assertEquals(len(sampleOHEDictManual.keys()), 7,
                            'incorrect number of keys in sampleOHEDictManual')
```

```
1 test passed.
1 test passed.
1 test passed.
```

1 test passed.

(1b) Sparse vectors

Data points can typically be represented with a small number of non-zero OHE features relative total number of features that occur in the dataset. By leveraging this sparsity and using sparse representations of OHE data, we can reduce storage and computational burdens. Below are a fe sample vectors represented as dense numpy arrays. Use SparseVector (https://spark.apache.org/docs/latest/api/python/pyspark.mllib.html#pyspark.mllib.linalg.Sparse to represent them in a sparse fashion, and verify that both the sparse and dense representation

the same results when computing dot products (http://en.wikipedia.org/wiki/Dot_product) (we v use MLlib to train classifiers via gradient descent, and MLlib will need to compute dot products between SparseVectors and dense parameter vectors).

Use SparseVector(size, *args) to create a new sparse vector where size is the length of the and args is either a dictionary, a list of (index, value) pairs, or two separate arrays of indices an values (sorted by index). You'll need to create a sparse vector representation of each dense vec aDense and bDense.

```
In [88]: import numpy as np
from pyspark.mllib.linalg import SparseVector
```

```
In [89]: # TODO: Replace <FILL IN> with appropriate code
    aDense = np.array([0., 3., 0., 4.])
    aSparse = SparseVector(4, [(1,3), (3,4)])

bDense = np.array([0., 0., 0., 1.])
    bSparse = SparseVector(4, [(3, 1)])

w = np.array([0.4, 3.1, -1.4, -.5])
    print aDense.dot(w)
    print bDense.dot(w)
    print bSparse.dot(w)
```

- 7.3
- 7.3
- -0.5
- -0.5

In [90]: # TEST Sparse Vectors (1b)

Test.assertTrue(isinstance(aSparse, SparseVector), 'aSparse needs to be an instanc
Test.assertTrue(isinstance(bSparse, SparseVector), 'aSparse needs to be an instanc
Test.assertTrue(aDense.dot(w) == aSparse.dot(w),

'dot product of aDense and w should equal dot product of aSparse a
Test.assertTrue(bDense.dot(w) == bSparse.dot(w),

'dot product of bDense and w should equal dot product of bSparse a

- 1 test passed.
- 1 test passed.
- 1 test passed.
- 1 test passed.

(1c) OHE features as sparse vectors

Now let's see how we can represent the OHE features for points in our sample dataset. Using the mapping defined by the OHE dictionary from Part (1a), manually define OHE features for the three sample data points using SparseVector format. Any feature that occurs in a point should have the value 1.0. For example, the DenseVector for a point with features 2 and 4 would be [0.0, 0.0, 1.0, 0.0, 1.0, 0.0, 0.0].

```
In [91]: # Reminder of the sample features
# sampleOne = [(0, 'mouse'), (1, 'black')]
# sampleTwo = [(0, 'cat'), (1, 'tabby'), (2, 'mouse')]
# sampleThree = [(0, 'bear'), (1, 'black'), (2, 'salmon')]
```

```
In [92]: # TODO: Replace <FILL IN> with appropriate code
    sampleOneOHEFeatManual = SparseVector(7,[(2,1),(3,1)])
    sampleTwoOHEFeatManual = SparseVector(7,[(1,1),(4,1),(5,1)])
    sampleThreeOHEFeatManual = SparseVector(7,[(0,1),(3,1),(6,1)])
```

```
In [93]:
         # TEST OHE Features as sparse vectors (1c)
         Test.assertTrue(isinstance(sampleOneOHEFeatManual, SparseVector),
                          'sampleOneOHEFeatManual needs to be a SparseVector')
         Test.assertTrue(isinstance(sampleTwoOHEFeatManual, SparseVector),
                          sampleTwoOHEFeatManual needs to be a SparseVector')
         Test.assertTrue(isinstance(sampleThreeOHEFeatManual, SparseVector),
                          'sampleThreeOHEFeatManual needs to be a SparseVector')
         Test.assertEqualsHashed(sampleOneOHEFeatManual.
                                  'ecc00223d141b7bd0913d52377cee2cf5783abd6',
                                  'incorrect value for sampleOneOHEFeatManual')
         Test.assertEqualsHashed(sampleTwoOHEFeatManual,
                                  '26b023f4109e3b8ab32241938e2e9b9e9d62720a',
                                  'incorrect value for sampleTwoOHEFeatManual')
         Test.assertEqualsHashed(sampleThreeOHEFeatManual,
                                  'c04134fd603ae115395b29dcabe9d0c66fbdc8a7',
                                  'incorrect value for sampleThreeOHEFeatManual')
```

```
1 test passed.
```

(1d) Define a OHE function

Next we will use the OHE dictionary from Part (1a) to programatically generate OHE features from the original categorical data. First write a function called oneHotEncoding that creates OHE feature vectors in SparseVector format. Then use this function to create OHE features for the first sample data point and verify that the result matches the result from Part (1c).

¹ test passed.

```
MIDS-w261-HW12-Lane
In [94]:
         # TODO: Replace <FILL IN> with appropriate code
         def oneHotEncoding(rawFeats, OHEDict, numOHEFeats):
             """Produce a one-hot-encoding from a list of features and an OHE dictionary.
             Note:
                 You should ensure that the indices used to create a SparseVector are sorte
             Args:
                 rawFeats (list of (int, str)): The features corresponding to a single obse
                     feature consists of a tuple of featureID and the feature's value. (e.g.
                 OHEDict (dict): A mapping of (featureID, value) to unique integer.
                 numOHEFeats (int): The total number of unique OHE features (combinations o
                     value).
             Returns:
                 SparseVector: A SparseVector of length numOHEFeats with indicies equal to
                     identifiers for the (featureID, value) combinations that occur in the
                     with values equal to 1.0.
             return SparseVector(numOHEFeats, [(OHEDict[(featureID, value)],1) for (feature
         # Calculate the number of features in sampleOHEDictManual
         numSampleOHEFeats = len(sampleOHEDictManual.keys())
         # Run oneHotEnoding on sampleOne
         sampleOneOHEFeat =oneHotEncoding(sampleOne, sampleOHEDictManual, numSampleOHEFeats
         print sampleOneOHEFeat
         (7,[2,3],[1.0,1.0])
In [95]: # TEST Define an OHE Function (1d)
         Test.assertTrue(sampleOneOHEFeat == sampleOneOHEFeatManual,
```

```
'sampleOneOHEFeat should equal sampleOneOHEFeatManual')
Test.assertEquals(sampleOneOHEFeat, SparseVector(7, [2,3], [1.0,1.0]),
                  'incorrect value for sampleOneOHEFeat')
Test.assertEquals(oneHotEncoding([(1, 'black'), (0, 'mouse')], sampleOHEDictManual
                                 numSampleOHEFeats), SparseVector(7, [2,3], [1.0,1]
                  'incorrect definition for oneHotEncoding')
```

```
1 test passed.
```

1 test passed.

1 test passed.

(1e) Apply OHE to a dataset

Finally, use the function from Part (1d) to create OHE features for all 3 data points in the sample dataset.

```
# TODO: Replace <FILL IN> with appropriate code
In [96]:
         sampleOHEData = sampleDataRDD.map(lambda v: oneHotEncoding(v, sampleOHEDictManual,
         print sampleOHEData.collect()
```

```
[SparseVector(7, {2: 1.0, 3: 1.0}), SparseVector(7, {1: 1.0, 4: 1.0, 5: 1.0}),
SparseVector(7, {0: 1.0, 3: 1.0, 6: 1.0})]
```

```
1 test passed.
1 test passed.
1 test passed.
```

1 test passed.

1 test passed.

Part 2: Construct an OHE dictionary

```
(2a) Pair RDD of (featureID, category)
```

To start, create an RDD of distinct (featureID, category) tuples. In our sample dataset, the 7 items in the resulting RDD are (0, 'bear'), (0, 'cat'), (0, 'mouse'), (1, 'black'), (1, 'tabby'), (2, 'mouse'), (2, 'salmon'). Notably 'black' appears twice in the dataset but only contributes one item to the RDD: (1, 'black'), while 'mouse' also appears twice and contributes two items: (0, 'mouse') and (2, 'mouse'). Use flatMap (https://spark.apache.org/docs/latest/api/python/pyspark.html#pyspark.RDD.flatMap) and distinct

(https://spark.apache.org/docs/latest/api/python/pyspark.html#pyspark.RDD.distinct).

1 test passed.

(2b) OHE Dictionary from distinct features

Next, create an RDD of key-value tuples, where each (featureID, category) tuple in sampleDistinctFeats is a key and the values are distinct integers ranging from 0 to (number of keys - 1). Then convert this RDD into a dictionary, which can be done using the collectAsMap action. Note that there is no unique mapping from keys to values, as all we require is that each (featureID, category) key be mapped to a unique integer between 0 and the number of keys. this exercise, any valid mapping is acceptable. Use zipWithIndex

(https://spark.apache.org/docs/latest/api/python/pyspark.html#pyspark.RDD.zipWithIndex) followed by collectAsMap

(https://spark.apache.org/docs/latest/api/python/pyspark.html#pyspark.RDD.collectAsMap).

```
In our sample dataset, one valid list of key-value tuples is: [((0, 'bear'), 0), ((2, 'salmon 1), ((1, 'tabby'), 2), ((2, 'mouse'), 3), ((0, 'mouse'), 4), ((0, 'cat'), 5), ((1,
```

'black'), 6)]. The dictionary defined in Part (1a) illustrates another valid mapping between ke and integers.

(2c) Automated creation of an OHE dictionary

y'): 1, (1, 'black'): 5, (0, 'mouse'): 2}

Now use the code from Parts (2a) and (2b) to write a function that takes an input dataset and outputs an OHE dictionary. Then use this function to create an OHE dictionary for the sample dataset, and verify that it matches the dictionary from Part (2b).

```
In [102]: # TODO: Replace <FILL IN> with appropriate code
def createOneHotDict(inputData):
    """Creates a one-hot-encoder dictionary based on the input data.

Args:
    inputData (RDD of lists of (int, str)): An RDD of observations where each
    made up of a list of (featureID, value) tuples.

Returns:
    dict: A dictionary where the keys are (featureID, value) tuples and map to
    unique integers.
    """
    return inputData.flatMap(lambda v: v).distinct().zipWithIndex().collectAsMap()

sampleOHEDictAuto = createOneHotDict(sampleDataRDD)
print sampleOHEDictAuto

{(2, 'mouse'): 3, (0, 'cat'): 6, (0, 'bear'): 0, (2, 'salmon'): 4, (1, 'tabb)
```

- 1 test passed.
 1 test passed.
- Part 3: Parse CTR data and generate OHE features

Before we can proceed, you'll first need to obtain the data from Criteo. If you have already completed this step in the setup lab, just run the cells below and the data will be loaded into the rawData variable.

Below is Criteo's data sharing agreement. After you accept the agreement, you can obtain the download URL by right-clicking on the "Download Sample" button and clicking "Copy link address" or "Copy Link Location", depending on your browser. Paste the URL into the # TODO cell below. The file is 8.4 MB compressed. The script below will download the file to the virtual machine (VM) and then extract the data.

If running the cell below does not render a webpage, open the <u>Criteo agreement</u> (http://labs.criteo.com/downloads/2014-kaggle-display-advertising-challenge-dataset/) in a separate browser tab. After you accept the agreement, you can obtain the download URL by right-clicking on the "Download Sample" button and clicking "Copy link address" or "Copy Link Location", depending on your browser. Paste the URL into the # TODO cell below.

Note that the download could take a few minutes, depending upon your connection speed.

The Criteo CTR data is for HW12.1 is available here (24.3 Meg, 100,000 Rows):

```
https://www.dropbox.com/s/m4jlnv6rdbqzzhu/dac sample.txt?dl=0
```

Alternatively you can download the sample data directly by following the instructions contained in the cell below (8M compressed).

In [104]

Run this code to view Criteo's agreement
from IPython.lib.display import IFrame

IFrame("http://labs.criteo.com/downloads/2014-kaggle-display-advertising-challenge
600, 350)

Out[104]:





Download Kaggle Display Advertising Challenge Dataset

CRITEO LABS DATA TERMS OF USE

In [105]:	

```
# TODO: Replace <FILL IN> with appropriate code
# Just replace <FILL IN> with the url for dac sample.tar.gz
import glob
import os.path
import tarfile
import urllib
import urlparse
# Paste url, url should end with: dac_sample.tar.gz
url = 'http://labs.criteo.com/wp-content/uploads/2015/04/dac sample.tar.gz'
url = url.strip()
baseDir = os.path.join('data')
inputPath = os.path.join('w261', 'dac_sample.txt')
fileName = os.path.join(baseDir, inputPath)
inputDir = os.path.split(fileName)[0]
def extractTar(check = False):
   # Find the zipped archive and extract the dataset
   tars = glob.glob('dac sample*.tar.gz*')
   if check and len(tars) == 0:
      return False
   if len(tars) > 0:
       try:
            tarFile = tarfile.open(tars[0])
        except tarfile.ReadError:
            if not check:
                print 'Unable to open tar.gz file. Check your URL.'
            return False
        tarFile.extract('dac sample.txt', path=inputDir)
        print 'Successfully extracted: dac sample.txt'
        return True
   else:
        print 'You need to retry the download with the correct url.'
        print ('Alternatively, you can upload the dac sample.tar.gz file to your J
              'directory')
        return False
if os.path.isfile(fileName):
   print 'File is already available. Nothing to do.'
elif extractTar(check = True):
   print 'tar.gz file was already available.'
elif not url.endswith('dac sample.tar.gz'):
   print 'Check your download url. Are you downloading the Sample dataset?'
else:
   # Download the file and store it in the same directory as this notebook
        urllib.urlretrieve(url, os.path.basename(urlparse.urlsplit(url).path))
   except IOError:
        print 'Unable to download and store: {0}'.format(url)
   extractTar()
```

File is already available. Nothing to do.

[u'0,1,1,5,0,1382,4,15,2,181,1,2,,2,68fd1e64,80e26c9b,fb936136,7b4723c4,25c83c9 8,7e0cccf,de7995b8,1f89b562,a73ee510,a8cd5504,b2cb9c98,37c9c164,2824a5f6,1adce 6ef,8ba8b39a,891b62e7,e5ba7672,f54016b9,21ddcdc9,b1252a9d,07b5194c,,3a171ecb,c5 c50484,e8b83407,9727dd16']

(3a) Loading and splitting the data

We are now ready to start working with the actual CTR data, and our first task involves splitting it into training, validation, and test sets. Use the randomSplit method (https://spark.apache.org/docs/latest/api/python/pyspark.html#pyspark.RDD.randomSplit) with the specified weights and seed to create RDDs storing each of these datasets, and then cache (<a href="https://spark.apache.org/docs/latest/api/python/pyspark.html#pyspark.RDD.cache) each of these RDDs, as we will be accessing them multiple times in the remainder of this lab. Finally, compute the size of each dataset.

```
In [108]: # TODO: Replace <FILL IN> with appropriate code
    weights = [.8, .1, .1]
    seed = 42
    # Use randomSplit with weights and seed
    rawTrainData, rawValidationData, rawTestData = rawData.randomSplit(weights, seed)
    # Cache the data
    rawTrainData.cache()
    rawValidationData.cache()
    rawTestData.cache()

    nTrain = rawTrainData.count()
    nVal = rawValidationData.count()
    nTest = rawTestData.count()
    print nTrain, nVal, nTest, nTrain + nVal + nTest
    print rawData.take(1)
```

79911 10075 10014 100000

[u'0,1,1,5,0,1382,4,15,2,181,1,2,,2,68fd1e64,80e26c9b,fb936136,7b4723c4,25c83c9 8,7e0ccccf,de7995b8,1f89b562,a73ee510,a8cd5504,b2cb9c98,37c9c164,2824a5f6,1adce 6ef,8ba8b39a,891b62e7,e5ba7672,f54016b9,21ddcdc9,b1252a9d,07b5194c,,3a171ecb,c5 c50484,e8b83407,9727dd16']

- 1 test passed.
- 1 test passed.
- 1 test passed.

(3b) Extract features

We will now parse the raw training data to create an RDD that we can subsequently use to create an OHE dictionary. Note from the take() command in Part (3a) that each raw data point is a string containing several fields separated by some delimiter. For now, we will ignore the first field (which is the 0-1 label), and parse the remaining fields (or raw features). To do this, complete the implemention of the parsePoint function.

```
In [110]:
          # TODO: Replace <FILL IN> with appropriate code
          def parsePoint(point):
               """Converts a comma separated string into a list of (featureID, value) tuples.
              Note:
                   featureIDs should start at 0 and increase to the number of features - 1.
              Args:
                   point (str): A comma separated string where the first value is the label a
                       are features.
               Returns:
                   list: A list of (featureID, value) tuples.
              return [(featureID, value) for (featureID, value) in enumerate(point.split(','
          parsedTrainFeat = rawTrainData.map(parsePoint)
          numCategories = (parsedTrainFeat
                            .flatMap(lambda x: x)
                            .distinct()
                            .map(lambda x: (x[0], 1))
                            .reduceByKey(lambda x, y: x + y)
                            .sortByKey()
                            .collect())
          print numCategories[2][1]
```

855

In [111]: # TEST Extract features (3b)

Test.assertEquals(numCategories[2][1], 855, 'incorrect implementation of parsePoin Test.assertEquals(numCategories[32][1], 4, 'incorrect implementation of parsePoint

- 1 test passed.
- 1 test passed.

(3c) Create an OHE dictionary from the dataset

Note that parsePoint returns a data point as a list of (featureID, category) tuples, which is the same format as the sample dataset studied in Parts 1 and 2 of this lab. Using this observation, create an OHE dictionary using the function implemented in Part (2c). Note that we will assume for simplicity that all features in our CTR dataset are categorical.

In [112]: # TODO: Replace <FILL IN> with appropriate code
 ctrOHEDict = createOneHotDict(parsedTrainFeat)
 numCtrOHEFeats = len(ctrOHEDict.keys())
 print numCtrOHEFeats
 print ctrOHEDict[(0, '')]

233286 36325

In [113]: # TEST Create an OHE dictionary from the dataset (3c)

Test.assertEquals(numCtrOHEFeats, 233286, 'incorrect number of features in ctrOHED Test.assertTrue((0, '') in ctrOHEDict, 'incorrect features in ctrOHEDict')

- 1 test passed.
- 1 test passed.

(3d) Apply OHE to the dataset

Now let's use this OHE dictionary by starting with the raw training data and creating an RDD of LabeledPoint

(http://spark.apache.org/docs/1.3.1/api/python/pyspark.mllib.html#pyspark.mllib.regression.Lat objects using OHE features. To do this, complete the implementation of the parseOHEPoint function from Part (3b) and it uses the oneHot function from Part (1d).

In [114]: **from** pyspark.mllib.regression **import** LabeledPoint

```
In [149]:
          # TODO: Replace <FILL IN> with appropriate code
          def parseOHEPoint(point, OHEDict, numOHEFeats):
               """Obtain the label and feature vector for this raw observation.
              Note:
                  You must use the function `oneHotEncoding` in this implementation or later
                   of this lab may not function as expected.
              Args:
                   point (str): A comma separated string where the first value is the label a
                       are features.
                  OHEDict (dict of (int, str) to int): Mapping of (featureID, value) to uniq
                   numOHEFeats (int): The number of unique features in the training dataset.
              Returns:
                   LabeledPoint: Contains the label for the observation and the one-hot-encod
                       raw features based on the provided OHE dictionary.
              return LabeledPoint(point.split(',')[0], oneHotEncoding(parsePoint(point), OHE
          OHETrainData = rawTrainData.map(lambda point: parseOHEPoint(point, ctrOHEDict, num
          OHETrainData.cache()
          print OHETrainData.take(1)
          # Check that oneHotEncoding function was used in parseOHEPoint
          backupOneHot = oneHotEncoding
          oneHotEncoding = None
          withOneHot = False
          try: parseOHEPoint(rawTrainData.take(1)[0], ctrOHEDict, numCtrOHEFeats)
          except TypeError: withOneHot = True
          oneHotEncoding = backupOneHot
          [LabeledPoint(0.0, (233286, 386, 3119, 5284, 6851, 8957, 11948, 15688, 16268, 18913, 519]
```

```
In [116]: # TEST Apply OHE to the dataset (3d)
    numNZ = sum(parsedTrainFeat.map(lambda x: len(x)).take(5))
    numNZAlt = sum(OHETrainData.map(lambda lp: len(lp.features.indices)).take(5))
    Test.assertEquals(numNZ, numNZAlt, 'incorrect implementation of parseOHEPoint')
    Test.assertTrue(withOneHot, 'oneHotEncoding not present in parseOHEPoint')
```

1 test passed.

1 test passed.

Visualization 1: Feature frequency

We will now visualize the number of times each of the 233,286 OHE features appears in the training data. We first compute the number of times each feature appears, then bucket the features by these counts. The buckets are sized by powers of 2, so the first bucket corresponds to features that appear exactly once (2^0), the second to features that appear twice (2^1), the third to features that occur between three and four (2^1) times, the fifth

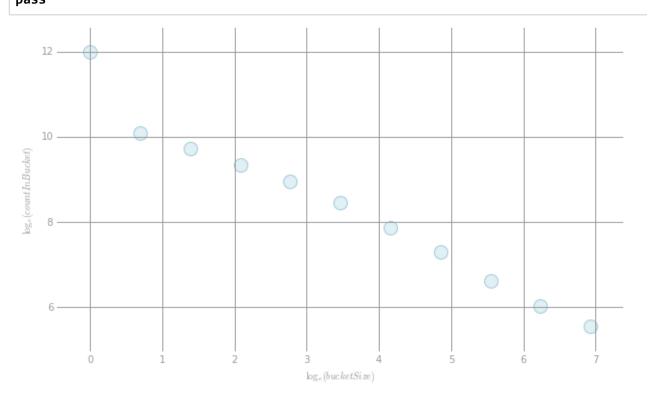
bucket is five to eight (2^3) times and so on. The scatter plot below shows the logarithm of the bucket thresholds versus the logarithm of the number of features that have counts that fall in the buckets.

```
In [117]: | def bucketFeatByCount(featCount):
               """Bucket the counts by powers of two."""
               for i in range(11):
                   size = 2 ** i
                   if featCount <= size:</pre>
                       return size
               return -1
           featCounts = (OHETrainData
                         .flatMap(lambda lp: lp.features.indices)
                         .map(lambda x: (x, 1))
                         .reduceByKey(lambda x, y: x + y))
           featCountsBuckets = (featCounts
                                 .map(lambda x: (bucketFeatByCount(x[1]), 1))
                                 .filter(lambda (k, v): k != -1)
                                 .reduceByKey(lambda x, y: x + y)
                                 .collect())
          print featCountsBuckets
```

```
[(32, 4755), (1024, 255), (2, 24076), (4, 16639), (8, 11440), (64, 2627), (128, 1476), (256, 748), (16, 7752), (512, 414), (1, 162813)]
```

In [118]:

```
%matplotlib inline
import matplotlib.pyplot as plt
x, y = zip(*featCountsBuckets)
x, y = np.log(x), np.log(y)
def preparePlot(xticks, yticks, figsize=(10.5, 6), hideLabels=False, gridColor='#9
                gridWidth=1.0):
    """Template for generating the plot layout."""
    plt.close()
    fig, ax = plt.subplots(figsize=figsize, facecolor='white', edgecolor='white')
    ax.axes.tick_params(labelcolor='#999999', labelsize='10')
    for axis, ticks in [(ax.get_xaxis(), xticks), (ax.get_yaxis(), yticks)]:
        axis.set ticks position('none')
        axis.set_ticks(ticks)
        axis.label.set color('#999999')
        if hideLabels: axis.set_ticklabels([])
    plt.grid(color=gridColor, linewidth=gridWidth, linestyle='-')
    map(lambda position: ax.spines[position].set visible(False), ['bottom', 'top',
    return fig, ax
# generate layout and plot data
fig, ax = preparePlot(np.arange(0, 10, 1), np.arange(4, 14, 2))
ax.set_xlabel(r'$\log_e(bucketSize)$'), ax.set_ylabel(r'$\log_e(countInBucket)$')
plt.scatter(x, y, s=14**2, c='#d6ebf2', edgecolors='#8cbfd0', alpha=0.75)
pass
```



(3e) Handling unseen features

We naturally would like to repeat the process from Part (3d), e.g., to compute OHE features for the validation and test datasets. However, we must be careful, as some categorical values will likely appear in new data that did not exist in the training data. To deal with this

situation, update the oneHotEncoding() function from Part (1d) to ignore previously unseen categories, and then compute OHE features for the validation data.

```
In [119]: # TODO: Replace <FILL IN> with appropriate code
          def oneHotEncoding(rawFeats, OHEDict, numOHEFeats):
             """Produce a one-hot-encoding from a list of features and an OHE dictionary.
             Note:
                 If a (featureID, value) tuple doesn't have a corresponding key in OHEDict
                 ignored.
             Args:
                 rawFeats (list of (int, str)): The features corresponding to a single obse
                     feature consists of a tuple of featureID and the feature's value. (e.g
                 OHEDict (dict): A mapping of (featureID, value) to unique integer.
                 numOHEFeats (int): The total number of unique OHE features (combinations o
                     value).
             Returns:
                 SparseVector: A SparseVector of length numOHEFeats with indicies equal to
                     identifiers for the (featureID, value) combinations that occur in the
                     with values equal to 1.0.
             return SparseVector(numOHEFeats, [(OHEDict[(featureID, value)],1) for (feature
                                                 if (featureID, value) in OHEDict])
          OHEValidationData = rawValidationData.map(lambda point: parseOHEPoint(point, ctrOH
          OHEValidationData.cache()
          print OHEValidationData.take(1)
          [LabeledPoint(0.0, (233286, [7639, 9282, 11431, 15655, 21732, 27739, 31381, 36325, 4681
          7,61951,66747,67374,68480,71539,76855,81587,85322,92157,97127,109795,110852,111
         015,117015,121552,141711,146496,147649,171128,184132,184687,185498,194763,19853
          7,201039,210717,213588,222162,222626,227709],[1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,
          In [120]:
         # TEST Handling unseen features (3e)
          numNZVal = (OHEValidationData
                     .map(lambda lp: len(lp.features.indices))
          Test.assertEquals(numNZVal, 372080, 'incorrect number of features')
          1 test passed.
```

Part 4: CTR prediction and logloss evaluation

(4a) Logistic regression

We are now ready to train our first CTR classifier. A natural classifier to use in this setting is log models the probability of a click-through event rather than returning a binary response, and wh probabilistic predictions are useful. First use <u>LogisticRegressionWithSGD</u> (https://spark.apache.org/docs/latest/api/python/pyspark.mllib.html#pyspark.mllib.classification

to train a model using OHETrainData with the given hyperparameter configuration. LogisticRes **LogisticRegressionModel**

(https://spark.apache.org/docs/latest/api/python/pyspark.mllib.html#pyspark.mllib.regression.L Next, use the LogisticRegressionModel.weights and LogisticRegressionModel.intercept at model's parameters. Note that these are the names of the object's attributes and should be callmodel.weights for a given model.

In [121]: | from pyspark.mllib.classification import LogisticRegressionWithSGD

fixed hyperparameters

numIters = 50

stepSize = 10.

regParam = 1e-6

regType = '12'

includeIntercept = True

In [122]:

TODO: Replace <FILL IN> with appropriate code model0 = model0 = LogisticRegressionWithSGD.train(OHETrainData,iterations=numIters ,regParam=regParam,regType=regTy sortedWeights = sorted(model0.weights)

print sortedWeights[:5], model0.intercept

/usr/local/spark/python/pyspark/mllib/classification.py:313: UserWarning: Depre cated in 2.0.0. Use ml.classification.LogisticRegression or LogisticRegressionW ithLBFGS.

"Deprecated in 2.0.0. Use ml.classification.LogisticRegression or "

[-0.45899236853575576, -0.37973707648623922, -0.3699655826675326, -0.3693496287 9928246, -0.32697945415010582] 0.56455084025

In [123]: # TEST Logistic regression (4a)

Test.assertTrue(np.allclose(model0.intercept, 0.56455084025), 'incorrect value fo Test.assertTrue(np.allclose(sortedWeights[0:5],

> [-0.45899236853575609, -0.37973707648623956, -0.36996558266753304, -0.36934962879928263, -0.32697945415010637]), 'incorrect value fo

- 1 test passed.
- 1 test passed.

(4b) Log loss

Throughout this lab, we will use log loss to evaluate the quality of models. Log loss is defined as:

$$\ell_{log}(p, y) = \begin{cases} -\log(p) & \text{if } y = 1\\ -\log(1 - p) & \text{if } y = 0 \end{cases}$$

where p is a probability between 0 and 1 and q is a label of either 0 or 1. Log loss is a standard evaluation criterion when predicting rare-events such as click-through rate prediction (it is also the criterion used in the Criteo Kaggle competition (https://www.kaggle.com/c/criteo-display-ad-challenge)). Write a function to compute log loss, and evaluate it on some sample inputs.

```
In [126]:
          # TODO: Replace <FILL IN> with appropriate code
          from math import log
          def computeLogLoss(p, y):
               """Calculates the value of log loss for a given probabilty and label.
              Note:
                   log(0) is undefined, so when p is 0 we need to add a small value (epsilon)
                   and when p is 1 we need to subtract a small value (epsilon) from it.
              Args:
                   p (float): A probabilty between 0 and 1.
                  y (int): A label. Takes on the values 0 and 1.
              Returns:
                   float: The log loss value.
              epsilon = 10e-12
              p = max(epsilon, min(1-epsilon,p))
              if (y == 1): return -\log(p)
              return -log(1-p)
          print computeLogLoss(.5, 1)
          print computeLogLoss(.5, 0)
          print computeLogLoss(.99, 1)
          print computeLogLoss(.99, 0)
          print computeLogLoss(.01, 1)
          print computeLogLoss(.01, 0)
          print computeLogLoss(0, 1)
          print computeLogLoss(1, 1)
          print computeLogLoss(1, 0)
          0.69314718056
          0.69314718056
          0.0100503358535
          4.60517018599
          4.60517018599
          0.0100503358535
          25.3284360229
          1.00000008275e-11
          25.3284359402
In [127]: # TEST Log loss (4b)
          Test.assertTrue(np.allclose([computeLogLoss(.5, 1), computeLogLoss(.01, 0), comput
                                       [0.69314718056, 0.0100503358535, 4.60517018599]),
                           'computeLogLoss is not correct')
          Test.assertTrue(np.allclose([computeLogLoss(0, 1), computeLogLoss(1, 1), computeLo
                                       [25.3284360229, 1.00000008275e-11, 25.3284360229]),
                           'computeLogLoss needs to bound p away from 0 and 1 by epsilon')
          1 test passed.
          1 test passed.
```

(4c) Baseline log loss

Next we will use the function we wrote in Part (4b) to compute the baseline log loss on the training data. A very simple yet natural baseline model is one where we always make the same prediction independent of the given datapoint, setting the predicted value equal to the fraction of training points that correspond to click-through events (i.e., where the label is one). Compute this value (which is simply the mean of the training labels), and then use it to compute the training log loss for the baseline model. The log loss for multiple observations is the mean of the individual log loss values.

In [128]: # TODO: Replace <FILL IN> with appropriate code # Note that our dataset has a very high click-through rate by design # In practice click-through rate can be one to two orders of magnitude lower classOneFracTrain = OHETrainData.map(lambda d: d.label).mean() print classOneFracTrain

> logLossTrBase = OHETrainData.map(lambda 1: computeLogLoss(classOneFracTrain, 1.lab print 'Baseline Train Logloss = {0:.3f}\n'.format(logLossTrBase)

0.22717773523

Baseline Train Logloss = 0.536

In [129]: # TEST Baseline log loss (4c)

Test.assertTrue(np.allclose(classOneFracTrain, 0.22717773523), 'incorrect value fo Test.assertTrue(np.allclose(logLossTrBase, 0.535844), 'incorrect value for logLoss

- 1 test passed.
- 1 test passed.

(4d) Predicted probability

In order to compute the log loss for the model we trained in Part (4a), we need to write code to generate predictions from this model. Write a function that computes the raw linear prediction from this logistic regression model and then passes it through a sigmoid function (http://en.wikipedia.org/wiki/Sigmoid function) $\sigma(t) = (1 + e^{-t})^{-1}$ to return the model's probabilistic prediction. Then compute probabilistic predictions on the training data.

Note that when incorporating an intercept into our predictions, we simply add the intercept to the value of the prediction obtained from the weights and features. Alternatively, if the intercept was included as the first weight, we would need to add a corresponding feature to our data where the feature has the value one. This is not the case here.

```
In [130]:
          # TODO: Replace <FILL IN> with appropriate code
          from math import exp # exp(-t) = e^{-t}
          def getP(x, w, intercept):
               """Calculate the probability for an observation given a set of weights and int
              Note:
                  We'll bound our raw prediction between 20 and -20 for numerical purposes.
              Args:
                  x (SparseVector): A vector with values of 1.0 for features that exist in t
                      observation and 0.0 otherwise.
                  w (DenseVector): A vector of weights (betas) for the model.
                  intercept (float): The model's intercept.
              Returns:
                  float: A probability between 0 and 1.
              rawPrediction = w.dot(x) + intercept
              # Bound the raw prediction value
              rawPrediction = min(rawPrediction, 20)
              rawPrediction = max(rawPrediction, -20)
              return 1 / (1 + exp(-rawPrediction))
          trainingPredictions = OHETrainData.map(lambda d: getP(d.features, model0.weights,
          print trainingPredictions.take(5)
```

[0.3026288202391113, 0.10362661997434096, 0.28363424783875607, 0.17846102057880 123, 0.538977537921885]

1 test passed.

(4e) Evaluate the model

We are now ready to evaluate the quality of the model we trained in Part (4a). To do this, first write a general function that takes as input a model and data, and outputs the log loss. Then run this function on the OHE training data, and compare the result with the baseline log loss.

```
In [135]: # TODO: Replace <FILL IN> with appropriate code
          def evaluateResults(model, data):
               """Calculates the log loss for the data given the model.
              Args:
                  model (LogisticRegressionModel): A trained logistic regression model.
                  data (RDD of LabeledPoint): Labels and features for each observation.
              Returns:
                  float: Log loss for the data.
              results = data.map(lambda d: computeLogLoss( getP(d.features, model0.weights,
              return results.mean()
          logLossTrLR0 = evaluateResults(model0, OHETrainData)
          print ('OHE Features Train Logloss:\n\tBaseline = {0:.3f}\n\tLogReg = {1:.3f}'
                  .format(logLossTrBase, logLossTrLR0))
          OHE Features Train Logloss:
                  Baseline = 0.536
                  LogReg = 0.457
```

In [136]: # TEST Evaluate the model (4e)

Test.assertTrue(np.allclose(logLossTrLR0, 0.456903), 'incorrect value for logLossT

1 test passed.

(4f) Validation log loss

Next, following the same logic as in Parts (4c) and 4(e), compute the validation log loss for both the baseline and logistic regression models. Notably, the baseline model for the validation data should still be based on the label fraction from the training dataset.

In [141]: # TEST Validation log loss (4f)

Test.assertTrue(np.allclose(logLossValBase, 0.527603), 'incorrect value for logLos Test.assertTrue(np.allclose(logLossValLR0, 0.456957), 'incorrect value for logLoss

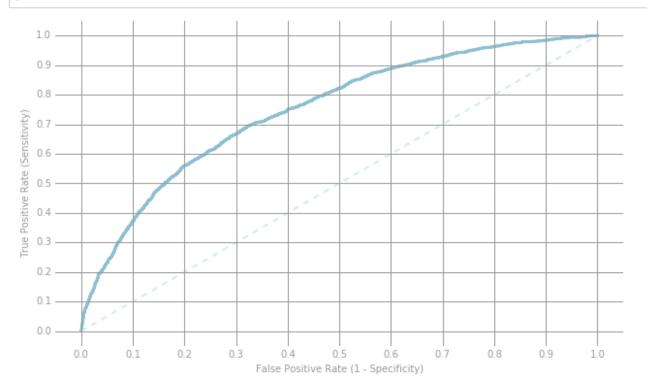
1 test passed.

1 test passed.

Visualization 2: ROC curve

We will now visualize how well the model predicts our target. To do this we generate a plot of the ROC curve. The ROC curve shows us the trade-off between the false positive rate and true positive rate, as we liberalize the threshold required to predict a positive outcome. A random model is represented by the dashed line.

```
labelsAndScores = OHEValidationData.map(lambda lp:
In [142]:
                                                       (lp.label, getP(lp.features, model0.we
          labelsAndWeights = labelsAndScores.collect()
          labelsAndWeights.sort(key=lambda (k, v): v, reverse=True)
          labelsByWeight = np.array([k for (k, v) in labelsAndWeights])
          length = labelsByWeight.size
          truePositives = labelsByWeight.cumsum()
          numPositive = truePositives[-1]
          falsePositives = np.arange(1.0, length + 1, 1.) - truePositives
          truePositiveRate = truePositives / numPositive
          falsePositiveRate = falsePositives / (length - numPositive)
          # Generate layout and plot data
          fig, ax = preparePlot(np.arange(0., 1.1, 0.1), np.arange(0., 1.1, 0.1))
          ax.set_xlim(-.05, 1.05), ax.set_ylim(-.05, 1.05)
          ax.set_ylabel('True Positive Rate (Sensitivity)')
          ax.set xlabel('False Positive Rate (1 - Specificity)')
          plt.plot(falsePositiveRate, truePositiveRate, color='#8cbfd0', linestyle='-', line
          plt.plot((0., 1.), (0., 1.), linestyle='--', color='#d6ebf2', linewidth=2.) # Bas
          pass
```



Part 5: Reduce feature dimension via feature hashing

(5a) Hash function

As we just saw, using a one-hot-encoding featurization can yield a model with good statistical accuracy. However, the number of distinct categories across all features is quite large -- recall that we observed 233K categories in the training data in Part (3c). Moreover, the full Kaggle training dataset includes more than 33M distinct categories, and the Kaggle dataset itself is just a small subset of Criteo's labeled data. Hence, featurizing via a one-hot-encoding representation would lead to a very large feature vector. To reduce the dimensionality of the feature space, we will use feature hashing.

####Below is the hash function that we will use for this part of the lab. We will first use this hash function with the three sample data points from Part (1a) to gain some intuition. Specifically, run code to hash the three sample points using two different values for numBuckets and observe the resulting hashed feature dictionaries.

```
from collections import defaultdict
In [143]:
          import hashlib
          def hashFunction(numBuckets, rawFeats, printMapping=False):
               """Calculate a feature dictionary for an observation's features based on hashi
              Note:
                  Use printMapping=True for debug purposes and to better understand how the
              Args:
                  numBuckets (int): Number of buckets to use as features.
                  rawFeats (list of (int, str)): A list of features for an observation.
                       (featureID, value) tuples.
                  printMapping (bool, optional): If true, the mappings of featureString to i
                      printed.
              Returns:
                  dict of int to float: The keys will be integers which represent the bucke
                      features have been hashed to. The value for a given key will contain
                       (featureID, value) tuples that have hashed to that key.
               .. .. ..
              mapping = {}
              for ind, category in rawFeats:
                  featureString = category + str(ind)
                  mapping[featureString] = int(int(hashlib.md5(featureString).hexdigest(), 1
              if(printMapping): print mapping
              sparseFeatures = defaultdict(float)
              for bucket in mapping.values():
                  sparseFeatures[bucket] += 1.0
              return dict(sparseFeatures)
          # Reminder of the sample values:
          # sampleOne = [(0, 'mouse'), (1, 'black')]
          # sampleTwo = [(0, 'cat'), (1, 'tabby'), (2, 'mouse')]
          # sampleThree = [(0, 'bear'), (1, 'black'), (2, 'salmon')]
```

```
# TODO: Replace <FILL IN> with appropriate code
In [144]:
          # Use four buckets
          sampOneFourBuckets = hashFunction(4, sampleOne, True)
          sampTwoFourBuckets = hashFunction(4, sampleTwo, True)
          sampThreeFourBuckets = hashFunction(4, sampleThree, True)
          # Use one hundred buckets
          sampOneHundredBuckets = hashFunction(100, sampleOne, True)
          sampTwoHundredBuckets = hashFunction(100, sampleTwo, True)
          sampThreeHundredBuckets = hashFunction(100, sampleThree, True)
          print '\t\t 4 Buckets \t\t\t 100 Buckets'
          print 'SampleOne:\t {0}\t\t {1}'.format(sampOneFourBuckets, sampOneHundredBuckets)
          print 'SampleTwo:\t {0}\t\t {1}'.format(sampTwoFourBuckets, sampTwoHundredBuckets)
          print 'SampleThree:\t {0}\t {1}'.format(sampThreeFourBuckets, sampThreeHundredBuck
          {'black1': 2, 'mouse0': 3}
          {'cat0': 0, 'tabby1': 0, 'mouse2': 2}
          {'bear0': 0, 'black1': 2, 'salmon2': 1}
          {'black1': 14, 'mouse0': 31}
          {'cat0': 40, 'tabby1': 16, 'mouse2': 62}
          {'bear0': 72, 'black1': 14, 'salmon2': 5}
                           4 Buckets
                                                            100 Buckets
          SampleOne:
                           {2: 1.0, 3: 1.0}
                                                            {14: 1.0, 31: 1.0}
          SampleTwo:
                           {0: 2.0, 2: 1.0}
                                                            {40: 1.0, 16: 1.0, 62: 1.0}
          SampleThree:
                           {0: 1.0, 1: 1.0, 2: 1.0}
                                                            {72: 1.0, 5: 1.0, 14: 1.0}
In [145]: # TEST Hash function (5a)
          Test.assertEquals(sampOneFourBuckets, {2: 1.0, 3: 1.0}, 'incorrect value for sampO
          Test.assertEquals(sampThreeHundredBuckets, {72: 1.0, 5: 1.0, 14: 1.0},
                             'incorrect value for sampThreeHundredBuckets')
```

1 test passed.

1 test passed.

(5b) Creating hashed features

Next we will use this hash function to create hashed features for our CTR datasets. First write a function that uses the hash function from Part (5a) with numBuckets = $2^{15} \approx 33 K$ to create a LabeledPoint with hashed features stored as a SparseVector. Then use this function to create new training, validation and test datasets with hashed features. Hint: parsedHashPoint is similar to parseOHEPoint from Part (3d).

```
In [150]:
          # TODO: Replace <FILL IN> with appropriate code
          def parseHashPoint(point, numBuckets):
               """Create a LabeledPoint for this observation using hashing.
              Args:
                   point (str): A comma separated string where the first value is the label a
                       features.
                   numBuckets: The number of buckets to hash to.
              Returns:
                   LabeledPoint: A LabeledPoint with a label (0.0 or 1.0) and a SparseVector
                       features.
               .. .. ..
              label = point.split(',')[0]
              features = parsePoint(point)
              return LabeledPoint(label, SparseVector(numBuckets, hashFunction(numBuckets, f
          numBucketsCTR = 2 ** 15
          hashTrainData = rawTrainData.map(lambda point: parseHashPoint(point,numBucketsCTR)
          hashTrainData.cache()
          hashValidationData = rawValidationData.map(lambda point: parseHashPoint(point,numB
          hashValidationData.cache()
          hashTestData = rawTestData.map(lambda point: parseHashPoint(point,numBucketsCTR))
          hashTestData.cache()
          print hashTrainData.take(1)
```

```
# TEST Creating hashed features (5b)
hashTrainDataFeatureSum = sum(hashTrainData
                           .map(lambda lp: len(lp.features.indices))
                           .take(20))
hashTrainDataLabelSum = sum(hashTrainData
                          .map(lambda lp: lp.label)
                          .take(100))
hashValidationDataFeatureSum = sum(hashValidationData
                                 .map(lambda lp: len(lp.features.indices))
                                .take(20))
hashValidationDataLabelSum = sum(hashValidationData
                               .map(lambda lp: lp.label)
                              .take(100))
hashTestDataFeatureSum = sum(hashTestData
                          .map(lambda lp: len(lp.features.indices))
                          .take(20))
hashTestDataLabelSum = sum(hashTestData
                        .map(lambda lp: lp.label)
                        .take(100))
Test.assertEquals(hashTrainDataFeatureSum, 772, 'incorrect number of features in h
Test.assertEquals(hashTrainDataLabelSum, 24.0, 'incorrect labels in hashTrainData'
Test.assertEquals(hashValidationDataFeatureSum, 776,
                   'incorrect number of features in hashValidationData')
Test.assertEquals(hashValidationDataLabelSum, 16.0, 'incorrect labels in hashValid
Test.assertEquals(hashTestDataFeatureSum, 774, 'incorrect number of features in ha
Test.assertEquals(hashTestDataLabelSum, 23.0, 'incorrect labels in hashTestData')
```

(5c) Sparsity

Since we have 33K hashed features versus 233K OHE features, we should expect OHE features to be sparser. Verify this hypothesis by computing the average sparsity of the OHE and the hashed training datasets.

Note that if you have a SparseVector named sparse, calling len(sparse) returns the total number of features, not the number features with entries. SparseVector objects have the attributes indices and values that contain information about which features are nonzero. Continuing with our example, these can be accessed using sparse.indices and sparse.values, respectively.

```
In []: # TODO: Replace <FILL IN> with appropriate code

def computeSparsity(data, d, n):
    """Calculates the average sparsity for the features in an RDD of LabeledPoints

Args:
    data (RDD of LabeledPoint): The LabeledPoints to use in the sparsity calcu
    d (int): The total number of features.
    n (int): The number of observations in the RDD.

Returns:
    float: The average of the ratio of features in a point to total features.

"""
    <FILL IN>

averageSparsityHash = computeSparsity(hashTrainData, numBucketsCTR, nTrain)
averageSparsityOHE = computeSparsity(OHETrainData, numCtrOHEFeats, nTrain)

print 'Average OHE Sparsity: {0:.7e}'.format(averageSparsityOHE)
print 'Average Hash Sparsity: {0:.7e}'.format(averageSparsityHash)
```

(5d) Logistic model with hashed features

Now let's train a logistic regression model using the hashed features. Run a grid search to find suitable hyperparameters for the hashed features, evaluating via log loss on the validation data. Note: This may take a few minutes to run. Use 1 and 10 for stepSizes and 1e-6 and 1e-3 for regParams.

```
In [ ]: numIters = 500
    regType = '12'
    includeIntercept = True

# Initialize variables using values from initial model training
    bestModel = None
    bestLogLoss = 1e10
```

```
In [ ]: | # TODO: Replace <FILL IN> with appropriate code
        stepSizes = <FILL IN>
        regParams = <FILL IN>
        for stepSize in stepSizes:
            for regParam in regParams:
                model = (LogisticRegressionWithSGD
                          .train(hashTrainData, numIters, stepSize, regParam=regParam, regT
                                 intercept=includeIntercept))
                 logLossVa = evaluateResults(model, hashValidationData)
                 print ('\tstepSize = {0:.1f}, regParam = {1:.0e}: logloss = {2:.3f}'
                        .format(stepSize, regParam, logLossVa))
                 if (logLossVa < bestLogLoss):</pre>
                     bestModel = model
                     bestLogLoss = logLossVa
        print ('Hashed Features Validation Logloss:\n\tBaseline = {0:.3f}\n\tLogReg = {1:.
                .format(logLossValBase, bestLogLoss))
```

```
In [ ]: # TEST Logistic model with hashed features (5d)
    Test.assertTrue(np.allclose(bestLogLoss, 0.4481683608), 'incorrect value for bestLogLoss')
```

Visualization 3: Hyperparameter heat map

We will now perform a visualization of an extensive hyperparameter search. Specifically, we will create a heat map where the brighter colors correspond to lower values of logLoss.

The search was run using six step sizes and six values for regularization, which required the training of thirty-six separate models. We have included the results below, but omitted the actual search to save time.

```
In [ ]: | from matplotlib.colors import LinearSegmentedColormap
        # Saved parameters and results. Eliminate the time required to run 36 models
        stepSizes = [3, 6, 9, 12, 15, 18]
        regParams = [1e-7, 1e-6, 1e-5, 1e-4, 1e-3, 1e-2]
        logLoss = np.array([[ 0.45808431,
                                           0.45808493, 0.45809113,
                                                                     0.45815333,
                                                                                  0.458792
                            [ 0.45188196, 0.45188306,
                                                        0.4518941,
                                                                     0.4520051,
                                                                                  0.453162
                            [ 0.44886478, 0.44886613, 0.44887974,
                                                                     0.44902096,
                                                                                  0.450561
                            [ 0.44706645, 0.4470698,
                                                        0.44708102, 0.44724251,
                                                                                  0.449055
                            [ 0.44588848, 0.44589365, 0.44590568, 0.44606631,
                                                                                  0.448071
                            [ 0.44508948, 0.44509474, 0.44510274, 0.44525007,
                                                                                  0.447383
        numRows, numCols = len(stepSizes), len(regParams)
        logLoss = np.array(logLoss)
        logLoss.shape = (numRows, numCols)
        fig, ax = preparePlot(np.arange(0, numCols, 1), np.arange(0, numRows, 1), figsize=
                              hideLabels=True, gridWidth=0.)
        ax.set_xticklabels(regParams), ax.set_yticklabels(stepSizes)
        ax.set xlabel('Regularization Parameter'), ax.set ylabel('Step Size')
        colors = LinearSegmentedColormap.from_list('blue', ['#0022ff', '#000055'], gamma=.
        image = plt.imshow(logLoss,interpolation='nearest', aspect='auto',
                            cmap = colors)
        pass
```

(5e) Evaluate on the test set

Finally, evaluate the best model from Part (5d) on the test set. Compare the resulting log loss with the baseline log loss on the test set, which can be computed in the same way that the

```
validation log loss was computed in Part (4f).
In [ ]: # TODO: Replace <FILL IN> with appropriate code
        # Log loss for the best model from (5d)
        logLossTest = <FILL IN>
        # Log loss for the baseline model
        logLossTestBaseline = <FILL IN>
        print ('Hashed Features Test Log Loss:\n\tBaseline = {0:.3f}\n\tLogReg = {1:.3f}\'
                .format(logLossTestBaseline, logLossTest))
        # TEST Evaluate on the test set (5e)
In [ ]:
        Test.assertTrue(np.allclose(logLossTestBaseline, 0.537438),
                         'incorrect value for logLossTestBaseline')
        Test.assertTrue(np.allclose(logLossTest, 0.455616931), 'incorrect value for logLos
```