DATASCI W261: Machine Learning at Scale

W261-4 Spring 2016
Week 12: Criteo CTR Project
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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 Criteo Labs (http://labs.criteo.com/) dataset that was used for a recent Kaggle competition (https://www.kaggle.com/c/criteo-display-ad-challenge).

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 Spark's Python API (https://spark.apache.org/docs/latest/api/python/pyspark.html#pyspark.RDD) and the relevant NumPy methods in the NumPy Reference (http://docs.scipy.org/doc/numpy/reference/index.html)

In [1]:

```
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 [2]:

```
# Data for manual OHE
# Note: the first data point does not include any value for the optional third
feature
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 [3]:

```
# 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 [4]:

```
# 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.
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 to the total number of features that occur in the dataset. By leveraging this sparsity and using sparse vector representations of OHE data, we can reduce storage and computational burdens. Below are a few sample vectors represented as dense numpy arrays. Use SparseVector (https://spark.apache.org/docs/latest/api/python/pyspark.mllib.html#pyspark.mllib.linalg.SparseVect to represent them in a sparse fashion, and verify that both the sparse and dense representations yield the same results when computing dot products (http://en.wikipedia.org/wiki/Dot_product) (we will later 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 vector and args is either a dictionary, a list of (index, value) pairs, or two separate arrays of indices and values (sorted by index). You'll need to create a sparse vector representation of each dense vector `aDense` and `bDense`.

In [5]:

```
import numpy as np
from pyspark.mllib.linalg import SparseVector
```

In [6]:

```
# 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 aSparse.dot(w)
print bDense.dot(w)
print bSparse.dot(w)
```

```
7.3
7.3
-0.5
-0.5
```

In [7]:

```
# TEST Sparse Vectors (1b)
Test.assertTrue(isinstance(aSparse, SparseVector), 'aSparse needs to be an inst
ance of SparseVector')
Test.assertTrue(isinstance(bSparse, SparseVector), 'aSparse needs to be an inst
ance of SparseVector')
Test.assertTrue(aDense.dot(w) == aSparse.dot(w),
                'dot product of aDense and w should equal dot product of aSpars
e and w')
Test.assertTrue(bDense.dot(w) == bSparse.dot(w),
                'dot product of bDense and w should equal dot product of bSpars
e and w')
```

```
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 [8]:

```
# 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 [9]:

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

In [10]:

```
# 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.
1 test passed.
1 test passed.
1 test passed.
1 test passed.
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).

```
In [11]:
```

```
# 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 so
rted.

    Args:
        rawFeats (list of (int, str)): The features corresponding to a single o
bservation.  Each
            feature consists of a tuple of featureID and the feature's value. (
e.g. sampleOne)
        OHEDict (dict): A mapping of (featureID, value) to unique integer.
        numOHEFeats (int): The total number of unique OHE features (combination
s of featureID and
            value).


    Returns:
        SparseVector: A SparseVector of length numOHEFeats with indicies equal
to the unique
            identifiers for the (featureID, value) combinations that occur in t
he observation and
            with values equal to 1.0.
    """
    sparse_index  = []
    for feature in rawFeats:
        if feature in OHEDict:
            sparse_index.append(OHEDict[feature])

    return SparseVector(numOHEFeats, sorted(sparse_index), [1.]*len(sparse_inde
x))

# Calculate the number of features in sampleOHEDictManual
numSampleOHEFeats = len(sampleOHEDictManual.keys())

# Run oneHotEnoding on sampleOne
sampleOneOHEFeat = oneHotEncoding(sampleOne, sampleOHEDictManual, numSampleOHEF
eats)

print sampleOneOHEFeat
```

```
(7,[2,3],[1.0,1.0])
```

In [12]:

```
# 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')], sampleOHEDictMan

ual,
                                 numSampleOHEFeats), SparseVector(7, [2,3], [1.
0,1.0]),
                  '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.

In [13]:

```
# TODO: Replace <FILL IN> with appropriate code
sampleOHEData = sampleDataRDD.map(lambda x: oneHotEncoding(x, sampleOHEDictManu
al, numSampleOHEFeats))
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})]
```

In [14]:

```
# TEST Apply OHE to a dataset (1e)
sampleOHEDataValues = sampleOHEData.collect()
Test.assertTrue(len(sampleOHEDataValues) == 3, 'sampleOHEData should have three
 elements')
Test.assertEquals(sampleOHEDataValues[0], SparseVector(7, {2: 1.0, 3: 1.0}),
                  'incorrect OHE for first sample')
Test.assertEquals(sampleOHEDataValues[1], SparseVector(7, {1: 1.0, 4: 1.0, 5: 1
.0}),
                  'incorrect OHE for second sample')
Test.assertEquals(sampleOHEDataValues[2], SparseVector(7, {0: 1.0, 3: 1.0, 6: 1
.0}),
                  'incorrect OHE for third sample')
```

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

In [15]:

```
# TODO: Replace <FILL IN> with appropriate code
sampleDistinctFeats = (sampleDataRDD
                       .flatMap(lambda x: x)
                       .distinct())
```

In [16]:

```
# TEST Pair RDD of (featureID, category) (2a)
Test.assertEquals(sorted(sampleDistinctFeats.collect()),
                  [(0, 'bear'), (0, 'cat'), (0, 'mouse'), (1, 'black'),
                   (1, 'tabby'), (2, 'mouse'), (2, 'salmon')],
                  'incorrect value for sampleDistinctFeats')
```

```
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. In this exercise, any valid mapping is acceptable. Use [zipWithIndex (https://spark.apache.org/docs/latest/api/python/pyspark.html#pyspark.RDD.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)](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 keys and integers.

In [17]:

```
# TODO: Replace <FILL IN> with appropriate code
sampleOHEDict = (sampleDistinctFeats
                           .zipWithIndex()
                           .collectAsMap()
                )
print sampleOHEDict
```

```
{(2, 'mouse'): 3, (0, 'cat'): 5, (0, 'bear'): 0, (2, 'salmon'): 1, (
1, 'tabby'): 2, (1, 'black'): 6, (0, 'mouse'): 4}
```

In [18]:

```
# TEST OHE Dictionary from distinct features (2b)
Test.assertEquals(sorted(sampleOHEDict.keys()),
                  [(0, 'bear'), (0, 'cat'), (0, 'mouse'), (1, 'black'),
                   (1, 'tabby'), (2, 'mouse'), (2, 'salmon')],
                  'sampleOHEDict has unexpected keys')
Test.assertEquals(sorted(sampleOHEDict.values()), range(7), 'sampleOHEDict has
unexpected values')
```

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

(2c) Automated creation of an OHE dictionary

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 [19]:

```
# 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 ea
ch observation is
            made up of a list of (featureID, value) tuples.

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

sampleOHEDictAuto = createOneHotDict(sampleDataRDD)
print sampleOHEDictAuto
```

```
{(2, 'mouse'): 3, (0, 'cat'): 5, (0, 'bear'): 0, (2, 'salmon'): 1, (
1, 'tabby'): 2, (1, 'black'): 6, (0, 'mouse'): 4}
```

In [20]:

```
# TEST Automated creation of an OHE dictionary (2c)
Test.assertEquals(sorted(sampleOHEDictAuto.keys()),
                  [(0, 'bear'), (0, 'cat'), (0, 'mouse'), (1, 'black'),
                   (1, 'tabby'), (2, 'mouse'), (2, 'salmon')],
                  'sampleOHEDictAuto has unexpected keys')
Test.assertEquals(sorted(sampleOHEDictAuto.values()), range(7),
                  'sampleOHEDictAuto has unexpected values')
```

```
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 [Criteo agreement (http://labs.criteo.com/downloads/2014-kaggle-display-advertising-challenge-dataset/)](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.

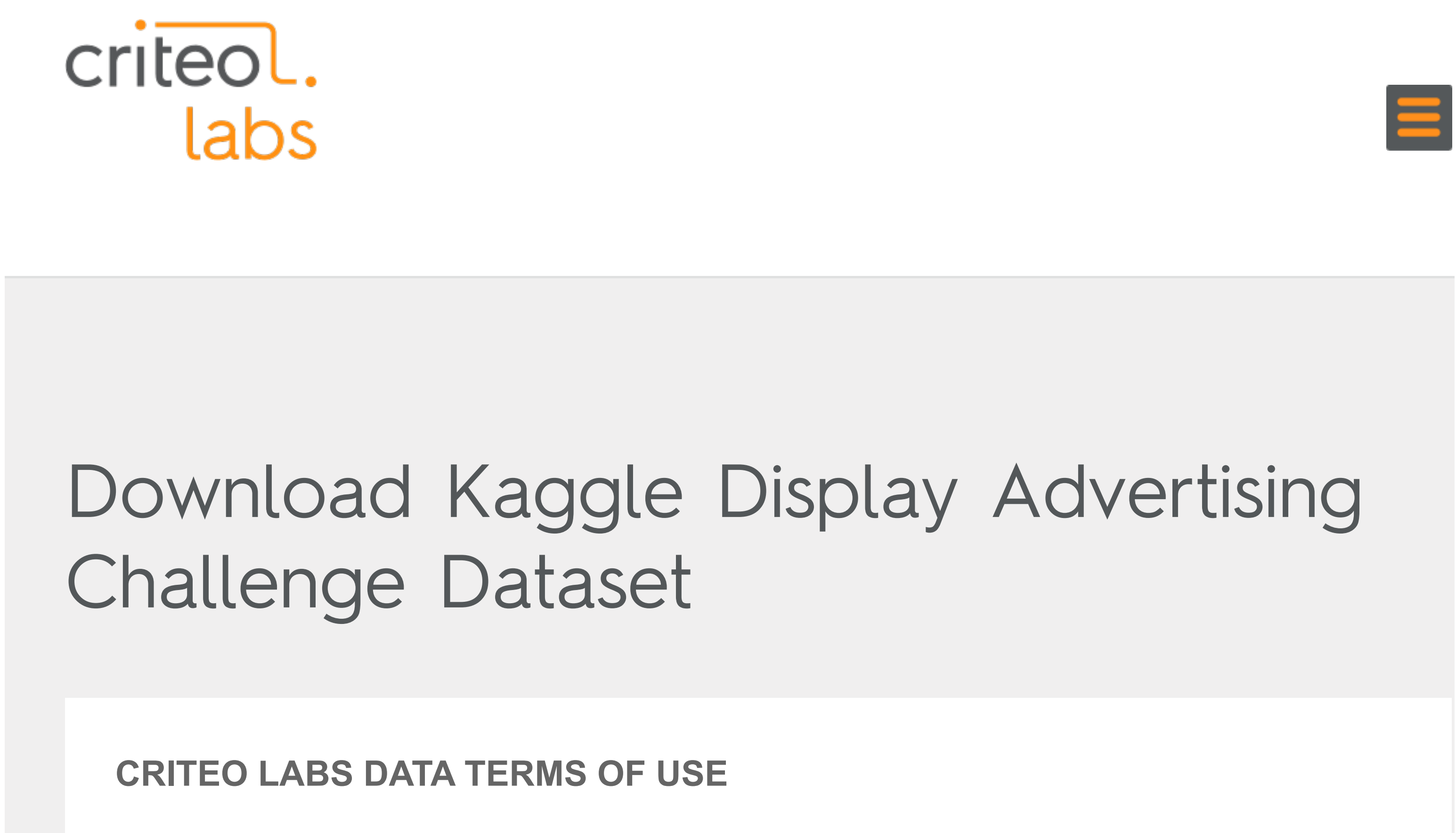
Note that the download could take a few minutes, depending upon your connection speed.

In [21]:

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

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

Out[21]:



In [22]:

```
# 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 = '<FILL IN>'


url = url.strip()
baseDir = os.path.join('data')
inputPath = os.path.join('cs190', '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 you
r Jupyter root ' +
              '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
    try:
        urllib.urlretrieve(url, os.path.basename(urlparse.urlsplit(url).path))
    except IOError:
        print 'Unable to download and store: {0}'.format(url)

    extractTar()
```

```
Check your download url.  Are you downloading the Sample dataset?
```

In [23]:

```
import os.path

# NOTE: the data file is on the HDFS file system, not the OS file system
baseDir = os.path.join('/user/rcordell/hw12')

inputPath = os.path.join('input', 'dac_sample.txt')
fileName = os.path.join(baseDir, inputPath)

#if os.path.isfile(fileName):
rawData = (sc
           .textFile(fileName, 2)
           .map(lambda x: x.replace('\t', ',')))  # work with either ',' or '\t
' separated data
print rawData.take(1)
```

```
[u'0,1,1,5,0,1382,4,15,2,181,1,2,,2,68fd1e64,80e26c9b,fb936136,7b472
3c4,25c83c98,7e0ccccf,de7995b8,1f89b562,a73ee510,a8cd5504,b2cb9c98,3
7c9c164,2824a5f6,1adce6ef,8ba8b39a,891b62e7,e5ba7672,f54016b9,21ddcd
c9,b1252a9d,07b5194c,,3a171ecb,c5c50484,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)](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 (https://spark.apache.org/docs/latest/api/python/pyspark.html#pyspark.RDD.cache)](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 [24]:

```
# 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, see
d)
# 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,fb936136,7b472
3c4,25c83c98,7e0ccccf,de7995b8,1f89b562,a73ee510,a8cd5504,b2cb9c98,3
7c9c164,2824a5f6,1adce6ef,8ba8b39a,891b62e7,e5ba7672,f54016b9,21ddcd
c9,b1252a9d,07b5194c,,3a171ecb,c5c50484,e8b83407,9727dd16']
```

In [25]:

```
# TEST Loading and splitting the data (3a)
Test.assertTrue(all([rawTrainData.is_cached, rawValidationData.is_cached, rawTe
stData.is_cached]),
                'you must cache the split data')
Test.assertEquals(nTrain, 79911, 'incorrect value for nTrain')
Test.assertEquals(nVal, 10075, 'incorrect value for nVal')
Test.assertEquals(nTest, 10014, 'incorrect value for nTest')
```

```
1 test passed.
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 [26]:

```
# TODO: Replace <FILL IN> with appropriate code
def parsePoint(point):
    """Converts a comma separated string into a list of (featureID, value) tupl
es.

    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 labe
l and the rest
            are features.

    Returns:
        list: A list of (featureID, value) tuples.
    """
    feature_tuples = []
    rawFeatures = point.split(',')
    for i,feature in enumerate(rawFeatures):
        if i > 0:
            feature_tuples.append((i-1, feature))

    return feature_tuples


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 [27]:

```
# TEST Extract features (3b)
Test.assertEquals(numCategories[2][1], 855, 'incorrect implementation of parseP
oint')
Test.assertEquals(numCategories[32][1], 4, 'incorrect implementation of parsePo
int')
```

```
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 [28]:

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

```
233286
36164
```

In [29]:

```
# TEST Create an OHE dictionary from the dataset (3c)
Test.assertEquals(numCtrOHEFeats, 233286, 'incorrect number of features in ctrO
HEDict')
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.LabeledI](http://spark.apache.org/docs/1.3.1/api/python/pyspark.mllib.html#pyspark.mllib.regression.LabeledPoint) objects using OHE features. To do this, complete the implementation of the `parseOHEPoint` function. Hint: `parseOHEPoint` is an extension of the `parsePoint` function from Part (3b) and it uses the `oneHotEncoding` function from Part (1d).

In [30]:

```
from pyspark.mllib.regression import LabeledPoint
```

In [31]:

```
# 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 la
ter portions
        of this lab may not function as expected.

    Args:
        point (str): A comma separated string where the first value is the labe
l and the rest
            are features.
        OHEDict (dict of (int, str) to int): Mapping of (featureID, value) to u
nique integer.
        numOHEFeats (int): The number of unique features in the training datase
t.

    Returns:
        LabeledPoint: Contains the label for the observation and the one-hot-en
coding of the
            raw features based on the provided OHE dictionary.
    """
    features = parsePoint(point)
    label = point.split(',')[0]
    OHEfeature_vector = oneHotEncoding(features, OHEDict, numOHEFeats)

    return LabeledPoint(label, OHEfeature_vector)

OHETrainData = rawTrainData.map(lambda point: parseOHEPoint(point, ctrOHEDict,
numCtrOHEFeats))
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,[386,3077,6799,8264,8862,11800,12802,1612
5,17551,18566,29331,33132,39525,55794,61786,81396,82659,93573,96929,
100677,109699,110646,112132,120260,128596,132397,132803,140620,16067
5,185498,190370,191146,195925,202664,204273,206055,222737,225958,229
942],[1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.
0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.
0,1.0,1.0,1.0,1.0,1.0,1.0]))]
```

In [32]:

```
# 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 (2^2) 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 [33]:

```
def bucketFeatByCount(featCount):
    """Bucket the counts by powers of two."""
    for i in range(11):
        size = 2 ** i
        if featCount <= size:
            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
```

```
[(256, 748), (1024, 255), (2, 24076), (4, 16639), (32, 4755), (8, 11
440), (64, 2627), (128, 1476), (16, 7752), (512, 414), (1, 162813)]
```

In [34]:

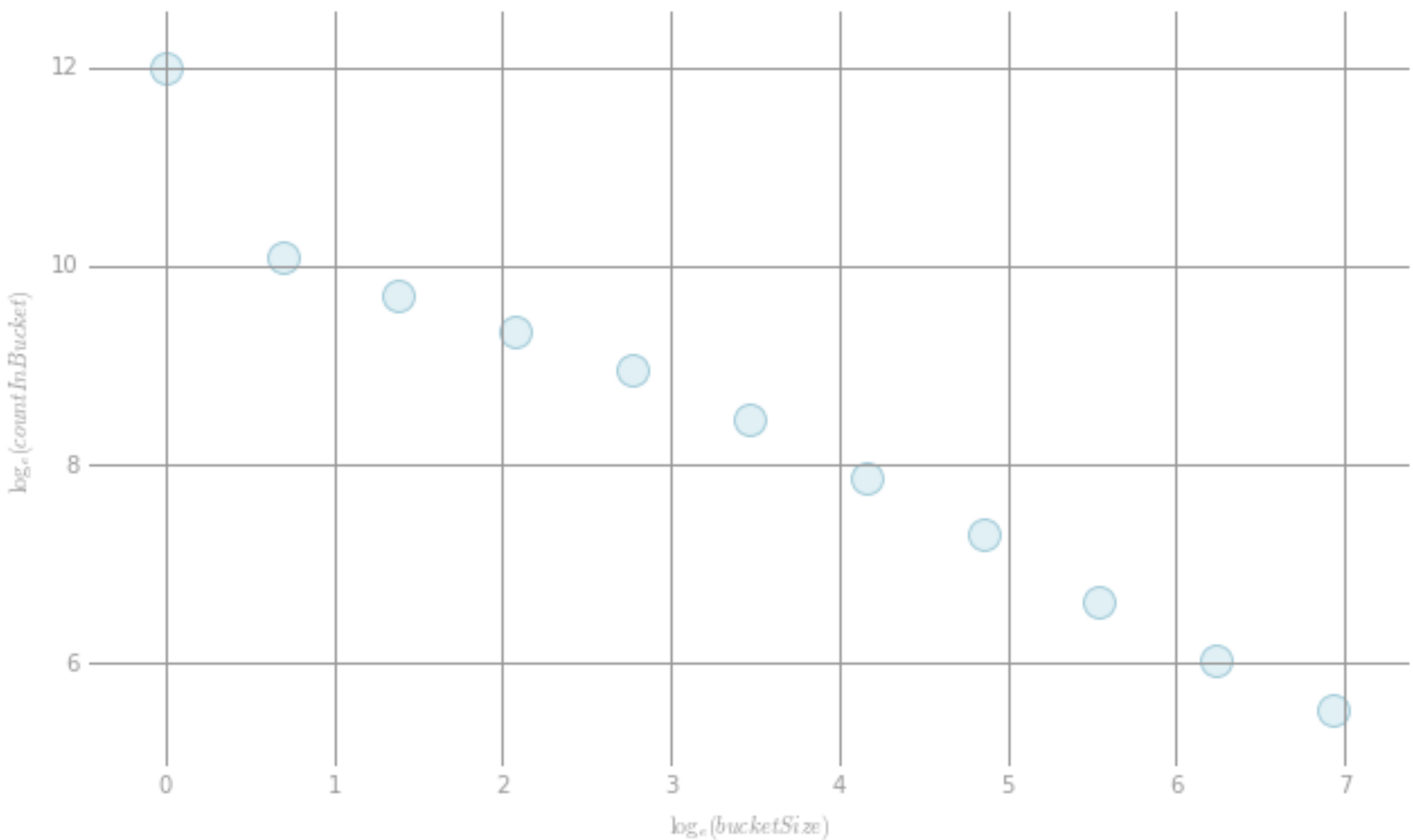
```
%matplotlib inline
import matplotlib.pyplot as plt

x, y = zip(*featCountsBuckets)
x, y = np.log(x), np.log(y)
```

```
def preparePlot(xticks, yticks, figsize=(10.5, 6), hideLabels=False, gridColor=
'#999999',
                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', 'to
p', 'left', 'right'])
    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
```

```
/Users/rcordell/Documents/MIDS/W261/W261env/lib/python2.7/site-packa
ges/matplotlib/font_manager.py:273: UserWarning: Matplotlib is build
ing the font cache using fc-list. This may take a moment.
  warnings.warn('Matplotlib is building the font cache using fc-list
. This may take a moment.')
```



(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 [35]:

```
# 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 OHEDi
ct it should be
        ignored.

    Args:
        rawFeats (list of (int, str)): The features corresponding to a single o
bservation.  Each
            feature consists of a tuple of featureID and the feature's value. (
e.g. sampleOne)
        OHEDict (dict): A mapping of (featureID, value) to unique integer.
        numOHEFeats (int): The total number of unique OHE features (combination
s of featureID and
            value).

    Returns:
        SparseVector: A SparseVector of length numOHEFeats with indicies equal
to the unique
            identifiers for the (featureID, value) combinations that occur in t
he observation and
            with values equal to 1.0.
    """
    sparse_index  = []
    for feature in rawFeats:
        if feature in OHEDict:
            sparse_index.append(OHEDict[feature])

    return SparseVector(numOHEFeats, sorted(sparse_index), [1.]*len(sparse_inde
x))



OHEValidationData = rawValidationData.map(lambda point: parseOHEPoint(point, ct
rOHEDict, numCtrOHEFeats))
OHEValidationData.cache()
print OHEValidationData.take(1)
```

```
[LabeledPoint(0.0, (233286,[7576,9187,15510,21585,31213,36164,39525,
49198,61786,66603,67218,68211,68311,73035,76672,81329,81396,91981,96
929,98450,109699,110946,117015,121552,141711,146496,147649,171128,18
4132,184687,185498,194763,198537,201039,210717,213588,222162,222626,
227709],[1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0
,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0
,1.0,1.0,1.0,1.0,1.0,1.0,1.0]))]
```

In [36]:

```
# TEST Handling unseen features (3e)
numNZVal = (OHEValidationData
            .map(lambda lp: len(lp.features.indices))
            .sum())
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 logistic regression, since it models the probability of a click-through event rather than returning a binary response, and when working with rare events, probabilistic predictions are useful. First use [LogisticRegressionWithSGD (https://spark.apache.org/docs/latest/api/python/pyspark.mllib.html#pyspark.mllib.classification.Log](https://spark.apache.org/docs/latest/api/python/pyspark.mllib.html#pyspark.mllib.classification.Log) to train a model using `OHETrainData` with the given hyperparameter configuration. `LogisticRegressionWithSGD` returns a [LogisticRegressionModel (https://spark.apache.org/docs/latest/api/python/pyspark.mllib.html#pyspark.mllib.regression.Logisti](https://spark.apache.org/docs/latest/api/python/pyspark.mllib.html#pyspark.mllib.regression.Logisti) Next, use the `LogisticRegressionModel.weights` and `LogisticRegressionModel.intercept` attributes to print out the model's parameters. Note that these are the names of the object's attributes and should be called using a syntax like `model.weights` for a given `model`.

In [37]:

```
from pyspark.mllib.classification import LogisticRegressionWithSGD

# fixed hyperparameters
numIters = 50
stepSize = 10.
regParam = 1e-6
regType = 'l2'
includeIntercept = True
```

In [38]:

```
# TODO: Replace <FILL IN> with appropriate code
model0 = LogisticRegressionWithSGD.train(OHETrainData,
                                         iterations = numIters,
                                         step=stepSize,
                                         miniBatchFraction=1.0,
                                         initialWeights=None,
                                         regParam=regParam,
                                         regType=regType,
                                         intercept=includeIntercept,
                                         validateData=True,
                                         convergenceTol=0.001)
sortedWeights = sorted(model0.weights)
print sortedWeights[:5], model0.intercept
```

```
[-0.45899236853575609, -0.37973707648623956, -0.36996558266753304, -
0.36934962879928263, -0.32697945415010637] 0.56455084025
```

In [39]:

```
# TEST Logistic regression (4a)
Test.assertTrue(np.allclose(model0.intercept,  0.56455084025), 'incorrect value
 for model0.intercept')
Test.assertTrue(np.allclose(sortedWeights[0:5],
                [-0.45899236853575609, -0.37973707648623956, -0.369965582667533
04,
                 -0.36934962879928263, -0.32697945415010637]), 'incorrect value
 for model0.weights')
```

```
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:

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

where p is a probability between 0 and 1 and y 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)](https://www.kaggle.com/c/criteo-display-ad-challenge)). Write a function to compute log loss, and evaluate it on some sample inputs.

In [40]:

```
# 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 (epsil
```

```
on) to it
        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
    operand = 0.0
    if y > 0:
        operand = p
    else:
        operand = 1-p
    if operand > 0.0:
        if operand < 1.0:
            return -log(operand)
        else:
            return -log(operand-epsilon)
    else:
        return -log(operand+epsilon)

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
25.3284360229
```

In [41]:

```
# TEST Log loss (4b)
Test.assertTrue(np.allclose([computeLogLoss(.5, 1), computeLogLoss(.01, 0), com
puteLogLoss(.01, 1)],
                            [0.69314718056, 0.0100503358535, 4.60517018599]),
                'computeLogLoss is not correct')
Test.assertTrue(np.allclose([computeLogLoss(0, 1), computeLogLoss(1, 1), comput
eLogLoss(1, 0)],
                            [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 [42]:

```
# 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 = float(OHETrainData.filter(lambda x: x.label == 1.0).count()
)/nTrain
print classOneFracTrain

logLossTrBase = (OHETrainData.map(lambda x: computeLogLoss(classOneFracTrain, x
.label))\
                            .reduce(lambda a,b: a+b))/nTrain
print 'Baseline Train Logloss = {0:.3f}\n'.format(logLossTrBase)
```

```
0.22717773523
Baseline Train Logloss = 0.536
```

In [43]:

```
# TEST Baseline log loss (4c)
Test.assertTrue(np.allclose(classOneFracTrain, 0.22717773523), 'incorrect value
 for classOneFracTrain')
Test.assertTrue(np.allclose(logLossTrBase, 0.535844), 'incorrect value for logL
ossTrBase')
```

```
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 [44]:

```
# 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
intercept.

    Note:
        We'll bound our raw prediction between 20 and -20 for numerical purpose
s.

    Args:
        x (SparseVector): A vector with values of 1.0 for features that exist i
n this
            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 = x.dot(w)+intercept

    # Bound the raw prediction value
    rawPrediction = min(rawPrediction, 20)
    rawPrediction = max(rawPrediction, -20)
    return 1.0/(1+exp(-rawPrediction))

trainingPredictions = OHETrainData.map(lambda p: getP(p.features, model0.weight
s, model0.intercept))

print trainingPredictions.take(5)
```

```
[0.3026288202391113, 0.10362661997434088, 0.283634247838756, 0.17846
102057880123, 0.5389775379218853]
```

In [45]:

```
# TEST Predicted probability (4d)
Test.assertTrue(np.allclose(trainingPredictions.sum(), 18135.4834348),
                'incorrect value for trainingPredictions')
```

```
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 [46]:

```
# 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.
    """

    log_loss = data.map(lambda p: computeLogLoss(getP(p.features,
                        model.weights, model.intercept), p.label))\
                    .reduce(lambda a,b: a+b)/(data.count())
    return log_loss

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 [47]:

```
# TEST Evaluate the model (4e)
Test.assertTrue(np.allclose(logLossTrLR0, 0.456903), 'incorrect value for logLo
ssTrLR0')
```

```
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 [48]:

```
# TODO: Replace <FILL IN> with appropriate code
logLossValBase = (OHEValidationData.map(lambda x: computeLogLoss(classOneFracTr
ain, x.label))\
                            .reduce(lambda a,b: a+b))/nVal

logLossValLR0 = evaluateResults(model0, OHEValidationData)
print ('OHE Features Validation Logloss:\n\tBaseline = {0:.3f}\n\tLogReg = {1:.
3f}'
       .format(logLossValBase, logLossValLR0))
```

```
OHE Features Validation Logloss:
        Baseline = 0.528
        LogReg = 0.457
```

In [49]:

```
# TEST Validation log loss (4f)
Test.assertTrue(np.allclose(logLossValBase, 0.527603), 'incorrect value for log
LossValBase')
Test.assertTrue(np.allclose(logLossValLR0, 0.456957), 'incorrect value for logL
ossValLR0')
```

```
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.

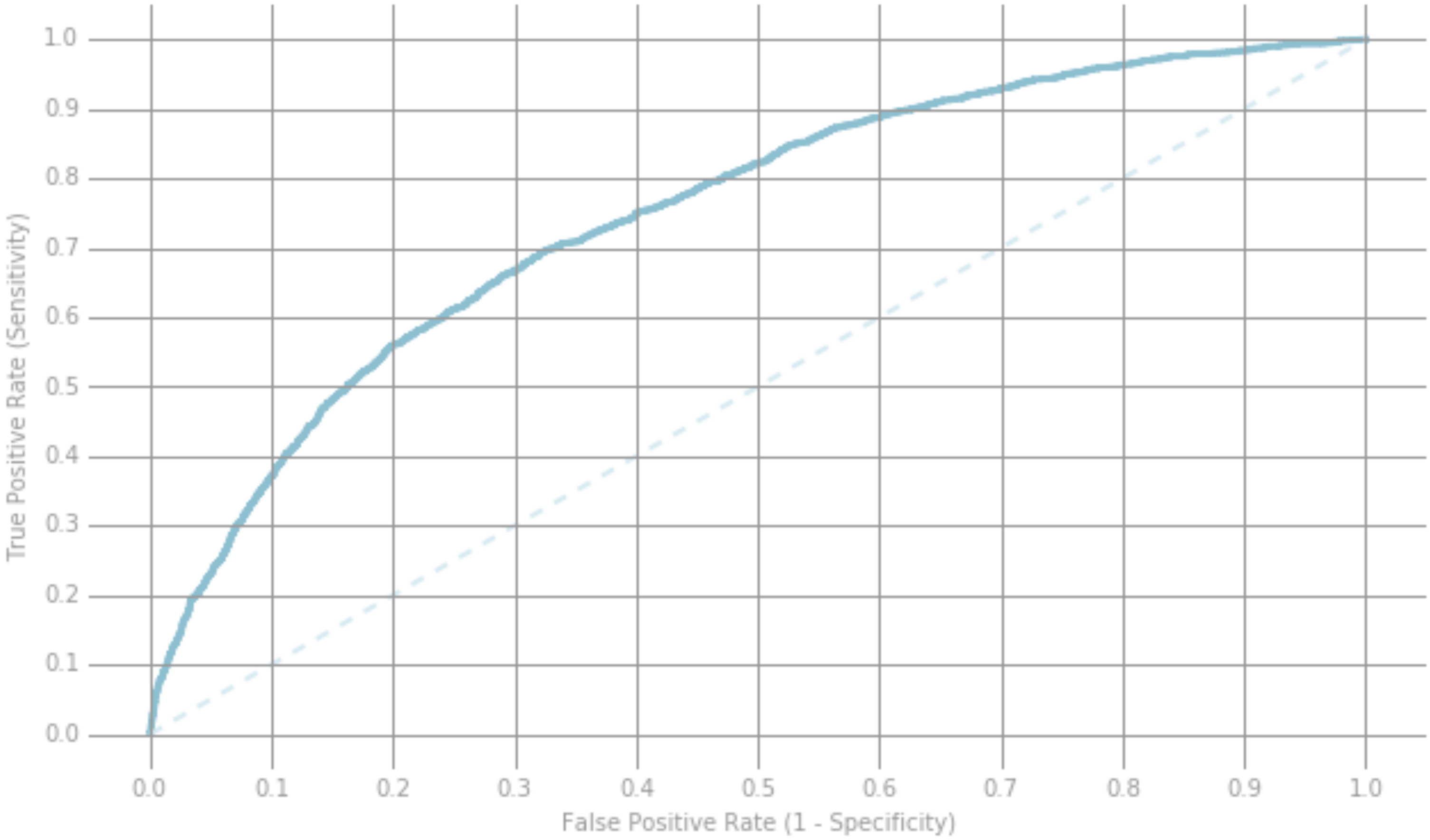
In [50]:

```
labelsAndScores = OHEValidationData.map(lambda lp:
                                        (lp.label, getP(lp.features, model0
.weights, model0.intercept)))
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='-', l
inewidth=3.)
plt.plot((0., 1.), (0., 1.), linestyle='--', color='#d6ebf2', linewidth=2.)  #
Baseline model
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.

In [51]:

```
from collections import defaultdict
import hashlib

def hashFunction(numBuckets, rawFeats, printMapping=False):
    """Calculate a feature dictionary for an observation's features based on ha
shing.

    Note:
        Use printMapping=True for debug purposes and to better understand how t
he hashing works.

    Args:
        numBuckets (int): Number of buckets to use as features.
        rawFeats (list of (int, str)): A list of features for an observation.
Represented as
            (featureID, value) tuples.
        printMapping (bool, optional): If true, the mappings of featureString t
o index will be
            printed.

    Returns:
        dict of int to float:  The keys will be integers which represent the bu
ckets that the
            features have been hashed to.  The value for a given key will conta
in the count of the
            (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()
, 16) % numBuckets)
    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')]
```

In [52]:

```
# TODO: Replace <FILL IN> with appropriate code
# 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, sampOneHundredBucke
ts)
print 'SampleTwo:\t {0}\t\t {1}'.format(sampTwoFourBuckets, sampTwoHundredBucke
ts)
print 'SampleThree:\t {0}\t {1}'.format(sampThreeFourBuckets, sampThreeHundredB
uckets)
```

```
{'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, 1
```

In [53]:

```
# TEST Hash function (5a)
Test.assertEquals(sampOneFourBuckets, {2: 1.0, 3: 1.0}, 'incorrect value for sa
mpOneFourBuckets')
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 = $2^{15} \approx 33K$ 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 [54]:

```
# 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 labe
l and the rest are
            features.
        numBuckets: The number of buckets to hash to.

    Returns:
        LabeledPoint: A LabeledPoint with a label (0.0 or 1.0) and a SparseVect
or of hashed
            features.
    """
    label = float(point.split(',')[0])
    hashed_features = hashFunction(numBuckets,
                                   parsePoint(point), False)
    sparse_features = SparseVector(numBuckets,
                                   sorted(hashed_features.keys()),
                                   [1.0]*len(hashed_features.keys()))
    return LabeledPoint(label, sparse_features)

numBucketsCTR = 2 ** 15
hashTrainData = rawTrainData.map(lambda point: parseHashPoint(point, numBuckets
CTR))
hashTrainData.cache()
hashValidationData = rawValidationData.map(lambda point: parseHashPoint(point,
numBucketsCTR))
hashValidationData.cache()
hashTestData = rawTestData.map(lambda point: parseHashPoint(point, numBucketsCT
R))
hashTestData.cache()

print hashTrainData.take(1)
```

```
[LabeledPoint(0.0, (32768,[1305,2883,3807,4814,4866,4913,6952,7117,9
985,10316,11512,11722,12365,13893,14735,15816,16198,17761,19274,2160
4,22256,22563,22785,24855,25202,25533,25721,26487,26656,27668,28211,
29152,29402,29873,30039,31484,32493,32708],[1.0,1.0,1.0,1.0,1.0,1.0,
1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,
1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0,1.0]))]
```

In [55]:

```
# 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 i
n hashTrainData')
Test.assertEquals(hashTrainDataLabelSum, 24.0, 'incorrect labels in hashTrainDa
ta')
Test.assertEquals(hashValidationDataFeatureSum, 776,
                  'incorrect number of features in hashValidationData')
Test.assertEquals(hashValidationDataLabelSum, 16.0, 'incorrect labels in hashVa
lidationData')
Test.assertEquals(hashTestDataFeatureSum, 774, 'incorrect number of features in
 hashTestData')
Test.assertEquals(hashTestDataLabelSum, 23.0, 'incorrect labels in hashTestData
')
```

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

(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 [56]:

```
# TODO: Replace <FILL IN> with appropriate code
def computeSparsity(data, d, n):
    """Calculates the average sparsity for the features in an RDD of LabeledPoi
nts.

    Args:
        data (RDD of LabeledPoint): The LabeledPoints to use in the sparsity ca
lculation.
        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 feature
s.
    """
    num_counted_features = data.map(lambda point: point.features.numNonzeros())
\
                               .reduce(lambda a,b: a+b)
    return float(num_counted_features)/float(d)/float(n)

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)
```

```
Average OHE Sparsity: 1.6717677e-04
Average Hash Sparsity: 1.1805561e-03
```

In [57]:

```
# TEST Sparsity (5c)
Test.assertTrue(np.allclose(averageSparsityOHE, 1.6717677e-04),
                'incorrect value for averageSparsityOHE')
Test.assertTrue(np.allclose(averageSparsityHash, 1.1805561e-03),
                'incorrect value for averageSparsityHash')
```

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

(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 [58]:

```
numIters = 500

regType = 'l2'
includeIntercept = True

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

In [59]:

```
# TODO: Replace <FILL IN> with appropriate code
stepSizes =[1,10]
regParams = [1e-6,1e-3]
for stepSize in stepSizes:
    for regParam in regParams:
        model = (LogisticRegressionWithSGD
                 .train(hashTrainData, numIters, stepSize, regParam=regParam, r
egType=regType,
                        intercept=includeIntercept))
        logLossVa = evaluateResults(model, hashValidationData)
        print ('\tstepSize = {0:.1f}, regParam = {1:.0e}: logloss = {2:.3f}'
               .format(stepSize, regParam, logLossVa))
        if (logLossVa < bestLogLoss):
            bestModel = model
            bestLogLoss = logLossVa

print ('Hashed Features Validation Logloss:\n\tBaseline = {0:.3f}\n\tLogReg = {
1:.3f}'
       .format(logLossValBase, bestLogLoss))
```

```
        stepSize = 1.0, regParam = 1e-06: logloss = 0.475
        stepSize = 1.0, regParam = 1e-03: logloss = 0.475
        stepSize = 10.0, regParam = 1e-06: logloss = 0.450
        stepSize = 10.0, regParam = 1e-03: logloss = 0.452
Hashed Features Validation Logloss:
        Baseline = 0.528
        LogReg = 0.450
```

In [60]:

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

```
1 test failed. incorrect value for bestLogLoss
```

The above tests fail but it is not understood why. The output is slightly different than the test value and this has been corroborated with several people in the class. The same is true for the last set of tests.

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 [61]:

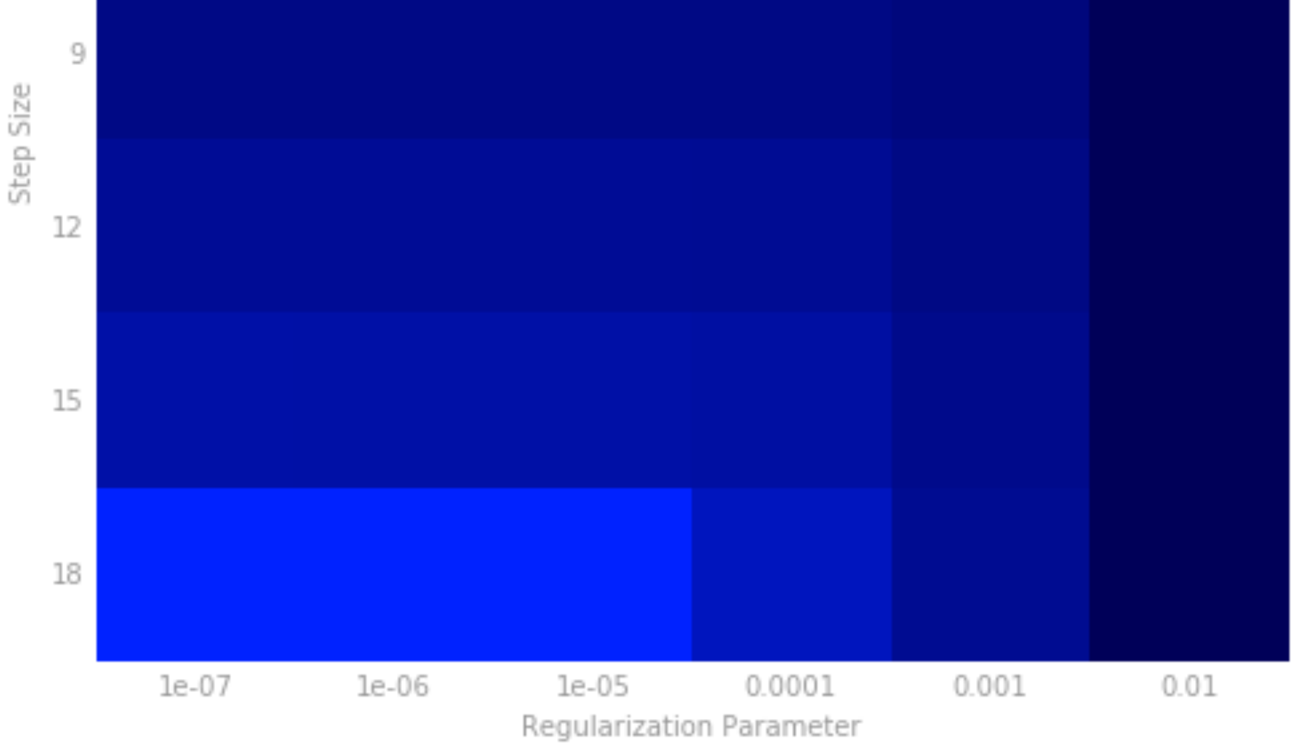
```
from matplotlib.colors import LinearSegmentedColormap

# 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,  0.458
79221,  0.46556321],
                    [ 0.45188196,  0.45188306,  0.4518941,   0.4520051,   0.453
16284,  0.46396068],
                    [ 0.44886478,  0.44886613,  0.44887974,  0.44902096,  0.450
5614,   0.46371153],
                    [ 0.44706645,  0.4470698,   0.44708102,  0.44724251,  0.449
05525,  0.46366507],
                    [ 0.44588848,  0.44589365,  0.44590568,  0.44606631,  0.448
07106,  0.46365589],
                    [ 0.44508948,  0.44509474,  0.44510274,  0.44525007,  0.447
38317,  0.46365405]])

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), figsi
ze=(8, 7),
                      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'], gamm
a=.2)
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 [62]:

```
stepSize = 10.0
regParam = 1e-06
model = (LogisticRegressionWithSGD
         .train(hashTrainData, numIters, stepSize, regParam=regParam, regType=r
egType,
                intercept=includeIntercept))
```

In [63]:

```
# TODO: Replace <FILL IN> with appropriate code
# Log loss for the best model from (5d)

logLossTest = evaluateResults(model, hashTestData)

# Log loss for the baseline model
logLossTestBaseline = (hashTestData.map(lambda x: computeLogLoss(classOneFracTr
ain, x.label))\
                            .reduce(lambda a,b: a+b))/nTest

print ('Hashed Features Test Log Loss:\n\tBaseline = {0:.3f}\n\tLogReg = {1:.8f
}'
       .format(logLossTestBaseline, logLossTest))
```

```
Hashed Features Test Log Loss:
        Baseline = 0.537
        LogReg = 0.45711746
```

In [64]:

```
# TEST Evaluate on the test set (5e)
Test.assertTrue(np.allclose(logLossTestBaseline, 0.537438),
                'incorrect value for logLossTestBaseline')
Test.assertTrue(np.allclose(logLossTest, 0.455616931), 'incorrect value for log
LossTest')
```

```
1 test passed.
1 test failed. incorrect value for logLossTest
```

This is another case of where a test failed without understanding why. The output value is corroborated by others in class, and despite many re-writes the algorithm gives the same output.

HW12.2 OPTIONAL Homework

Implement a decision tree algorithm for regression for two input continuous variables and one categorical input variable on a single core computer using Python. Use the IRIS dataset to evaluate your code, where the input variables are:

```
Petal.Length
Petal.Width
Species
```

and the target or output variable is

```
Sepal.Length.
```

Use the same dataset to train and test your implementation. Stop expanding nodes once you have less than ten (10) examples (along with the usual stopping criteria). Report the mean squared error for your implementation and contrast that with the MSE from scikit-learn's implementation on this dataset (http://scikit-learn.org/stable/modules/generated/sklearn.tree.DecisionTreeRegressor.html)

My simple approach is to first create a structure to encode the tree based on the table structure we reviewed in class. Then a simple algorithms to move across the tree one level at a time computing a split of the node based on the variance computation.

In [65]:

```
from sklearn import datasets
iris = datasets.load_iris()
```

In [66]:

```
# Transform the iris data set features and labels into the train data set
# The first position in the train_data vector is the label
# Species is encoded into dummy variables, or OHE. With only 3 species
# the OHE is simple so dictionary is used.
train_data = []
for i,feature_vector in enumerate(iris.data[:, :5]):
    species = [0.,0.,0.]
    species[iris.target[i]] = 1.
    new_feature = [feature_vector[0],
                   feature_vector[2],
                   feature_vector[3]]
    new_feature.extend(species)
    train_data.append(new_feature)
print train_data[0]
```

```
[5.0999999999999996, 1.3999999999999999, 0.20000000000000001, 1.0, 0
.0, 0.0]
```

In [67]:

```
from operator import itemgetter
import numpy as np
from collections import namedtuple

# Define a namedtuple to help with tracking the data
# attr_split_val = (MSE, index) - the index into the sorted data set for the mi
numum MSE
# attr_idx       = the attribute index on which the MSE was minimized
# data_left      = the left data set from the split
# data_right     = the right data set from the split
Split = namedtuple('Split', ['attr_split_val', 'attr_idx', 'data_left', 'data_r
ight'])

# Define a namedtuple for the tree table
# attr_idx       = the attribute index for the node split
# attr_split_val = the attribute value for the node split
# left_child     = node id of the left child, -1 if leaf node
# right_child    = node id of the right child, -1 if leaf node
# pred_value     = predicted value (mean of the data in that node), None if int
erior node
Node = namedtuple('Node', ['attr_idx','attr_split_val','left_child','right_chil
d','pred_value'])

# At each node we need to determine the best split by finding the split point f
or each variable.
# We do this by finding the lowest average label value we get by splitting into
 two groups by that
# variable. We do this for each variable.
def find_best_split(data):
    """Given a list of feature vectors find the best split across the features
by minimizing the
        mean squared error
```

```
        mean squared error:

        Input: [label, feature1, feature2, feature3, feature4, feature5]

        Output: (((MSE, index), variable_idx))

    """

    splits = []
    for j in range(1,len(data[0])):
        data = sorted(data, key=itemgetter(j))
        mse = []
        for s, feature_vector in enumerate(data):
            r_l = data[:s+1]
            r_r = data[s:]
            mu_l = np.mean(r_l[0])
            mu_r = np.mean(r_r[0])
            mse_l = float(len(r_l))/float(len(data))*sum(map(lambda x: (x[0]-mu_l)**2, r_l))
            mse_r = float(len(r_r))/float(len(data))*sum(map(lambda x: (x[0]-mu_r)**2, r_r))
            mse.append(mse_l + mse_r)

        splits.append(min((v, idx) for (idx, v) in enumerate(mse)))

    # the minimum out of all the attributes is one we want to use
    # the idx is the attribute value, so it would be feature_vector[idx+1]
    split_point = min((v, idx) for (idx, v) in enumerate(splits))

    # recover the right and left data sets sorted by that chosen attribute
    data = sorted(data, key=itemgetter(split_point[1]))
    data_l = data[:split_point[0][1]]
    data_r = data[split_point[0][1]:]
    split_val = data[split_point[0][1]][split_point[1]]

    return Split(split_val, split_point[1], data_l, data_r)

def find_prediction(data):
    return np.mean([x[0] for x in data])
```

In [68]:

```
# test the find_best_split function
data_split = find_best_split(train_data)

print data_split.attr_split_val
print data_split.attr_idx

# test the find_prediction function
print find_prediction(data_split.data_left)
print find_prediction(data_split.data_right)
```

```
0.0
3
5.87674418605
5.82990654206
```

In [69]:

```
def build_subtree(tree, node, data, leaf_size_limit):
    """
    Recursive algorithm
    Given a node and the data for that node, build the subtree for that node. E
valuate the stop
    criteria to decide whether to continue building more subtrees.

    Input: node - int, node id
           data - feature vectors associated with that node

    Output: builds the Tree table

    """
    # find the best split for this data set
    split = find_best_split(data)
    tree[node] = Node(split.attr_idx, split.attr_split_val, node+1, node+2, Non
e)

    if len(split.data_left) < leaf_size_limit:
        left_prediction = find_prediction(split.data_left)
        tree[node+1] = Node(-1, -1, -1, -1, left_prediction)
    else:
        build_subtree(tree, node+1, split.data_left, leaf_size_limit)

    if len(split.data_right) < leaf_size_limit:
        right_prediction = find_prediction(split.data_right)
        tree[node+2] = Node(-1, -1, -1, -1, right_prediction)
        return
    else:
        build_subtree(tree, node+2, split.data_right, leaf_size_limit)

    return
```

In [104]:

```
tree = {}

build_subtree(tree, 1, train_data, 10)
```

In [105]:

```
tree
```

Out[105]:

```
{1: Node(attr_idx=3, attr_split_val=0.0, left_child=2, right_child=3
, pred_value=None),
 2: Node(attr_idx=0, attr_split_val=5.7999999999999998, left_child=3
, right_child=4, pred_value=None),
 3: Node(attr_idx=3, attr_split_val=0.0, left_child=4, right_child=5
, pred_value=None),
 4: Node(attr_idx=4, attr_split_val=0.0, left_child=5, right_child=6
, pred_value=None),
 5: Node(attr_idx=2, attr_split_val=0.20000000000000001, left_child=
6, right_child=7, pred_value=None),
 6: Node(attr_idx=0, attr_split_val=4.7999999999999998, left_child=7
, right_child=8, pred_value=None),
 7: Node(attr_idx=2, attr_split_val=0.20000000000000001, left_child=
8, right_child=9, pred_value=None),
 8: Node(attr_idx=0, attr_split_val=5.0, left_child=9, right_child=1
0, pred_value=None),
 9: Node(attr_idx=2, attr_split_val=0.29999999999999999, left_child=
10, right_child=11, pred_value=None),
 10: Node(attr_idx=1, attr_split_val=1.6000000000000001, left_child=
11, right_child=12, pred_value=None),
 11: Node(attr_idx=2, attr_split_val=0.40000000000000002, left_child
=12, right_child=13, pred_value=None),
 12: Node(attr_idx=-1, attr_split_val=-1, left_child=-1, right_child
=-1, pred_value=5.3428571428571425),
 13: Node(attr_idx=2, attr_split_val=2.2999999999999998, left_child=
14, right_child=15, pred_value=None),
 14: Node(attr_idx=-1, attr_split_val=-1, left_child=-1, right_child
=-1, pred_value=5.7999999999999998),
 15: Node(attr_idx=1, attr_split_val=5.7000000000000002, left_child=
16, right_child=17, pred_value=None),
 16: Node(attr_idx=-1, attr_split_val=-1, left_child=-1, right_child
=-1, pred_value=6.25),
 17: Node(attr_idx=-1, attr_split_val=-1, left_child=-1, right_child
=-1, pred_value=7.0428571428571445)}
```

In [106]:

```
# now create a prediction function
def dt_predict(tree, X):
    """
    Given a tree as defined in the table structure and the feature vector
    predict y_hat

    Input: tree table
           a feature vector, X

    Output: predicted value, y_hat
    """

    node = 1

    while tree[node].attr_idx != -1:
        v = X[tree[node].attr_idx]
        t = tree[node].attr_split_val
        if v>=t:
            node = node+2
        else:
            node = node+1

    return tree[node].pred_value
```

In [107]:

```
from sklearn import metrics

X = [x[1:] for x in train_data]
Y = [y[0] for y in train_data]

y_hat = [dt_predict(tree, x) for x in X]

print metrics.mean_squared_error(Y, y_hat)
```

1.10391972789

In [108]:

```
from sklearn.tree import DecisionTreeRegressor

X = [x[1:] for x in train_data]
Y = [y[0] for y in train_data]

dt = DecisionTreeRegressor(min_samples_split=10, min_samples_leaf = 10, max_fea
tures=1)

model = dt.fit(X,Y)
y_hat = [model.predict(x)[0] for x in X]

print  metrics.mean_squared_error(Y, y_hat)
```

```
0.189286562054
```

The best MSE from my tree regressor is 0.8465 with a leaf size limit of 5 while that from the Scikit-Learn module is 0.121.

I used different parameters for both tree algorithms and it seems that setting fewer items per leaf node does not improve the accuracy lower than a leaf size limit of 5 and actually accuracy gets worse with smaller leaf limits.

In [ ]:

```

```