```
In [1]: % matplotlib inline

import numpy as np
import scipy as sp
import math
import scipy.stats as scp
import matplotlib as mpl
from numpy import linalg as la
from numpy import random as rand
from matplotlib import pyplot as plt
from sklearn import svm
from sklearn.metrics import mean_squared_error
```

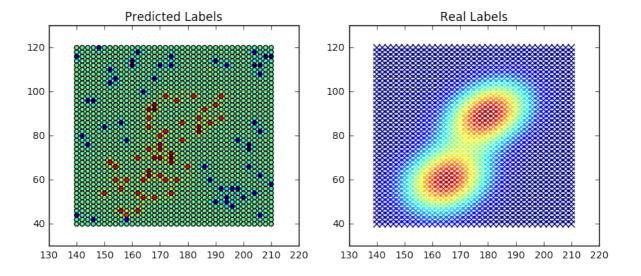
```
In [45]: Data = np.genfromtxt("TrainingRidge.csv", delimiter=',', skip_heade
    r=True,dtype='float').T

X = Data[0:2,:].T
Y = Data[2,:][np.newaxis,:].T

Valid = np.genfromtxt("ValidationRidge-Y.csv", delimiter=',', skip_header=True,dtype='float').T

Xvalid = Valid[0:2,:].T
Yvalid = Valid[2,:][np.newaxis,:].T
```

```
# train
In [49]:
         nsvm = svm.NuSVR(kernel='rbf',nu=0.5)
         nsvm.fit(X,Y)
         #predict
         pred y = nsvm.predict(Xvalid)
         pred_y.astype(int)
         fig = plt.figure()
         fig.set size inches(10,4)
         plt.subplot(121)
         plt.scatter(Xvalid[:,0],Xvalid[:,1], c=pred_y )
         plt.title('Predicted Labels')
         plt.subplot(122)
         plt.scatter(Xvalid[:,0], Xvalid[:,1], c=Yvalid, marker = 'x', s = 50
         plt.title('Real Labels')
         plt.show()
```



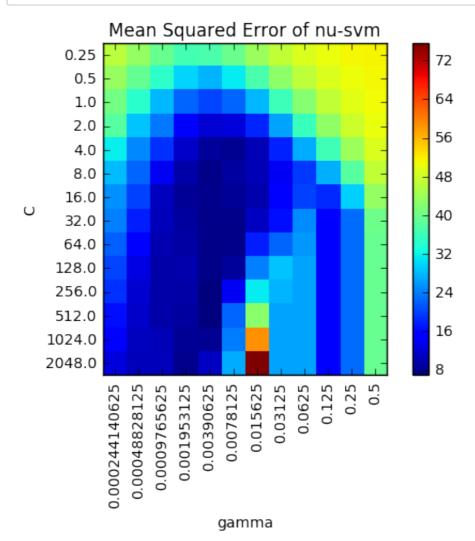
```
In [50]: def cross_valid(X, y, g,c, n_folds=10, nu=0.5):
              Synopsis:
                  w opt, b opt, lambda opt = cross validation(X, Y, L, n fold
         s=10)
             Arguments:
                  X:
                                2D array of data (features x samples)
                  Y:
                                Vector of true labels (1 x samples)
                  L:
                                List of lambdas to cross validate (1 x #lambd
         as)
                                Number of nested folds
                  n folds:
              Output:
                  w opt:
                                optimal weight vector
                                optimal bias
                  b opt:
                  lambda opt: the lambda with the lowest MSE
              , , ,
             X = X.T
             y = y.T
             d, n = X.shape
             samples per fold = int(float(n)/ float(n folds))
             rates = np.empty(n folds)
             idx = np.arange(n) # np.random.permutation(n) # np.arange(n)
              for j in range(n folds):
                  # extract one fold for testing
                  idx te = idx[j*samples per fold:(j+1)*samples per fold]
                  # get the train data
                  X tr = np.delete(X, idx te, axis=1)
                  y tr = np.delete(y, idx te, axis=1)
                  # get the test data
                  X \text{ te} = X[:,idx \text{ te}]
                  y_te = y[:,idx_te]
                  # train the model
                  c svm = svm.NuSVR(kernel='rbf',nu=nu,gamma = g,C = c)
                  c_svm.fit(X_tr.T,y_tr.T)
                  # predict the label
                  y pred = c svm.predict(X te.T)
                  rates[j] = mean squared error(y te.T,y pred)
             return np.min(rates)
```

```
In [73]: C = 2.**np.arange(-2,12,1)
    gamma = 2.** np.arange(-12,0,1)
    nu = 0.5

mse = np.zeros((gamma.shape[0],C.shape[0]))

for j,g in enumerate(gamma):
    for i, c in enumerate(C):
        mse[j][i] = cross_valid(X,Y,g,c)
```

```
In [102]: plt.imshow(mse.T, interpolation='nearest')
    plt.colorbar()
    plt.title("Mean Squared Error of nu-svm")
    plt.xlabel("gamma")
    plt.ylabel("C")
    plt.xticks(np.arange(gamma.shape[0]),gamma, rotation = "vertical")
    plt.yticks(np.arange(C.shape[0]),C, rotation = "horizontal")
    plt.show()
```



```
In [99]: # find best combination of C and gamma
   idx_g, idx_c = (np.unravel_index(mse.argmin(), mse.shape))
   opt_g = gamma[idx_g]
   opt_c = C[idx_c]
```

