

1. (a) Loading Red Wine data set...

In [62]:

```
import urllib
urllib.urlretrieve('https://archive.ics.uci.edu/ml/machine-learning-databases/wine-quality/winequality-
red.csv', 'winequality-red.csv')
import numpy as np
with open('winequality-red.csv') as f:
    lines = (line for line in f)
    data = np.loadtxt(lines, delimiter=';', skiprows=1)
q = data[:,11]
```

1. (b) Loading White Wine data set...

In [63]:

```
urllib.urlretrieve('https://archive.ics.uci.edu/ml/machine-learning-databases/wine-quality/winequality-
white.csv', 'winequality-white.csv')
with open('winequality-white.csv') as fw:
    lines_white = (line for line in fw)
    data_white = np.loadtxt(lines_white, delimiter=';', skiprows=1)
q_white = data_white[:,11]
```

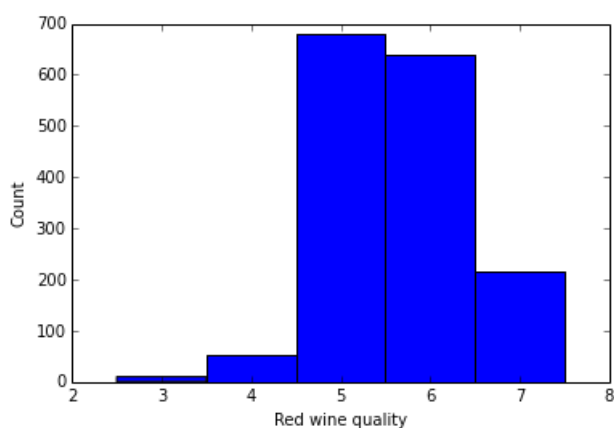
2. (a) Plotting Red Wine Quality distribution...

In [64]:

```
%matplotlib inline
import pylab as plt
plt.hist(q, bins=np.arange(q.min(), q.max()+1), align='left')
plt.xlabel('Red wine quality')
plt.ylabel('Count')
```

Out[64]:

<matplotlib.text.Text at 0x7f67db0>



2. (b) Plotting White Wine Quality distribution...

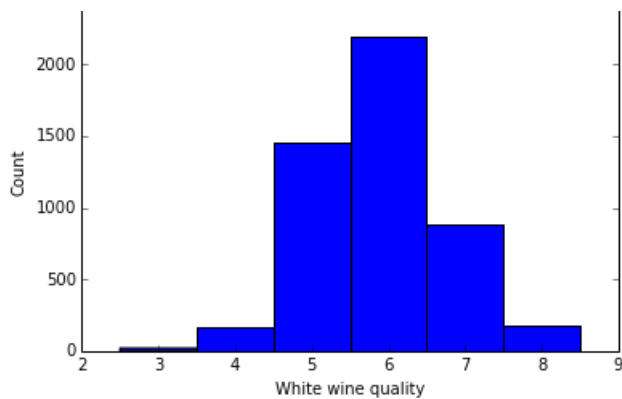
In [65]:

```
%matplotlib inline
import pylab as plt
plt.hist(q_white, bins=np.arange(q_white.min(), q_white.max()+1), align='left')
plt.xlabel('White wine quality')
plt.ylabel('Count')
```

Out[65]:

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3. Comments on these distributions

4. Linear regression

In [66]:

```
from numpy.linalg import inv

def fitRegressionModelAndGetSquareError(showPlot, order):
    N = data.shape[0] #get tuple (numRows, numCols)
    np.random.shuffle(data) # randomly shuffle the data

    # split the data in 70% train and 30% test
    train = data[:int(N*0.7)]
    test = data[int(N*0.7):]

    X_train = train[:, :11]
    X_train = X_train ** (order)
    X_train = np.c_[np.ones(train.shape[0]), X_train] # append 1s as first column
    q_train = train[:, 11]
    # data is now splitted into train and test

    XtX = np.dot(X_train.T, X_train)
    XtXI = inv(XtX)
    XtXIXt = np.dot(XtXI, X_train.T)
    # Fitting the regression model to the training data.
    # Depending on the order parameter, we can fit linear (order = 1), square or higher order
    # regression models.
    wbar = np.dot(XtXIXt, q_train)
    # Optimal values for the wbar are now found.
    # For these values we are minimizing the Loss against the train data

    X_test = test[:, :11]
    X_test = X_test ** (order)
    X_test = np.c_[np.ones(test.shape[0]), X_test]
    q_test = test[:, 11]
    f_test = np.dot(X_test, wbar) # getting the prediction values

    meanSquareError = ((q_test - f_test) ** 2).mean()

    # Part 4(c) Scatter plot showing predictions vs true values
    if showPlot:
        print "Linear regression with minimised square loss. Mean Square Error =", meanSquareError
        plt.scatter(f_test, q_test, color='blue')
        plt.xlabel('Red wine predicted quality')
        plt.ylabel('Red wine test data quality')
    return meanSquareError
```

4 (a), (b) and (c) The function above splits the data in 70% train and 30% test. Fits the linear regression to the training data and prints the Mean Square Error

In [67]:

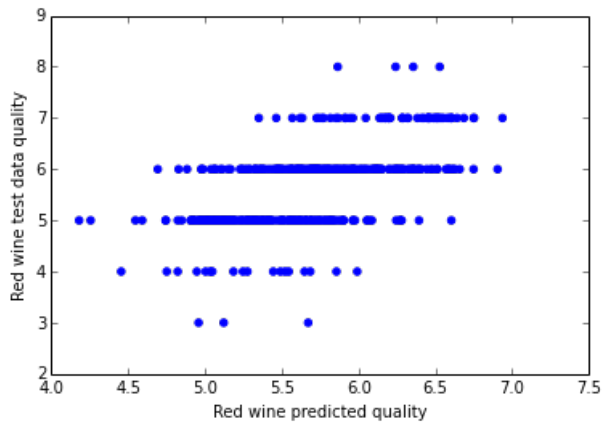
```
fitRegressionModelAndGetSquareError(True, 1)
```

Linear regression with minimised square loss. Mean Square Error = 0.413728282497

Out[67]:

Out[67]:

0.41372828249704313



4 (d) Benchmark Suggestion

4 (e) Benchmark Code

In [68]:

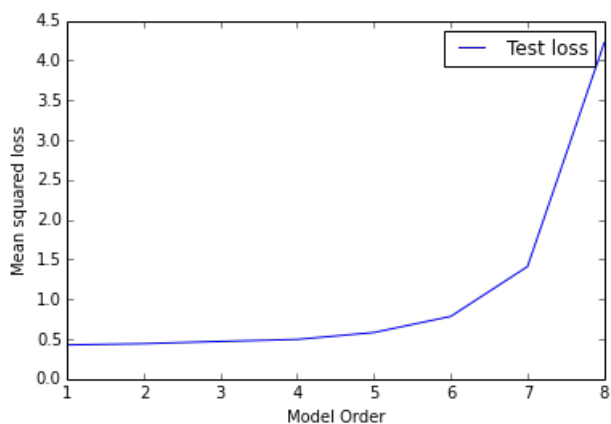
```
max_order = 7
runs = 100
error_runs = np.zeros((max_order+1,100))
errors = []
orders = [i+1 for i in range(max_order+1)]

for k in range(max_order+1):
    for i in range(100):
        err = fitRegressionModelAndGetSquareError(False, k+1)
        error_runs[k,i] = err

plt.figure()
plt.plot(np.array(orders), error_runs.mean(axis=1), 'b-', label="Test loss")
plt.legend()
plt.xlabel('Model Order')
plt.ylabel('Mean squared loss')
```

Out[68]:

<matplotlib.text.Text at 0x80cce10>



4 (f) Linear Regression Performance compared to the benchmark

5 (a) Implementing Regularised least square and plotting lambda parameter against the error.

In [69]:

```
import pylab as plt
from sklearn.linear_model import Ridge
```

```

from numpy.linalg import inv

#np.random.shuffle(data)
N = data.shape[0] #get tuple (numRows, numCols)

train = data[:int(N*0.7)]
test = data[int(N*0.7):]
X_train = train[:,1:]
X_train = np.c_[np.ones(train.shape[0]), X_train] # append 1s as first column
q_train = train[:,1:]

X_test = test[:,1:]
X_test = np.c_[np.ones(test.shape[0]), X_test]
q_test = test[:,1:]

lambdas = [0,0.01,0.02,0.04,0.08,0.16,0.32,0.64, 1.28]
errors = []
for lamb in lambdas:
    XtX = np.dot(X_train.T, X_train)
    XtXlam = XtX + X_train.shape[0]*lamb*np.identity(12)
    XtXlamI = inv(XtXlam)
    XtXIXt = np.dot(XtXlamI, X_train.T)
    w = np.dot(XtXIXt, q_train)

    f_test = np.dot(X_test,w)
    meanSquareError = ((q_test-f_test)**2).mean()/2.0
    errors += [meanSquareError]
    print "lambda", lamb, "Mean Square Error =", meanSquareError

plt.legend()
plt.xlabel('Lambda')
plt.ylabel('Mean squared loss')
plt.plot(lambdas, errors, '-o')

```

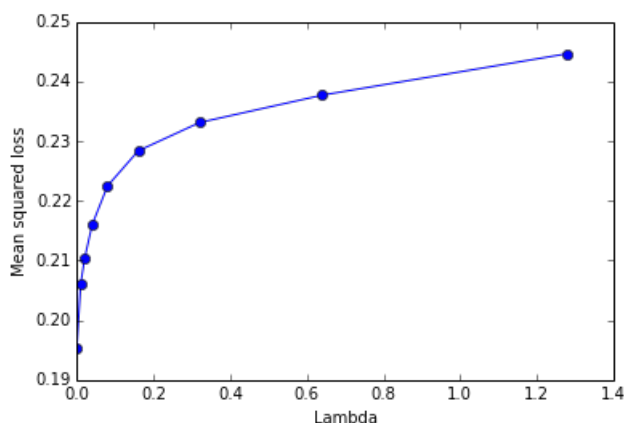
```

lambda 0 Mean Square Error = 0.195429756891
lambda 0.01 Mean Square Error = 0.206040917983
lambda 0.02 Mean Square Error = 0.210273456116
lambda 0.04 Mean Square Error = 0.216005524788
lambda 0.08 Mean Square Error = 0.222484022256
lambda 0.16 Mean Square Error = 0.228357543001
lambda 0.32 Mean Square Error = 0.233097168956
lambda 0.64 Mean Square Error = 0.237688160855
lambda 1.28 Mean Square Error = 0.244638129586

```

Out[69]:

[<matplotlib.lines.Line2D at 0x80e85f0>]



5 (b) Why this is not a good way to determine the value of lambda

TODO

5 (c) Implementation of 10-fold Cross Validation

In [70]:

```

N = train.shape[0]
K = 10

```

```

sizes = np.tile(np.floor(N/10), (1,K))
sizes[-1] = sizes[-1] + N - sizes.sum()
c_sizes = np.hstack((0,np.cumsum(sizes)))
X = np.copy(train[:,11])
X = np.c_[np.ones(train.shape[0]), X]
t = np.copy(train[:,11])
X_test = np.copy(test[:,11])
X_test = np.c_[np.ones(test.shape[0]), X_test]
t_test = np.copy(test[:,11])
lambdas = [0,0.01,0.02,0.04,0.08,0.16,0.32,0.64, 1.28]
cv_loss = np.zeros((K, len(lambdas)))
ind_loss = np.zeros((K, len(lambdas)))
train_loss = np.zeros((K, len(lambdas)))

k = 0
for lamb in lambdas:
    for fold in range(K):
        X_fold = X[c_sizes[fold]:c_sizes[fold+1],:]
        X_train = np.delete(X,np.arange(c_sizes[fold],c_sizes[fold+1],1),0)

        t_fold = t[c_sizes[fold]:c_sizes[fold+1]]
        t_train = np.delete(t,np.arange(c_sizes[fold],c_sizes[fold+1],1),0)
        XtX = np.dot(X_train.T, X_train)
        XtXlam = XtX + X_train.shape[0]*lamb*np.identity(12)
        XtXlamI = inv(XtXlam)
        XtXIXt = np.dot(XtXlamI, X_train.T)
        w = np.dot(XtXIXt, t_train)

        fold_pred = np.dot(X_fold,w)
        cv_loss[fold,k] = ((fold_pred - t_fold)**2).mean()/2.0
        ind_pred = np.dot(X_test,w)
        ind_loss[fold,k] = ((ind_pred - t_test)**2).mean()/2.0
        train_pred = np.dot(X_train,w)
        train_loss[fold,k] = ((train_pred - t_train)**2).mean()/2.0

    k += 1

print "Min cv_loss error is at lampda = 0. The error is =", cv_loss.mean(axis=0)[0]
print "Min train_loss error is at lampda = 0. The error is =", train_loss.mean(axis=0)[0]
print "Min ind_loss error is at lampda = 0. The error is =", ind_loss.mean(axis=0)[0]
plt.figure()
plt.plot(lambdas,train_loss.mean(axis=0),'b-',label="Training loss")
plt.plot(lambdas,cv_loss.mean(axis=0),'r-',label="CV loss")
plt.plot(lambdas,ind_loss.mean(axis=0),'k',label="Independent test loss")
plt.legend()
plt.xlabel('Lambda')
plt.ylabel('Mean squared loss')

```

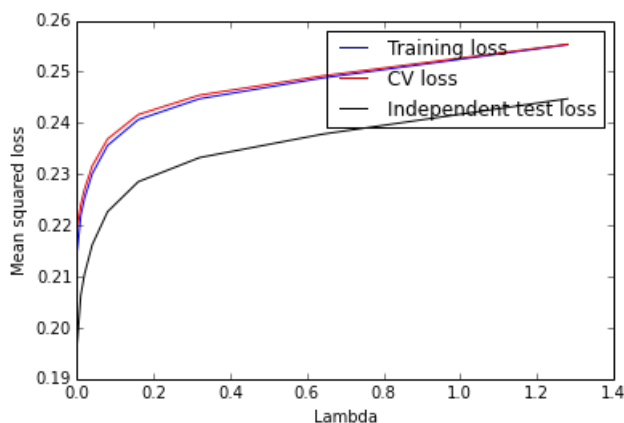
```

Min cv_loss error is at lampda = 0. The error is = 0.218976200163
Min train_loss error is at lampda = 0. The error is = 0.214178807735
Min ind_loss error is at lampda = 0. The error is = 0.195922649176

```

Out[70]:

<matplotlib.text.Text at 0x8c5f050>



5 (d) Compare the performance with the standard linear regression case.

TODO

6 Classification.

6 (a) Limitation of using regression

TODO

6 (b) Positive and negative features of KNN classifier with respect to our data

TODO

6 (c) Data pre-processing

TODO

6 (d) Implementing KNN classifier

In [71]:

```
import scipy.spatial.distance as ssd

# def read_data():
#     #read in red wine data
#     urllib.urlretrieve('https://archive.ics.uci.edu/ml/machine-learning-databases/wine-quality/winequality-red.csv', 'winequality-red.csv')
#     with open('winequality-red.csv') as f:
#         lines = (line for line in f)
#         data = np.loadtxt(lines, delimiter=';', skiprows=1)
#     return data

def knn(k, X_train, X_test, q_train):
    """ k-nearest neighbors """
    # initialize list to store predicted class
    pred_class = []
    # for each instance in data testing,
    # calculate distance in respect to data training
    for ii, di in enumerate(X_test):
        distances = [] # initialize list to store distance
        for ij, dj in enumerate(X_train):
            # calculate distances
            distances.append((ssd.euclidean(di,dj), ij))
        # k-neighbors
        k_nn = sorted(distances)[:k]
        # predict the class for the instance
        pred_class.append(classify(k_nn, q_train))
    # return prediction class
    return pred_class

def calc_dist(di,dj):
    return ssd.euclidean(di,dj) # built-in Euclidean fn

def evaluate(result):
    # create eval result array to store evaluation result
    eval_result = np.zeros(2,int)
    for x in result:
        # increment the correct prediction by 1
        if x == 0:
            eval_result[0] += 1
        # increment the wrong prediction by 1
        else:
            eval_result[1] += 1
    # return evaluation result
    return eval_result

def classify(k_nn, q_train):
    qlabel = []
    for dist, idx in k_nn:
        # retrieve label class and store into qlabel
        qlabel.append(q_train[idx])
    # return prediction class
    return np.argmax(np.bincount(qlabel))
```

```

def main():
    # read dataset of red wine
    #data = read_data()
    N = data.shape[0] #get tuple (numRows, numCols)
    np.random.shuffle(data)

    Nfolds = 10
    sizes = np.tile(np.floor(N/10), (1,Nfolds))
    sizes[-1] = sizes[-1] + N - sizes.sum()
    c_sizes = np.hstack((0,np.cumsum(sizes)))
    X = np.copy(data[:, :11]) # change to data here if you dont want to run the cv on independent test data
    t = np.copy(data[:, 11])

    # initialize K
    K = [1,3,7,11,19]
    cv_loss = np.zeros((Nfolds, len(K)))
    print "Started Cross Validation for 10 folds. This may take a few minutes..."
    for i in range(len(K)):
        for fold in range(Nfolds):
            X_fold = X[c_sizes[fold]:c_sizes[fold+1],:]
            X_train = np.delete(X,np.arange(c_sizes[fold],c_sizes[fold+1],1),0)

            t_fold = t[c_sizes[fold]:c_sizes[fold+1]]
            t_train = np.delete(t,np.arange(c_sizes[fold],c_sizes[fold+1],1),0)

            # predict the data test into class
            pred_class = knn(K[i], X_train, X_fold, t_train)
            # evaluate the predicted result
            eval_result = evaluate(pred_class-t_fold)
            # Calculate accuracy
            cv_loss[fold,i] = float(eval_result[1])/float(eval_result[0]+eval_result[1])
        print "Processing... K,",K[i]

    plt.plot(K,cv_loss.mean(axis=0), 'r-', label="CV results")
    plt.legend()
    plt.xlabel('K')
    plt.ylabel('Accuracy')
    return 3 #numpy.argmin(cv_loss, axis=None)

def confusionMatrix(optimalK):
    # read dataset of red wine
    #data = read_data()
    N = data.shape[0] #get tuple (numRows, numCols)
    np.random.shuffle(data)

    train = data[:int(N*0.7)]
    test = data[int(N*0.7):]
    X_train = train[:, :11]
    q_train = train[:, 11]

    X_test = test[:, :11]
    q_test = test[:, 11]

    confusion_matrix = np.zeros((6,6)) # map class 3 to 0, 4 to 1, 5 to 2, 6 to 3, 7 to 4, 8 to 5
    m = {3:0, 4:1, 5:2, 6:3, 7:4, 8:5}

    # predict the data test into class
    pred_class = knn(optimalK, X_train, X_test, q_train)
    # build the confusion matrix
    for p in range(len(pred_class)):
        confusion_matrix[m[pred_class[p]]][m[q_test[p]]] += 1.0

    print "Confusion Matrix"

    print " ",3," ",4," ",5," ",6," ",7," ",8
    for i in range(6):
        line = str(i+3) + " "
        for j in range(6):
            line += str(confusion_matrix[i][j]) + " "
        print line

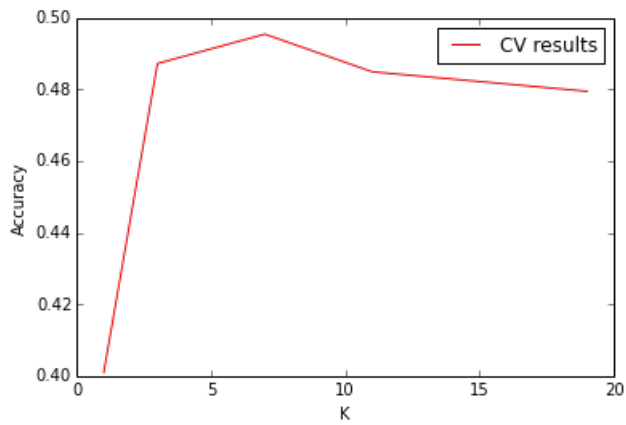
    optimalK = main()
    confusionMatrix(optimalK)

```

Started Cross Validation for 10 folds. This may take a few minutes...
Processing... K, 1

```
Processing... K, 3
Processing... K, 7
Processing... K, 11
Processing... K, 19
Confusion Matrix
```

	3	4	5	6	7	8
3	0.0	1.0	1.0	0.0	3.0	0.0
4	1.0	1.0	9.0	7.0	3.0	0.0
5	0.0	11.0	126.0	68.0	13.0	3.0
6	0.0	5.0	65.0	95.0	27.0	3.0
7	0.0	0.0	2.0	15.0	20.0	1.0
8	0.0	0.0	0.0	0.0	0.0	0.0



6 (e) Display the confusion matrix

6 (f) Discuss the performance of KNN

In []:

In []: