# PROJECT 1: PROBABILITY DISTRIBUTIONS AND BAYESIAN NETWORKS

Avinash Kommineni, 50248877

October 23, 2017

#### Introduction

The csv file is read and loaded into the workspace by the *read\_csvl* query of *pandas* library as a dataframe. Since there is no header and the values start right from row 0, the argument *header=None* is used. The given data is divided into training set, validation set and test set of sizes 80%, 10%, 10% respectively and stored into their respective X's and y's.

```
1 import random
 2 import numpy as np
 3 import pandas as pd
 4 from scipy.stats import multivariate_normal
 5 import matplotlib.pyplot as plt
 6 from sklearn import preprocessing, cluster, model_selection, ←
      metrics
 7
 8 synInputData = pd.read_csv('input.csv',header=None).values
9 synOutputData = pd.read_csv('output.csv',header=None).values
10 letorInputData = pd.read_csv('Querylevelnorm_X.csv',header=None). ←
  letorOutputData = pd.read_csv('Querylevelnorm_t.csv',header=None).↔
11
      values
12
13 # X_train, X_test, y_train, y_test = model_selection. ←
      train_test_split(synInputData, synOutputData, test_size=0.20, ←
      shuffle=False)
14 # X_validate, X_test, y_validate, y_test = model_selection. ←
      train_test_split(X_test, y_test, test_size=0.50, shuffle=False)
15
16 X_train, X_test, y_train, y_test = model_selection.train_test_split↔
       (letorInputData, letorOutputData, test_size=0.20, shuffle=False)
17 X_validate, X_test, y_validate, y_test = model_selection. ←
```

# **Design Matrix**

The design matrx,  $\Phi$  shown belowis calculated from

$$\phi_j(\mathbf{x}) = \exp\left(-\frac{1}{2} (x - \mu_i)^T \Sigma_j^{-1} (x - \mu_i)\right)$$

and substituted accordingly in the below form. Use of vector methods help code this pretty easily.

$$\Phi = \begin{bmatrix} \phi_0(\mathbf{x}_1) & \phi_1(\mathbf{x}_1) & \phi_2(\mathbf{x}_1) & \dots & \phi_{M-1}(\mathbf{x}_1) \\ \phi_0(\mathbf{x}_2) & \phi_1(\mathbf{x}_2) & \phi_2(\mathbf{x}_2) & \dots & \phi_{M-1}(\mathbf{x}_2) \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ \phi_0(\mathbf{x}_N) & \phi_1(\mathbf{x}_N) & \phi_2(\mathbf{x}_N) & \dots & \phi_{M-1}(\mathbf{x}_N) \end{bmatrix}$$

```
def compute_design_matrix(X_train, k_clusters):
 1
 2
       N,D = X_{train.shape}
 3
       kmeans = cluster.KMeans(k_clusters).fit(X_train)
 4
       centers = kmeans.cluster_centers_
 5
       centers = centers[:,np.newaxis,:]
 6
       spreads = []
 7
       covM = np.cov(X_train.T)
8
       for _ in range(0,k_clusters): spreads.append(covM)
9
       spreads = np.array(spreads)
       X = X_train[np.newaxis,:,:]
10
11
12
       basis_func_outputs = np.exp(np. sum(np.matmul(X - centers, ←
           spreads) * (X - centers), axis=2) / (-2) ).T
       return np.insert(basis_func_outputs, 0, 1, axis=1)
13
```

## **Closed Form Solution**

The closed form solution with the regularisation is calculated from

$$\mathbf{w}_{\mathrm{ML}} = \left(\lambda \mathbf{I} + \Phi^T \Phi\right)^{-1} \Phi^T t$$

The equation has been implemented in a effecient way by the use of broadcasting.

As shown, it takes the design matrix, truth values and the regularisation factor  $\lambda$ .

## **Stochastic Gradient Descent**

```
1 def SGD(learningRate, l2_lamda, designMatrix, outValues, ←
       miniBatchSize, numEpochs,k_clusters):
 2
       N=designMatrix.shape[0]
       weights = np.zeros([1, k_clusters+1])
 3
       for epoch in range(numEpochs):
 4
 5
       for i in range(int(N / miniBatchSize)):
 6
       lower_bound = i * miniBatchSize
 7
       upper_bound = min((i+1)*miniBatchSize, N)
       Phi = designMatrix[lower_bound : upper_bound, :]
 8
9
       t = outValues[lower_bound : upper_bound, :]
       E_D = np.matmul((np.matmul(Phi, weights.T)-t).T,Phi )
10
11
       E = (E D + 12 lamda * weights) / miniBatchSize
12
       weights = weights - learningRate * E
       return weights.flatten()
13
```

#### **Error - RMS**

The RMS error has been calculated from the given formula,

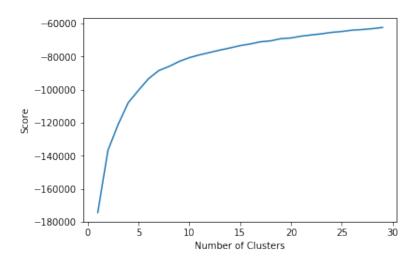
$$E_{RMS} = \sqrt{2E\left(w^*\right)/N_v}$$

where E(w) is defined as ...

$$E(w) = E_D(\mathbf{w}) + \frac{1}{2}\lambda E_W(\mathbf{w})$$

## **Results 1**

The hyper-parameter M needs to be set. I calculated it by plotting a graph of number of cluster vs scores.



From the above graph, I took a safe value of 10 as the value of M.

```
1 k_cluster = 10
2
3 12_lamda = 0.9
```

```
4
 5 designMatrix = compute_design_matrix(X_train,k_cluster)
 6 weights_train0 = closed_form_solution(12_lamda, designMatrix, \leftarrow
       y_train)
7 train_error0 = rms_error(weights_train0,designMatrix,y_train, ←
       12_lamda)
8 print("Training error:",train_error0)
9
10 designMatrix2 = compute_design_matrix(X_validate,k_cluster)
11 train_error2 = rms_error(weights_train0,designMatrix2,y_validate, ←
       12_lamda)
12 print("Test error (Validation set):",train_error2)
14
15 #Test error
16 designMatrix3 = compute_design_matrix(X_test,k_cluster)
17 train_error3 = rms_error(weights_train0,designMatrix3,y_test,←
       12_lamda)
18 print("Test error (test set):",train_error3)
```

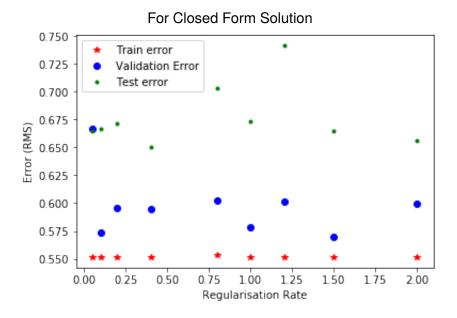
# **Code Output**

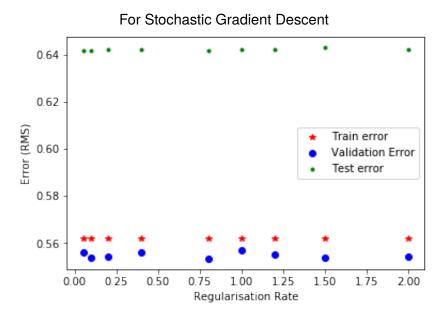
## Listing 1: Code output.

```
1 From closed form solution.
2 Training error: [[ 0.54948601]]
3 Test error (Validation set): [[ 0.64268395]]
4 Test error (test set): [[ 0.68269302]]
5 From SGD.
6 Training error: 0.560374755594
7 Test error (Validation set): 0.556667218125
8 Test error (test set): 0.639586215758
```

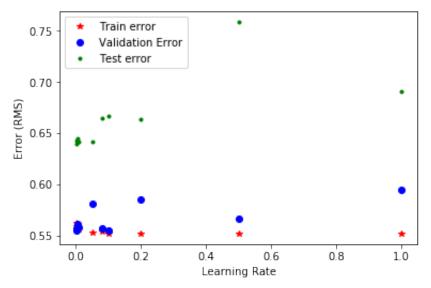
# **Results 2**

The following plots are drawn to better understand and decide the variation between the 3 errors and hyper-parameters such as regularization  $\lambda$  and learning rate  $\eta$ .

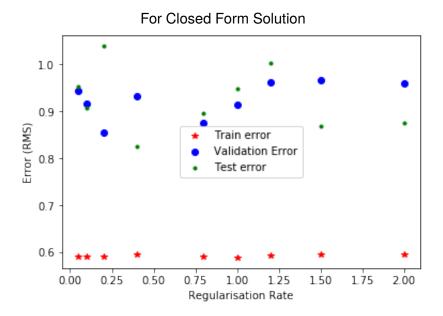


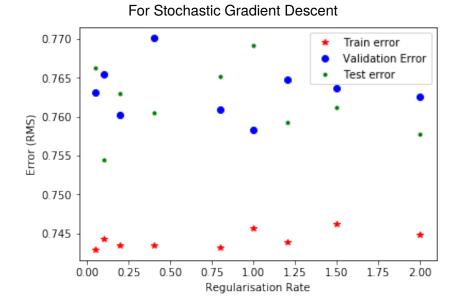


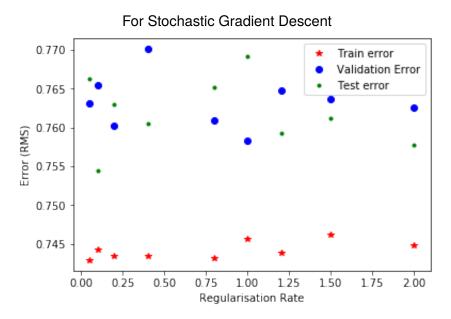




The below graphs are for the synthetic dataset.







- The stochastic graient descent appears to have a better error rate than closed form solution even though test error is high when compared to train and validation set error but has a little consistency.
- ullet The learning time increases significantly with the increase in  ${
  m M}$ , the number of basis functions.