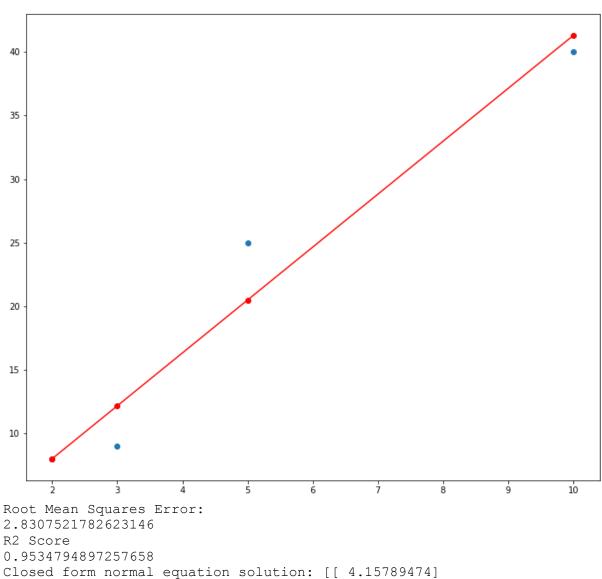
1.b) 
$$\theta = (x^T x)^{-1} x^T y = \begin{pmatrix} 1/38 & -5/38 \\ -5/38 & 69/76 \end{pmatrix} \begin{pmatrix} 568 \\ 82 \end{pmatrix} = \begin{pmatrix} 4.15789474 \\ -0.289473684 \end{pmatrix}$$

c) Gradient Descent is iterative in nature meaning it takes multiple steps to get global optimum but there is a case of linear regression where there is a way to get optimal values of parameters theta in single step. This is nothing but normal equation. Here, we get the best parameters with a formula that comprises of matrix multiplications and inversions.

d)

```
#Importing libraries
import pandas as pd
import matplotlib.pyplot as plt
import numpy as np
#Preprocessing the Data
x=[2,5,3,10]
y = [8, 25, 9, 40]
#Model building
mean x = np.mean(x)
mean_y = np.mean(y)
#total number of values
n = len(x)
#Calculating m and c
numer = 0
denom = 0
for i in range(n) :
  numer += (x[i] - mean_x) * (y[i] - mean_y)
  denom += (x[i] - mean x) ** 2
m = numer/denom
c = mean y - (m * mean x)
#coefficients
print("coefficients:")
print(m,c)
#Prediction
y_pred = m * np.asarray(x) + c
plt.scatter(x,y) #actual
plt.plot([min(x), max(x)], [min(y_pred), max(y_pred)], color ='red')
plt.scatter(x, y pred, color = 'red')
plt.show()
#Calculating RMSE
rmse = 0
```

```
for i in range(n):
 y pred = c + m * x[i]
  rmse += (y[i] - y_pred) ** 2
rmse = np.sqrt(rmse/n)
print("Root Mean Squares Error:")
print(rmse)
#Calculate R2 Score
ss tot = 0
ss res = 0
for i in range(n) :
 y pred = c + m * x[i]
ss tot += (y[i] - mean y) ** 2
 ss_res += (y[i] - y_pred) ** 2
r2 = 1 - (ss res/ ss tot)
print("R2 Score:")
print(r2)
#Closed form Normal Equation
a = np.asarray(x). shape[0]
x = np.append(np.asarray(x).reshape(-1,1),np.ones((a,1)),axis = 1)
x = np.array(x, dtype = 'int16')
y = np.array(y).reshape(-1,1)
theta = np.dot(np.linalg.inv(np.dot(x.T, x)), np.dot(x.T, y))
print("Closed form normal equation:", theta)
coefficients:
4.157894736842105 -0.28947368421052744
```



[-0.28947368]]

## 2)

```
#Importing libraries
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
import random
from mnist import MNIST
{\tt from \ sklearn.preprocessing \ import \ StandardScaler}
from scipy.linalg import eigh
#Loading MNIST Dataset
mndata = MNIST('C:\\Users\\anujeeth\\OneDrive\\Documents\\mnist')
training images, training labels = mndata.load training()
```

```
test images, test labels = mndata.load testing()
print("shape of training images data is : ", np.array(training images).
print("shape of training labels data is : ", np.array(training labels).
print("shape of test images data is : ", np.array(test images).shape)
print("shape of test labels data is : ", np.array(test_labels).shape)
#Extracting column labels from training set
label = training labels
ind = np.random.randint(0,20000)
plt.figure(figsize = (20,5))
grid data = np.array(pd.DataFrame(training images).iloc[ind]).reshape(2
8,28)
#Plotting
plt.imshow(grid data, interpolation = None, cmap = 'gray')
plt.show()
#Column standardization using standardScalar class. after column standa
rdization, the mean of every attribute becomes 0 and variance 1.
scalar = StandardScaler()
std df = scalar.fit transform(training images)
print ("Shape of the dataset after the column standardization:", std df.
shape)
#Finding the co-variance matrix
covar mat = np.matmul(std df.T, std df)
print("the dimensions of co-
variance matrix after multiplication", covar mat.shape)
#Finding the top two eigen-
values and corresponding eigen vectors. It generates only top 2 (782 an
d 783) eigen values. Converting eigen vectors into (2,d) form
values, vectors = eigh(covar mat, eigvals = (782,783))
print("Dimensions of eigen vector:", vectors.shape)
vectors= vectors.T
print("Dimensions of eigen vector:", vectors.shape)
#Finding PC1 and PC2
final df = np.matmul(vectors, stu df.T)
print("vectors:", vectors.shape, "n", "std_df:", std_df.T.shape,"n", "f
inal_df:", final_df.shape)
#Transposing final df
```

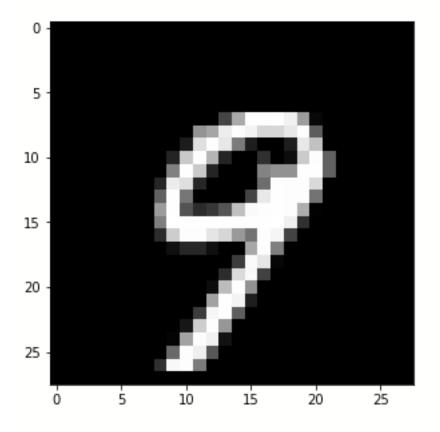
```
#Now converted 60000 * 784 data to 60000*4

final_dfT = np.vstack(final_df, label).T

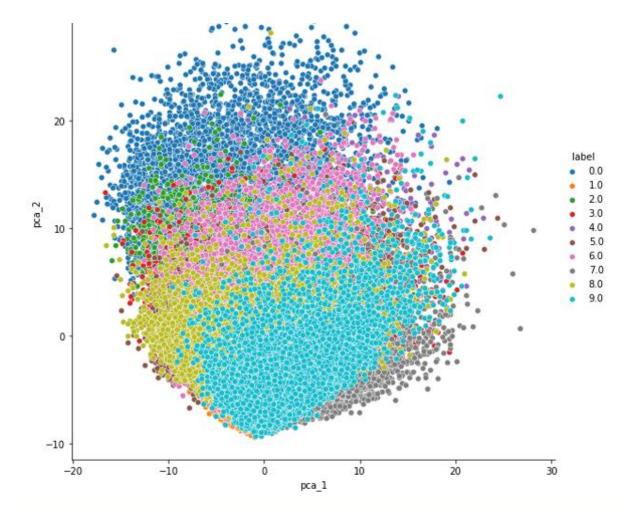
dataFrame = pd.DataFrame(final_dfT, columns = ['pca_1', 'pca_2', 'label
'])
print(dataFrame)

#Visualizing final data using seaborn Facet Grid

sns.FacetGrid(dataFrame, hue = 'label', size = 8)\
.map(sns.scatterplot, 'pca_1', 'pca_2')\
.add_legend()
plt.show()
```



	pca_1	pca_2	label
0	-4.814790	-0.922159	5.0
1	-7.754403	8.708977	0.0
2	9.431338	2.328389	4.0
3	-3.746318	-6.582173	1.0
4	3.133297	-5.183251	9.0
59995	-5.119129	-2.039339	8.0
59996	-6.498440	0.607841	3.0
59997	-3.230564	-3.777212	5.0
59998	-4.948125	1.722369	6.0
59999	-6.175386	-1.427251	8.0



## 2.a)

```
#Importing libraries
from mnist import MNIST
import numpy as np
import matplotlib.pyplot as plt
#Loading MNIST dataset
mnist = MNIST('C:\\Users\\anujeeth\\OneDrive\\Documents\\mnist')
X_train, y_train = mnist.load_training()
M = np.zeros(28, 28, 10)
S = np.zeros(28, 28, 10)
for i in range(9):
    X subset = X train[y train == i]
    M[:,:,i+1] = np.mean(X subset, axis = 0)
    S[:,:, i+1] = np.mean(X subset, axis = 0)
    plt.subplot(2,10,i+1)
    plt.imshow(M[:,:,i+1])
    plt.subplot(2,10,i+11)
    plt.imshow(S[:,:,i+1])
```

```
#Visualizing
plt.show()
```





3)

## #Importing libraries

```
from mnist import MNIST
import numpy as np
import matplotlib.pyplot as plt
#defining function
def pcs_svd(im_train, i):
    PC = np.dot(im train, im train.T)
    reconn = np.dot(PC, im train)
    normal = np.dot(reconn, PC)
    dif = np.sum(normal=im train)
    dif = dif/(60000*784)
    diff[i] = dif
    if(_name__ == '__main__'):
        im_train = np.loadtxt('C:\\Users\\anujeeth\\OneDrive\\Desktop\\
mnist\\t10k-images-idx3-ubyte')
        diff = np.zeros(784)
    for i in range (784)
    pcs svd(i train, i)
    plt.plot(diff)
    plt.show #Plotting
```