```
In [1]:
```

```
import pandas as pd
import matplotlib
from matplotlib import pyplot as plt
```

In [2]:

```
data = pd.read_csv("F:/assignments/Sem 6 Assignments/ML Assignment 1/Q1_1/Iris.csv")
```

In [3]:

```
data.columns
```

Out[3]:

In [4]:

data

Out[4]:

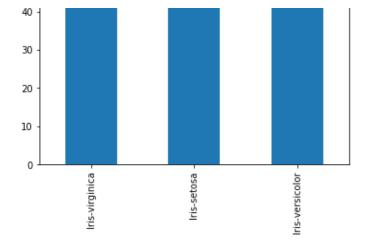
	ld	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm	Species
0	1	5.1	3.5	1.4	0.2	Iris-setosa
1	2	4.9	3.0	1.4	0.2	Iris-setosa
2	3	4.7	3.2	1.3	0.2	Iris-setosa
3	4	4.6	3.1	1.5	0.2	Iris-setosa
4	5	5.0	3.6	1.4	0.2	Iris-setosa
145	146	6.7	3.0	5.2	2.3	Iris-virginica
146	147	6.3	2.5	5.0	1.9	Iris-virginica
147	148	6.5	3.0	5.2	2.0	Iris-virginica
148	149	6.2	3.4	5.4	2.3	Iris-virginica
149	150	5.9	3.0	5.1	1.8	Iris-virginica

150 rows × 6 columns

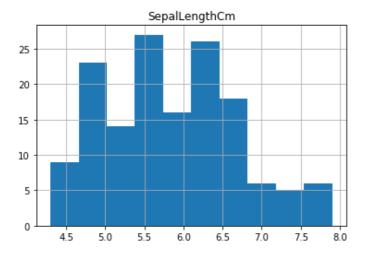
In [5]:

```
df2=pd.DataFrame(data)
df2['Species'].value_counts().plot(kind='bar')
plt.figure()
df2.hist(column='SepalLengthCm')
plt.figure()
df2.hist(column='SepalWidthCm')
plt.figure()
df2.hist(column='PetalLengthCm')
plt.figure()
df2.hist(column='PetalWidthCm')
```

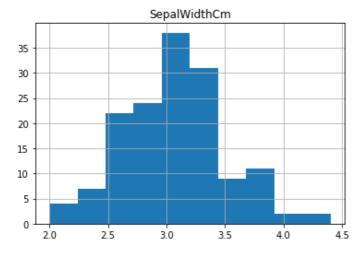
Out[5]:



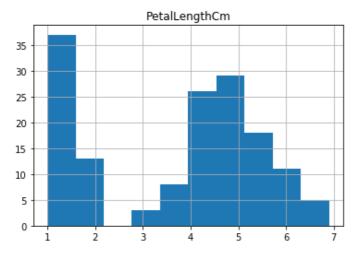
<Figure size 432x288 with 0 Axes>



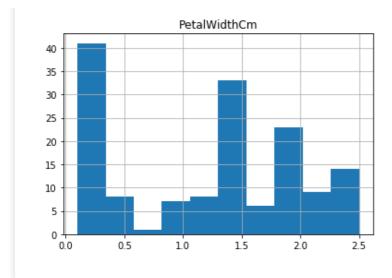
<Figure size 432x288 with 0 Axes>



<Figure size 432x288 with 0 Axes>



<Figure size 432x288 with 0 Axes>



In [1]:

```
import numpy as np
import idx2numpy
import random
from matplotlib import pyplot as plt
import cv2
```

In [2]:

```
train_images = idx2numpy.convert_from_file('train-images.idx3-ubyte')
train_labels = idx2numpy.convert_from_file('train-labels.idx1-ubyte')
test_images = idx2numpy.convert_from_file('t10k-images.idx3-ubyte')
test_labels = idx2numpy.convert_from_file('t10k-labels.idx1-ubyte')
```

In [3]:

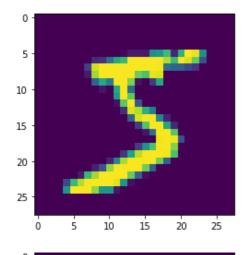
```
im1=train_images[0]
im2=train_images[56]
```

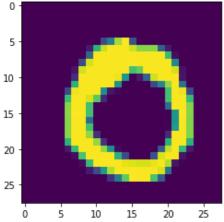
In [4]:

```
cv2.imwrite('color_img.jpg', im1)
plt.imshow(im1)
plt.figure()
cv2.imwrite('color_img.jpg', im2)
plt.imshow(im2)
```

Out[4]:

<matplotlib.image.AxesImage at 0x2473c0eaf88>





In [5]:

```
new_train=[]
for i in range(0,10):
    c=0
```

```
for j in range(0,len(train_labels)):
    if(train_labels[j]==i and c<=1000):
        new_train.append(train_images[j])
        c+=1</pre>
```

```
In [6]:
```

```
train_new=[]
#flatten images to a 1d array
for i in new_train:
    i=np.array(i)
    train_new.append(i.flatten())
train_images=train_new
```

Using TSNE for dimensionality reduction

```
In [7]:
```

```
from sklearn.manifold import TSNE
embedded=X_embedded = TSNE(n_components=2, init='random').fit_transform(train_images)
```

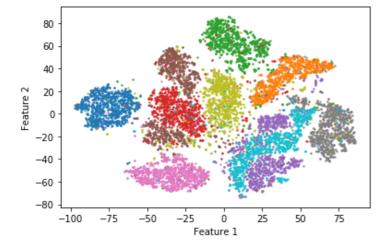
```
In [8]:
```

```
x=[]
y=[]
for i in embedded:
    x.append(i[0])
    y.append(i[1])
```

In [11]:

```
for i in range(0,10):
    x1=[]
    y1=[]
    for j in range(i*1000,(i+1)*1000):
        x1.append(x[j])
        y1.append(y[j])
    plt.scatter(x1,y1,s=1.5)

plt.xlabel('Feature 1')
plt.ylabel('Feature 2')
plt.show()
```



The Data is separable barring four classes that have seemed to mix up with each other (blue and purple, red and brown)

Linear Regression

```
In [1]:
```

```
import pandas as pd
import sklearn as sk
from sklearn import model_selection
from matplotlib import pyplot as plt
import numpy as np
from sklearn.metrics import r2_score
```

In [2]

data = pd.read_csv("F:/assignments/Sem 6 Assignments/ML Assignment 1/Q2/abalone.data")

Raw Data

In [3]:

data

Out[3]:

	Sex	Length	Diameter	Height	Whole Weight	Shucked weight	Viscera weight	Shell weight	Rings
0	М	0.455	0.365	0.095	0.5140	0.2245	0.1010	0.1500	15
1	М	0.350	0.265	0.090	0.2255	0.0995	0.0485	0.0700	7
2	F	0.530	0.420	0.135	0.6770	0.2565	0.1415	0.2100	9
3	M	0.440	0.365	0.125	0.5160	0.2155	0.1140	0.1550	10
4	I	0.330	0.255	0.080	0.2050	0.0895	0.0395	0.0550	7
4172	F	0.565	0.450	0.165	0.8870	0.3700	0.2390	0.2490	11
4173	M	0.590	0.440	0.135	0.9660	0.4390	0.2145	0.2605	10
4174	M	0.600	0.475	0.205	1.1760	0.5255	0.2875	0.3080	9
4175	F	0.625	0.485	0.150	1.0945	0.5310	0.2610	0.2960	10
4176	М	0.710	0.555	0.195	1.9485	0.9455	0.3765	0.4950	12

4177 rows × 9 columns

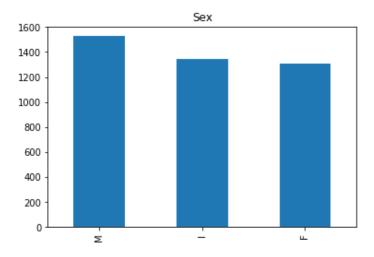
Visualization of dataset

```
In [4]:
```

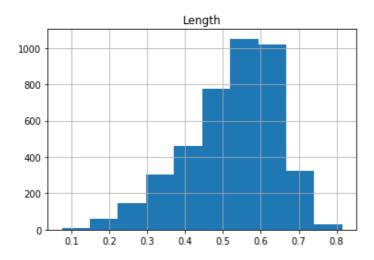
```
data['Sex'].value_counts().plot(kind='bar')
plt.title('Sex')
plt.figure()
data.hist(column='Length')
plt.figure()
data.hist(column='Diameter')
plt.figure()
data.hist(column='Height')
plt.figure()
data.hist(column='Whole Weight')
plt.figure()
data.hist(column='Shucked weight')
plt.figure()
data.hist(column='Shucked weight')
plt.figure()
data.hist(column='Viscera weight')
```

```
plt.figure()
data.hist(column='Shell weight')
```

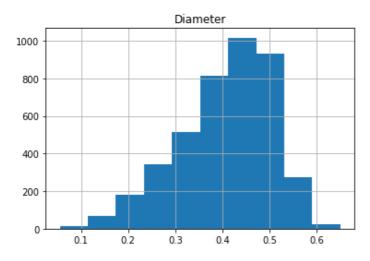
Out[4]:



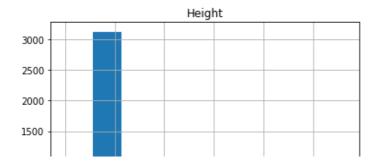
<Figure size 432x288 with 0 Axes>

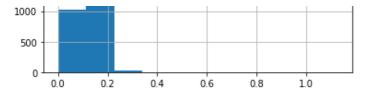


<Figure size 432x288 with 0 Axes>

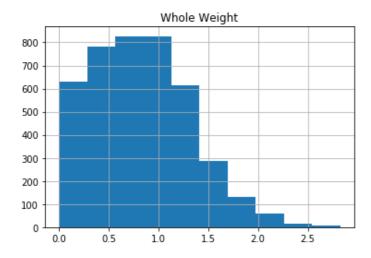


<Figure size 432x288 with 0 Axes>

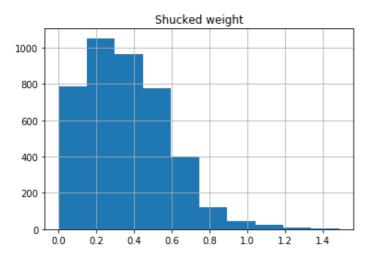




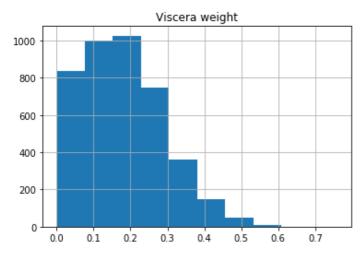
<Figure size 432x288 with 0 Axes>



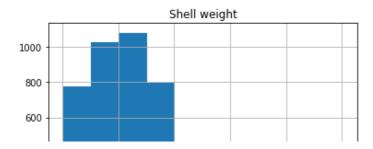
<Figure size 432x288 with 0 Axes>



<Figure size 432x288 with 0 Axes>



<Figure size 432x288 with 0 Axes>



```
200 0.0 0.2 0.4 0.6 0.8 1.0
```

In [5]:

```
#one hot encoding for Sex
one_hot = pd.get_dummies(data['Sex'])
data = data.drop('Sex',axis = 1)
data = data.join(one_hot)
```

Data after One Hot Encoding for Discrete Values (Sex)

In [6]:

data

Out[6]:

	Length	Diameter	Height	Whole Weight	Shucked weight	Viscera weight	Shell weight	Rings	F	ı	M
0	0.455	0.365	0.095	0.5140	0.2245	0.1010	0.1500	15	0	0	1
1	0.350	0.265	0.090	0.2255	0.0995	0.0485	0.0700	7	0	0	1
2	0.530	0.420	0.135	0.6770	0.2565	0.1415	0.2100	9	1	0	0
3	0.440	0.365	0.125	0.5160	0.2155	0.1140	0.1550	10	0	0	1
4	0.330	0.255	0.080	0.2050	0.0895	0.0395	0.0550	7	0	1	0
4172	0.565	0.450	0.165	0.8870	0.3700	0.2390	0.2490	11	1	0	0
4173	0.590	0.440	0.135	0.9660	0.4390	0.2145	0.2605	10	0	0	1
4174	0.600	0.475	0.205	1.1760	0.5255	0.2875	0.3080	9	0	0	1
4175	0.625	0.485	0.150	1.0945	0.5310	0.2610	0.2960	10	1	0	0
4176	0.710	0.555	0.195	1.9485	0.9455	0.3765	0.4950	12	0	0	1

4177 rows × 11 columns

Min-Max Scaling

```
In [7]:
```

```
#min max normalization
for column in data.columns:
    data[column] = (data[column] - data[column].min()) / (data[column] .max() - data[column] .min())
```

Splitting Dataset into 90% train 10% test

```
In [8]:
```

```
train_, test=model_selection.train_test_split(data, test_size=0.1, train_size=0.9)
```

Note: Training is done using the 5-fold validation. RMSE is reported on the validation set. Best Model from the 5 folds is determined using the RMSE value. Lower the RMSE, better is

tne model

Linear Regression w/o Regularization LR=0.00001

```
In [9]:
```

```
def linear(w,x_train,y_train,x_test,y_test):
    lr=0.00001
    x_train=np.array(x_train)
    y_train=np.array(y_train)
    x_test=np.array(x_test)
    y_test=np.array(y_test)

for i in range(x_train.shape[0]): #3007
    grad=0
    for j in range(x_train.shape[1]): #10
        grad=grad+(np.dot(x_train[i],w)-y_train[i])*x_train[i][j]
        w[j]=w[j]-lr*grad

#validation
    y_pred=np.dot(x_test,w)
    mse=np.sqrt(np.sum(np.square(y_pred-y_test)))/x_test.shape[0]
    return(w,mse)
```

In [10]:

```
best rmse=10000
kf=model selection.KFold(n splits=5)
# train=kf.get n splits(train )
# print(train .shape)
# print(y.shape)
y = train ['Rings']
new_train = train_.drop(['Rings'], axis=1)
# print(train .shape)
# print(y.shape)
for train_index, test_index in kf.split(train ):
     print("TRAIN:", train_index, "TEST:", test index)
    x train, x test = new train.iloc[train index], new train.iloc[test index]
    w=np.zeros((x train.shape[1],1))
     print(X train.shape)
     print(X test.shape)
   y_train, y_test = y.iloc[train index], y.iloc[test index]
     print(y test.shape)
    for i in range (50):
        w,mse=linear(w,x train,y train,x test,y test)
        rmse.append (mse)
    plt.plot(rmse)
    if (sum(rmse) <best_rmse):</pre>
        w best=w
        best rmse=sum(rmse)
plt.figure()
```

Out[10]:

<Figure size 432x288 with 0 Axes>



RMSE for model without penalty on test set

```
In [11]:
```

```
y=test['Rings']
new_test=test.drop(['Rings'],axis=1)
new_test=np.array(new_test)
y_pred=np.dot(new_test,w)
y=np.array(y)
# np.sqrt(np.sum(np.square((y-y_pred))))/y.size
# r2_score(y,y_pred)
np.sqrt(np.sum(np.square(y_pred-y)))/len(y)
```

Out[11]:

0.14695609463470916

L2 Regularization LR:0.00001 lambda=0.01

Notice the loss function here. The square of the norm of the weights of the model are added to the loss function.

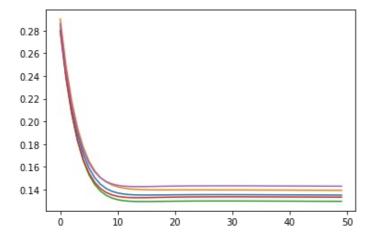
```
In [12]:
```

```
def 12(w,x_train,y_train,x_test,y_test):
   lr=0.00001
   x_train=np.array(x_train)
    y_train=np.array(y train)
   x test=np.array(x test)
    y test=np.array(y_test)
    lamb=0.0001
    for i in range(x train.shape[0]): #3007
        grad=0
        for j in range(x_train.shape[1]): #10
            grad=grad+(np.dot(x train[i],w)-y train[i]) *x train[i][j] + lamb*w[j]
            w[j]=w[j]-lr*grad
    #validation
    y pred=np.dot(x test,w)
   mse=(np.sqrt(np.sum(np.square(y pred-y test))) + lamb*np.square(np.linalg.norm(w)))
/x test.shape[0] #notice the change in loss function
   return (w, mse)
```

In [13]:

Out[13]:

<Figure size 432x288 with 0 Axes>



<Figure size 432x288 with 0 Axes>

RMSE for L2 on test set

```
In [14]:
```

```
y=test['Rings']
new_test=test.drop(['Rings'],axis=1)
new_test=np.array(new_test)
y_pred=np.dot(new_test,w_best_12)
y=np.array(y)
# np.sqrt(np.sum(np.square((y-y_pred))))/y.size
# r2_score(y,y_pred)
np.sqrt(np.sum(np.square(y_pred-y)))/len(y)
```

Out[14]:

0.14705283270813393

L1 Regularization LR:0.00001, lambda=0.001

Notice the loss function here. The absolute value of the weights of the model are added to the loss function

```
In [15]:
```

```
def l1(w,x_train,y_train,x_test,y_test):
    lr=0.00001
    x_train=np.array(x_train)
    y_train=np.array(y_train)
    x_test=np.array(x_test)
    y_test=np.array(y_test)
```

```
constant=0.5
lamb=0.001

for i in range(x_train.shape[0]): #3007
    grad=0
    for j in range(x_train.shape[1]): #10
        grad=grad+(np.dot(x_train[i],w)-y_train[i])*x_train[i][j] + lamb*constant
        w[j]=w[j]-lr*grad

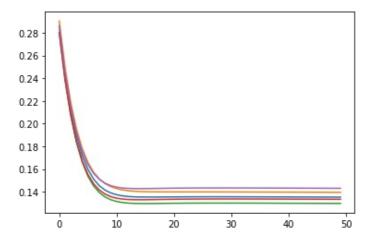
#validation
y_pred=np.dot(x_test,w)
mse=(np.sqrt(np.sum(np.square(y_pred-y_test))) + lamb*np.sum(np.abs(w)))/x_test.shap
e[0]#notice the change in loss function here
return(w,mse)
```

In [16]:

```
best rmse=10000
kf=model selection.KFold(n splits=5)
# train=kf.get_n_splits(train_)
# print(train .shape)
# print(y.shape)
y = train ['Rings']
new train = train .drop(['Rings'], axis=1)
# print(train .shape)
# print(y.shape)
for train index, test index in kf.split(train ):
    rmse=[]
     print("TRAIN:", train_index, "TEST:", test_index)
    x train, x test = new train.iloc[train index], new train.iloc[test index]
    wl1=np.zeros((x_train.shape[1],1))
     print(X train.shape)
     print(X test.shape)
    y train, y test = y.iloc[train index], y.iloc[test index]
     print(y test.shape)
    for i in range (50):
        wl1, mse=l1(wl1, x_train, y_train, x_test, y_test)
        rmse.append(mse)
    plt.plot(rmse)
    if (sum(rmse) < best rmse):</pre>
        w best l1=wl1
        best rmse=sum(rmse)
plt.figure()
```

Out[16]:

<Figure size 432x288 with 0 Axes>



<Figure size 432x288 with 0 Axes>

RMSE for L1 on test set

```
In [17]:

y=test['Rings']
new_test=test.drop(['Rings'],axis=1)
new_test=np.array(new_test)
y_pred=np.dot(new_test,w_best_l1)
y=np.array(y)
# np.sqrt(np.sum(np.square((y-y_pred))))/y.size
np.sqrt(np.sum(np.square(y_pred-y)))/len(y)

Out[17]:
0.1471399932030919
```

SkLearn implementation of Linear Regression no Penalty

In [18]:

```
from sklearn.linear model import LinearRegression
reg=LinearRegression()
kf=model selection.KFold(n splits=5)
# train=kf.get n splits(train )
# print(train .shape)
# print(y.shape)
y = train ['Rings']
new train = train .drop(['Rings'], axis=1)
# v1=test['Rings']
# new test=test.drop(['Rings'],axis=1)
# new test=np.array(new test)
best_rmse=10000
# print(train .shape)
# print(y.shape)
for train index, test index in kf.split(train ):
     print("TRAIN:", train index, "TEST:", test index)
   x train, x test = new train.iloc[train index], new train.iloc[test index]
     wl1=np.zeros((x train.shape[1],1))
     print(X train.shape)
     print(X test.shape)
   y train, y test = y.iloc[train index], y.iloc[test index]
     print(y test.shape)
    reg.fit(x_train,y_train)
   y pred=reg.predict(x test)
   rmse=np.sqrt(np.sum(np.square(y pred-y test)))/len(y pred)
    print(rmse)
    if(rmse<best rmse):</pre>
       best_rmse=rmse
       best model=reg
     print(reg.score(new test,y1))
y=test['Rings']
new_test=test.drop(['Rings'],axis=1)
```

RMSE for inbuilt Linear Regression on test set

```
In [19]:
```

O--- [101.

```
from sklearn.metrics import mean_squared_error

y=test['Rings']
new_test=test.drop(['Rings'],axis=1)
y_pred=best_model.predict(new_test)
rms = mean_squared_error(y, y_pred,squared=False)
rms
```

```
Out[19]:
0.08471284911075605
```

SkLearn Implementation of Ridge (L2)

```
In [20]:
```

```
from sklearn.linear model import Ridge
reg=Ridge(alpha=0.000001)
kf=model selection.KFold(n splits=5)
# train=kf.get_n_splits(train_)
# print(train_.shape)
# print(y.shape)
y = train ['Rings']
new train = train .drop(['Rings'], axis=1)
# y1=test['Rings']
# new test=test.drop(['Rings'],axis=1)
# new test=np.array(new_test)
best rmse=10000
# print(train_.shape)
# print(y.shape)
for train index, test index in kf.split(train ):
     rmse=[]
     print("TRAIN:", train index, "TEST:", test index)
   x train, x test = new train.iloc[train index], new train.iloc[test index]
     wl1=np.zeros((x train.shape[1],1))
#
    print(X_train.shape)
#
     print(X test.shape)
   y_train, y_test = y.iloc[train_index], y.iloc[test_index]
    print(y test.shape)
   reg.fit(x train, y train)
   y pred=reg.predict(x test)
   rmse=np.sqrt(np.sum(np.square(y pred-y test)))/len(y pred)
    print(rmse)
   if(rmse<best rmse):</pre>
       best rmse=rmse
       best model=reg
     print(reg.score(new test,y1))
y=test['Rings']
new test=test.drop(['Rings'],axis=1)
```

RMSE for Ridge on test set

```
In [21]:
```

```
from sklearn.metrics import mean_squared_error

y=test['Rings']
new_test=test.drop(['Rings'],axis=1)
y_pred=best_model.predict(new_test)
rms = mean_squared_error(y, y_pred,squared=False)
rms
```

0.08471285141421189

SkLearn Implementation of Lasso (L1)

```
In [22]:
```

Out[21]:

```
from sklearn.linear model import Lasso
```

```
reg=Lasso(alpha=0.00001)
kf=model selection.KFold(n splits=5)
# train=kf.get_n_splits(train_)
# print(train_.shape)
# print(y.shape)
y = train ['Rings']
new_train = train_.drop(['Rings'], axis=1)
# y1=test['Rings']
# new test=test.drop(['Rings'],axis=1)
# new test=np.array(new test)
best_rmse=10000
# print(train .shape)
# print(y.shape)
for train index, test index in kf.split(train ):
     print("TRAIN:", train index, "TEST:", test index)
   x train, x test = new train.iloc[train index], new train.iloc[test index]
     wl1=np.zeros((x_train.shape[1],1))
     print(X_train.shape)
#
      print(X test.shape)
#
   y train, y test = y.iloc[train index], y.iloc[test index]
     print(y test.shape)
   reg.fit(x train, y train)
   y pred=reg.predict(x test)
   rmse=np.sqrt(np.sum(np.square(y pred-y test)))/len(y pred)
     print(rmse)
   if(rmse<best rmse):</pre>
       best rmse=rmse
       best model=reg
     print(reg.score(new test,y1))
y=test['Rings']
new test=test.drop(['Rings'],axis=1)
```

RMSE for Lasso on test set

```
In [23]:
```

```
from sklearn.metrics import mean_squared_error

y=test['Rings']
new_test=test.drop(['Rings'],axis=1)
y_pred=best_model.predict(new_test)
rms = mean_squared_error(y, y_pred,squared=False)
rms
```

```
Out[23]:
0.08480590164862717
```

The introduction of penalty (L1 and L2) does not seem to have much effect on the RMSE value. After running multiple runs of this code, no conclusive best model can be found. Sometimes L1 performs the best, sometimes L2 and sometimes no penalty is the best model. The RMSE values are very close to each other.

The inbuilt regression models give slightly lower RMSE (about 0.06)

Closed Form RMSE on validation set

```
In [24]:
```

```
kf=model selection.KFold(n splits=5)
```

```
y = train_['Rings']
new train = train .drop(['Rings'], axis=1)
# y1=test['Rings']
# new_test=test.drop(['Rings'],axis=1)
# new test=np.array(new test)
idx=1
for train index, test index in kf.split(train ):
    x train, x test = new train.iloc[train index], new train.iloc[test index]
    y train, y test = y.iloc[train index], y.iloc[test index]
    w=np.dot(np.linalg.inv(np.dot(np.transpose(x train), x train)), np.dot(np.transpose(x
train),y train))
    y_pred=np.dot(x_test,w)
   y_test=np.array(y_test)
   print("RMSE of fold "+str(idx)+":", np.sqrt(np.sum(np.square(y pred-y test)))/len(y t
est))
   idx+=1
RMSE of fold 1: 0.0027954060476873955
RMSE of fold 2: 0.003147262374071904
RMSE of fold 3: 0.002681020281461619
RMSE of fold 4: 0.0026932049414070255
RMSE of fold 5: 0.0029880871585471888
```

Logistic Regression on Ionosphere Dataset

```
In [1]:
```

```
import pandas as pd
import sklearn as sk
from sklearn import model_selection
from matplotlib import pyplot as plt
import numpy as np
from sklearn.metrics import r2_score
from sklearn.linear_model import LogisticRegression
```

In [2]:

```
data = pd.read_csv("F:/assignments/Sem 6 Assignments/ML Assignment 1/Q3/ionosphere_data_k
aggle.csv")
data
```

Out[2]:

	feature1	feature2	feature3	feature4	feature5	feature6	feature7	feature8	feature9	feature10	 feature26	feature27
0	1	0	0.99539	0.05889	0.85243	0.02306	0.83398	0.37708	1.00000	0.03760	 -0.51171	0.41078
1	1	0	1.00000	- 0.18829	0.93035	- 0.36156	- 0.10868	- 0.93597	1.00000	-0.04549	 -0.26569	-0.20468
2	1	0	1.00000	- 0.03365	1.00000	0.00485	1.00000	- 0.12062	0.88965	0.01198	 -0.40220	0.58984
3	1	0	1.00000	- 0.45161	1.00000	1.00000	0.71216	1.00000	0.00000	0.00000	 0.90695	0.51613
4	1	0	1.00000	- 0.02401	0.94140	0.06531	0.92106	- 0.23255	0.77152	-0.16399	 -0.65158	0.13290
346	1	0	0.83508	0.08298	0.73739	- 0.14706	0.84349	- 0.05567	0.90441	-0.04622	 -0.04202	0.83479
347	1	0	0.95113	0.00419	0.95183	0.02723	0.93438	- 0.01920	0.94590	0.01606	 0.01361	0.93522
348	1	0	0.94701	0.00034	0.93207	0.03227	0.95177	- 0.03431	0.95584	0.02446	 0.03193	0.92489
349	1	0	0.90608	- 0.01657	0.98122	- 0.01989	0.95691	0.03646	0.85746	0.00110	 -0.02099	0.89147
350	1	0	0.84710	0.13533	0.73638	- 0.06151	0.87873	0.08260	0.88928	-0.09139	 -0.15114	0.81147

Replacing g and b with 1 and 0 in the label column

```
In [3]:
```

351 rows × 35 columns

```
data['label'].replace({'g':1,'b':0},inplace=True)
data
```

Out[3]:

	feature1	feature2	feature3	feature4	feature5	feature6	feature7	feature8	feature9	feature10	 feature26	feature27
0	1	0	0.99539	0.05889	0.85243	0.02306	0.83398	0.37708	1.00000	0.03760	 -0.51171	0.41078

_	feature1	feature2		feature4		feature6	feature7	feature8		feature10	 feature26	feature27
1	i	0	1.00000	0.18829	0.93035	0.36156	0.10868	0.93597	1.00000	-0.04549	 -0.26569	-0.20468
2	1	0	1.00000	0.03365	1.00000	0.00485	1.00000	- 0.12062	0.88965	0.01198	 -0.40220	0.58984
3	1	0	1.00000	- 0.45161	1.00000	1.00000	0.71216	1.00000	0.00000	0.00000	 0.90695	0.51613
4	1	0	1.00000	0.02401	0.94140	0.06531	0.92106	0.23255	0.77152	-0.16399	 -0.65158	0.13290
			•••					•••			 	
346	1	0	0.83508	0.08298	0.73739	- 0.14706	0.84349	- 0.05567	0.90441	-0.04622	 -0.04202	0.83479
347	1	0	0.95113	0.00419	0.95183	- 0.02723	0.93438	- 0.01920	0.94590	0.01606	 0.01361	0.93522
348	1	0	0.94701	0.00034	0.93207	- 0.03227	0.95177	- 0.03431	0.95584	0.02446	 0.03193	0.92489
349	1	0	0.90608	- 0.01657	0.98122	- 0.01989	0.95691	0.03646	0.85746	0.00110	 -0.02099	0.89147
350	1	0	0.84710	0.13533	0.73638	- 0.06151	0.87873	0.08260	0.88928	-0.09139	 -0.15114	0.81147

351 rows × 35 columns

The state of the s

Splitting Dataset into 90% train 10% test

```
In [4]:
```

```
train_,test=model_selection.train_test_split(data, test_size=0.1, train_size=0.9)
```

In [5]:

```
stats=pd.DataFrame()
stats["mean"]=train_.mean()
stats["Var"]=train_.var()
stats
```

Out[5]:

	mean	Var
feature1	0.892063	0.096593
feature2	0.000000	0.000000
feature3	0.642893	0.238592
feature4	0.051843	0.192843
feature5	0.608174	0.254543
feature6	0.123541	0.213319
feature7	0.555684	0.226843
feature8	0.118215	0.278598
feature9	0.501287	0.260881
feature10	0.189137	0.244225
feature11	0.471601	0.309997
feature12	0.171377	0.239788
feature13	0.395721	0.388738
feature14	0.110508	0.245293
feature15	0.333867	0.430329

```
feature16 0.075898 0.210811
feature17 0.366644 0.385899
feature18 0.004513 0.249589
feature19 0.348317 0.394385
feature20 -0.008207 0.274522
feature21 0.336343 0.364187
feature22 0.024012 0.267117
feature23 0.361513 0.361545
feature24 -0.051329 0.277417
feature25 0.378971 0.336360
feature26 -0.058471 0.261215
feature27 0.540448 0.263351
feature28 -0.065516 0.304881
feature29 0.370661 0.326723
feature30 -0.005855 0.253571
feature31 0.340722 0.326127
feature32 -0.000377 0.264168
feature33 0.354652 0.264820
feature34 0.014154 0.222551
    label 0.638095 0.231665
```

In [6]:

```
stats['Var'].nlargest(5)
```

Out[6]:

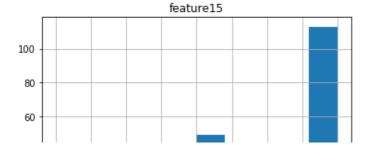
```
feature15 0.430329
feature19 0.394385
feature13 0.388738
feature17 0.385899
feature21 0.364187
Name: Var, dtype: float64
```

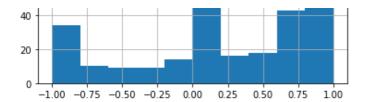
Plotting Histograms of the Features with the highest variances (top 5)

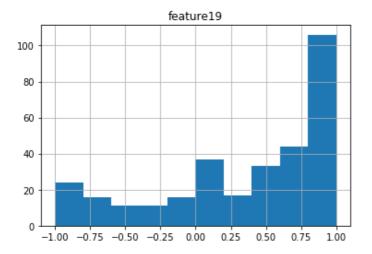
In [7]:

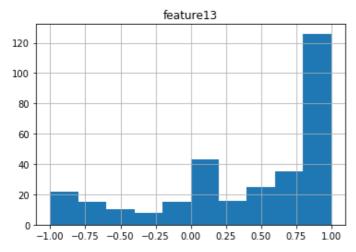
```
train_.hist(column='feature15')
train_.hist(column='feature19')
train_.hist(column='feature13')
train_.hist(column='feature17')
train_.hist(column='feature21')
```

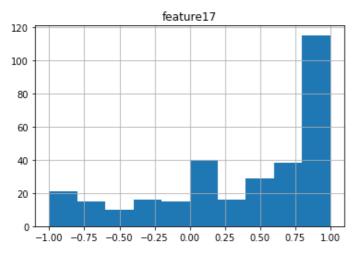
Out[7]:

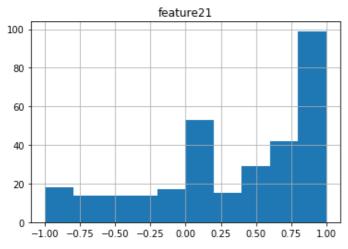












```
In [8]:
```

```
reg=LogisticRegression(penalty='none', max iter=1000)
kf=model selection.KFold(n splits=5)
# train=kf.get_n_splits(train_)
# print(train_.shape)
# print(y.shape)
y1=test['label']
new test=test.drop(['label'],axis=1)
new test=np.array(new test)
from sklearn.metrics import precision recall fscore support
from sklearn.metrics import accuracy score
y = train ['label']
new train = train .drop(['label'], axis=1)
# print(train_.shape)
# print(y.shape)
stats1=[]
for train_index, test_index in kf.split(train_):
    rmse=[]
     print("TRAIN:", train index, "TEST:", test index)
   x_train, x_test = new_train.iloc[train_index], new train.iloc[test index]
   y_train, y_test = y.iloc[train_index], y.iloc[test_index]
    print(y test.shape)
   reg.fit(x train, y train)
   ac=accuracy score(reg.predict(new test), y1)
   stats1.append(list(precision recall fscore support(reg.predict(new test), y1, averag
e='micro')))
   stats1[-1].append(ac)
# y=test['label']
# new test=test.drop(['label'],axis=1)
# new_test=np.array(new_test)
# from sklearn.metrics import precision recall fscore support
stats1=pd.DataFrame(stats1)
stats1.columns =['Precision', 'Recall', 'F1-Score', 'Support','Accuracy']
stats1=stats1.drop(['Support'],axis=1)
print ("Logistic Regression Stats without PCA")
print(stats1)
Logistic Regression Stats without PCA
  Precision Recall F1-Score Accuracy
   0.861111 0.861111 0.861111 0.861111
```

```
Precision Recall F1-Score Accuracy
0 0.861111 0.861111 0.861111
1 0.861111 0.861111 0.861111
2 0.861111 0.861111 0.861111
3 0.888889 0.888889 0.888889
4 0.861111 0.861111 0.861111 0.861111
```

PCA for dimensionality reduction

```
In [9]:
```

```
from sklearn.decomposition import PCA
kf=model_selection.KFold(n_splits=5)
# train=kf.get_n_splits(train_)
# print(train_.shape)
# print(y.shape)
for val in range(90,100,1):
    pca = PCA(val/100)
    yl=test['label']
    new_test=test.drop(['label'],axis=1)
    new_test=np.array(new_test)
    # print(new_test.shape)
    y = train_['label']
    new_train = train_.drop(['label'], axis=1)
# print(train_.shape)
# print(y.shape)
```

```
stats1=[]
    ind=0
    for train index, test index in kf.split(train ):
       x train, x test = new train.iloc[train index], new train.iloc[test index]
       y train, y test = y.iloc[train index], y.iloc[test index]
         print(y test.shape)
         components=pca.fit transform(x train)
         pca.fit(x train)
         x train= pca.transform(x train)
    #
         pca.fit(x train)
         x train= pca.transform(x train)
       pca.fit(x train)
       x_train= pca.transform(x train)
       new test1=pca.transform(new test)
         ind=10
    #
         y train=y train.values.reshape(-1, 1)
        y_test=y_test.values.reshape(-1, 1)
        print(y_test.shape)
    #
         y_train=pca.transform(y_train)
         y test=pca.transform(y test)
       reg=LogisticRegression(penalty='none', max_iter=1000)
       reg.fit(x_train,y_train)
        ac=accuracy score(reg.predict(new test1), y1)
       stats1.append(list(precision recall fscore support(reg.predict(new test1), y1, a
verage='micro')))
       stats1[-1].append(ac)
    # y=test['label']
    # new test=test.drop(['label'],axis=1)
    # new test=np.array(new test)
    # from sklearn.metrics import precision recall fscore support
    stats1=pd.DataFrame(stats1)
    stats1.columns =['Precision', 'Recall', 'F1-Score', 'Support','Accuracy']
    stats1=stats1.drop(['Support'],axis=1)
    print("Logistic Regression Stats with PCA with variance", val/100)
    print(stats1)
   print()
Logistic Regression Stats with PCA with variance 0.9
  Precision Recall F1-Score Accuracy
   0.861111 0.861111
                      0.861111
                                0.861111
   0.861111 0.861111 0.861111 0.861111
   0.861111 0.861111 0.861111 0.861111
3
   0.888889 0.888889 0.888889
   0.888889 0.888889 0.888889 0.888889
Logistic Regression Stats with PCA with variance 0.91
  Precision Recall F1-Score Accuracy
0
  0.861111 0.861111 0.861111 0.861111
1
   0.861111 0.861111 0.861111 0.861111
   0.861111 0.861111 0.861111 0.861111
   0.888889 0.888889 0.888889 0.888889
   0.888889 0.888889 0.888889 0.888889
Logistic Regression Stats with PCA with variance 0.92
  Precision Recall F1-Score Accuracy
   0.916667 0.916667 0.916667 0.916667
\cap
   0.861111 0.861111 0.861111 0.861111
1
   0.861111 0.861111
                      0.861111 0.861111
   0.888889 0.888889
3
                      0.888889 0.888889
   0.916667 0.916667 0.916667 0.916667
Logistic Regression Stats with PCA with variance 0.93
  Precision Recall F1-Score Accuracy
\cap
  0.888889 0.888889 0.888889 0.888889
1
   0.861111 0.861111 0.861111 0.861111
   0.861111 0.861111 0.861111 0.861111
3
   0.916667 0.916667 0.916667 0.916667
   0.916667 0.916667 0.916667 0.916667
```

```
Logistic Regression Stats with PCA with variance 0.94
  Precision Recall F1-Score Accuracy
  0.916667 0.916667 0.916667 0.916667
   0.888889 0.888889 0.888889 0.888889
  0.888889 0.888889 0.888889 0.888889
  0.916667 0.916667 0.916667 0.916667
   0.916667 0.916667 0.916667 0.916667
Logistic Regression Stats with PCA with variance 0.95
  Precision Recall F1-Score Accuracy
  0.916667 0.916667 0.916667 0.916667
 0.916667 0.916667 0.916667 0.916667
2 0.888889 0.888889 0.888889
3 0.944444 0.944444 0.944444 0.944444
4 0.916667 0.916667 0.916667 0.916667
Logistic Regression Stats with PCA with variance 0.96
  Precision Recall F1-Score Accuracy
  0.916667 0.916667 0.916667 0.916667
  0.916667 0.916667 0.916667 0.916667
  0.888889 0.888889 0.888889 0.888889
  0.916667 0.916667 0.916667 0.916667
  0.916667 0.916667 0.916667 0.916667
Logistic Regression Stats with PCA with variance 0.97
  Precision Recall F1-Score Accuracy 0.916667 0.916667 0.916667 0.916667
1
   0.888889 0.888889 0.888889 0.888889
  0.888889 0.888889 0.888889 0.888889
3 0.916667 0.916667 0.916667 0.916667
4 0.916667 0.916667 0.916667 0.916667
Logistic Regression Stats with PCA with variance 0.98
  Precision Recall F1-Score Accuracy
  0.888889 0.888889 0.888889 0.888889
 0.888889 0.888889 0.888889 0.888889
 0.916667 0.916667 0.916667 0.916667
  0.916667 0.916667 0.916667 0.916667
  0.888889 0.888889 0.888889 0.888889
Logistic Regression Stats with PCA with variance 0.99
  Precision Recall F1-Score Accuracy
  0.888889 0.888889 0.888889 0.888889
```

The best model among the 5 folds is determined using the accuracy, higher the accuracy, better is the model. Again, F1 Score/Accuracy is first reported on the validation set and testing is done on the 10% test dataset

Logistic Regression w/o penalty

```
In [10]:
```

```
reg=LogisticRegression(penalty='none', max_iter=1000)
kf=model_selection.KFold(n_splits=5)
# train=kf.get_n_splits(train_)
# print(train_.shape)
# print(y.shape)
# y1=test['label']
# new_test=test.drop(['label'], axis=1)
# new_test=np.array(new_test)
from sklearn.metrics import precision_recall_fscore_support
```

```
from sklearn.metrics import accuracy_score
y = train ['label']
new_train = train_.drop(['label'], axis=1)
# print(train .shape)
# print(y.shape)
stats1=[]
best accuracy=0
for train index, test index in kf.split(train ):
    rmse=[]
      print("TRAIN:", train index, "TEST:", test index)
    x train, x test = new train.iloc[train index], new train.iloc[test index]
    y train, y test = y.iloc[train index], y.iloc[test index]
     print(y test.shape)
    reg.fit(x train, y train)
    ac=accuracy score(reg.predict(x test), y test)
    stats1.append(list(precision recall fscore support(reg.predict(x test), y test, aver
age='micro')))
    stats1[-1].append(ac)
    if(ac>best accuracy):
        best model nopenalty=reg
        best accuracy=ac
# y=test['label']
# new test=test.drop(['label'],axis=1)
# new test=np.array(new test)
# from sklearn.metrics import precision recall fscore support
stats1=pd.DataFrame(stats1)
stats1.columns =['Precision', 'Recall', 'F1-Score', 'Support','Accuracy']
stats1=stats1.drop(['Support'],axis=1)
print("Logistic Regression Stats without Penalty")
print(stats1)
Logistic Regression Stats without Penalty
              Recall F1-Score Accuracy
  Precision
   0.841270 0.841270 0.841270 0.841270
   0.873016 0.873016 0.873016 0.873016
1
   0.888889 0.888889 0.888889 0.888889
   0.936508 0.936508 0.936508 0.936508
3
   0.888889 0.888889 0.888889 0.888889
In [11]:
y=test['label']
y numpy=np.array(y)
new_test=test.drop(['label'],axis=1)
new test=np.array(new test)
ac=accuracy score(best model nopenalty.predict(new test), y)
fscore=precision recall fscore support (best model nopenalty.predict(new test), y, averag
e='micro')
print("Testing Best Model with no Penalty")
print("Accuracy=", ac)
print("Precision=", fscore[0])
print("Recall=", fscore[1])
print("F1-Score=", fscore[2])
Testing Best Model with no Penalty
```

```
Logistic Regression with L1 penalty
```

Accuracy= 0.8611111111111112 Precision= 0.8611111111111112 Recall= 0.861111111111112 F1-Score= 0.8611111111111112

```
In [12]:
    reg=LogisticRegression(penalty='11' may iter=1000 solve)
```

```
reg=LogisticRegression(penalty='l1', max_iter=1000, solver='liblinear')
kf=model_selection.KFold(n_splits=5)
```

```
# train=kf.get_n_splits(train_)
# print(train_.shape)
# print(y.shape)
# y1=test['label']
# new test=test.drop(['label'],axis=1)
# new test=np.array(new test)
from sklearn.metrics import precision recall fscore support
from sklearn.metrics import accuracy score
y = train ['label']
new train = train .drop(['label'], axis=1)
# print(train .shape)
# print(y.shape)
stats1=[]
best accuracy=0
for train index, test index in kf.split(train ):
     print("TRAIN:", train index, "TEST:", test index)
    x_train, x_test = new_train.iloc[train_index], new_train.iloc[test_index]
   y_train, y_test = y.iloc[train_index], y.iloc[test index]
     print(y test.shape)
   reg.fit(x_train,y_train)
    ac=accuracy score(reg.predict(x test), y test)
   stats1.append(list(precision recall_fscore_support(reg.predict(x_test), y_test, aver
age='micro')))
    stats1[-1].append(ac)
    if(ac>best accuracy):
        best_model l1=reg
       best accuracy=ac
# y=test['label']
# new test=test.drop(['label'],axis=1)
# new test=np.array(new test)
# from sklearn.metrics import precision_recall_fscore_support
stats1=pd.DataFrame(stats1)
stats1.columns =['Precision', 'Recall', 'F1-Score', 'Support','Accuracy']
stats1=stats1.drop(['Support'],axis=1)
print("Logistic Regression Stats L1")
print(stats1)
Logistic Regression Stats L1
  Precision Recall F1-Score Accuracy
   0.873016 0.873016 0.873016 0.873016
   0.857143 0.857143 0.857143 0.857143
1
   0.888889 0.888889 0.888889
3
   0.904762 0.904762 0.904762 0.904762
   0.841270 0.841270 0.841270 0.841270
In [13]:
y=test['label']
y numpy=np.array(y)
new test=test.drop(['label'],axis=1)
new test=np.array(new test)
ac=accuracy_score(best_model_l1.predict(new_test), y)
fscore=precision recall fscore support(best model 11.predict(new test), y, average='micr
0')
print("Testing Best Model with L1 Penalty")
print("Accuracy=", ac)
print("Precision=", fscore[0])
print("Recall=", fscore[1])
print("F1-Score=", fscore[2])
Testing Best Model with L1 Penalty
```

Logistic Regression with L2

```
In [14]:
reg=LogisticRegression(penalty='12', max iter=1000, solver='liblinear')
kf=model selection.KFold(n splits=5)
# train=kf.get n splits(train )
# print(train .shape)
# print(y.shape)
# y1=test['label']
# new test=test.drop(['label'],axis=1)
# new test=np.array(new test)
from sklearn.metrics import precision recall fscore support
from sklearn.metrics import accuracy_score
y = train ['label']
new_train = train_.drop(['label'], axis=1)
# print(train .shape)
# print(y.shape)
stats1=[]
best accuracy=0
for train index, test index in kf.split(train ):
   rmse=[]
      print("TRAIN:", train index, "TEST:", test index)
   x train, x test = new train.iloc[train index], new train.iloc[test index]
    y_train, y_test = y.iloc[train_index], y.iloc[test index]
     print(y test.shape)
    reg.fit(x train, y train)
    ac=accuracy score(reg.predict(x test), y test)
    stats1.append(list(precision recall fscore support(reg.predict(x test), y test, aver
age='micro')))
    stats1[-1].append(ac)
    if(ac>best accuracy):
       best model 12=reg
        best accuracy=ac
         print("YAY")
# y=test['label']
# new test=test.drop(['label'],axis=1)
# new test=np.array(new test)
# from sklearn.metrics import precision recall fscore support
stats1=pd.DataFrame(stats1)
stats1.columns =['Precision', 'Recall', 'F1-Score', 'Support', 'Accuracy']
stats1=stats1.drop(['Support'],axis=1)
print("Logistic Regression Stats L2")
print(stats1)
Logistic Regression Stats L2
  Precision Recall F1-Score Accuracy
   0.841270 0.841270 0.841270 0.841270
   0.841270 0.841270 0.841270 0.841270
1
   0.873016 0.873016 0.873016 0.873016
2
   0.888889 0.888889 0.888889 0.888889
3
   0.873016 0.873016 0.873016 0.873016
In [15]:
y=test['label']
y numpy=np.array(y)
new test=test.drop(['label'],axis=1)
new test=np.array(new test)
ac=accuracy score(best model 12.predict(new test), y)
fscore=precision recall fscore support(best model 12.predict(new test), y, average='micr
print("Testing Best Model with L2 Penalty")
print("Accuracy=", ac)
print("Precision=", fscore[0])
print("Recall=", fscore[1])
print("F1-Score=", fscore[2])
```

Introduction of L1 and L2 penalty has improved the accuracy/F1 Score.

```
In [16]:
```

```
print(best model nopenalty)
print(best model 11)
print(best model 12)
LogisticRegression(C=1.0, class weight=None, dual=False, fit intercept=True,
                   intercept scaling=1, 11 ratio=None, max iter=1000,
                   multi class='auto', n jobs=None, penalty='none',
                   random state=None, solver='lbfgs', tol=0.0001, verbose=0,
                   warm start=False)
LogisticRegression(C=1.0, class weight=None, dual=False, fit intercept=True,
                   intercept scaling=1, 11 ratio=None, max iter=1000,
                   multi_class='auto', n_jobs=None, penalty='11',
                   random state=None, solver='liblinear', tol=0.0001, verbose=0,
                   warm start=False)
LogisticRegression(C=1.0, class weight=None, dual=False, fit intercept=True,
                   intercept_scaling=1, l1_ratio=None, max_iter=1000,
                   multi_class='auto', n_jobs=None, penalty='12',
                   random state=None, solver='liblinear', tol=0.0001, verbose=0,
                   warm start=False)
```

Plotting ROC-AUC curve for the three best models

In [17]:

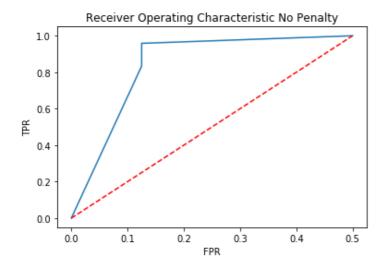
```
y=test['label']
y numpy=np.array(y)
new test=test.drop(['label'],axis=1)
new test=np.array(new test)
probab 12=best model 12.predict proba(new test)
probab l1=best model l1.predict proba(new test)
probab no=best model nopenalty.predict proba(new test)
#second column is 1 first column is 0
curve no x=[]
curve no y=[]
curve 11 x=[]
curve 11 y=[]
curve 12 x=[]
curve_12 y=[]
# print(y)
for i in range (0,110,10):
   threshold=i/100
   predicted=[]
     print(threshold)
    for j in probab no:
       if (j[1]>threshold):
           predicted.append(1)
           predicted.append(0)
     print(predicted)
    tpr=0
    fpr=0
    for j in range(len(predicted)):
        if (predicted[j] == y_numpy[j] and y_numpy[j] == 1):
            tpr+=1
        if (predicted[j]!=y numpy[j] and predicted[j]==1):
            fpr+=1
    tpr=tpr/sum(y numpy)
```

```
fpr=fpr/sum(y_numpy)
    curve_no_x.append(fpr)
    curve_no_y.append(tpr)

plt.plot(curve_no_x,curve_no_y)
plt.plot([0, max(curve_no_x)], [0, max(curve_no_y)],'r--')
plt.title('Receiver Operating Characteristic No Penalty')
plt.ylabel('TPR')
plt.xlabel('FPR')
```

Out[17]:

Text(0.5, 0, 'FPR')



In [18]:

```
for i in range (0,110,10):
    threshold=i/100
    predicted=[]
      print(threshold)
    for j in probab_l1:
        if(j[1]>threshold):
            predicted.append(1)
        else:
            predicted.append(0)
      print(predicted)
    tpr=0
    fpr=0
    for j in range(len(predicted)):
        if(predicted[j]==y_numpy[j] and predicted[j]==1):
        if (predicted[j]!=y_numpy[j] and predicted[j]==1):
            fpr+=1
    tpr=tpr/sum(y_numpy)
    fpr=fpr/sum(y_numpy)
    curve_l1_x.append(fpr)
    curve_l1_y.append(tpr)
plt.plot(curve_l1_x,curve_l1_y)
plt.plot([0, max(curve_l1_x)], [0, max(curve_l1_y)], 'r--')
plt.title('Receiver Operating Characteristic L1')
plt.ylabel('TPR')
plt.xlabel('FPR')
```

Out[18]:

Text(0.5, 0, 'FPR')



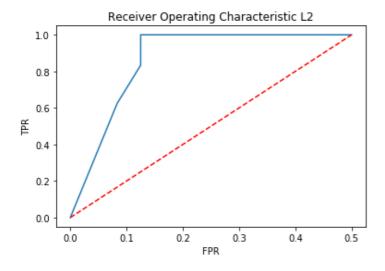
```
0.4 - 0.2 - 0.3 0.4 0.5 EPR
```

In [19]:

```
for i in range (0,110,10):
    threshold=i/100
    predicted=[]
      print(threshold)
    for j in probab 12:
        if(j[1]>threshold):
            predicted.append(1)
        else:
            predicted.append(0)
      print (predicted)
    tpr=0
    fpr=0
    for j in range(len(predicted)):
        if (predicted[j] == y_numpy[j] and y_numpy[j] == 1):
        if(predicted[j]!=y_numpy[j] and predicted[j]==1):
            fpr+=1
    tpr=tpr/sum(y_numpy)
    fpr=fpr/sum(y_numpy)
    curve 12 x.append(fpr)
    curve_12_y.append(tpr)
plt.plot(curve_12_x,curve_12_y)
plt.plot([0, max(curve_12_x)], [0, max(curve_12_y)], 'r--')
plt.title('Receiver Operating Characteristic L2')
plt.ylabel('TPR')
plt.xlabel('FPR')
```

Out[19]:

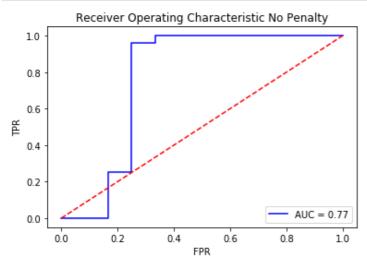
Text(0.5, 0, 'FPR')



In [20]:

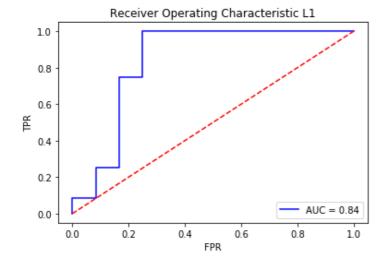
```
import sklearn.metrics as metrics
preds = probab_no[:,1]
fpr, tpr, threshold = metrics.roc_curve(y_numpy, preds)
roc_auc = metrics.auc(fpr, tpr)
plt.title('Receiver Operating Characteristic No Penalty')
plt.plot(fpr, tpr, 'b', label = 'AUC = %0.2f' % roc_auc)
plt.legend(loc = 'lower right')
plt.plot([0, 1], [0, 1], 'r--')
plt.ylabel('TPR')
```

```
plt.xlabel('FPR')
plt.show()
```



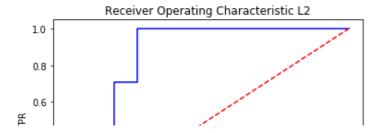
In [21]:

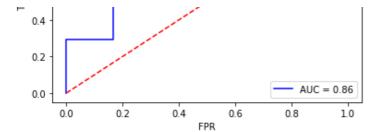
```
preds = probab_l1[:,1]
fpr, tpr, threshold = metrics.roc_curve(y_numpy, preds)
roc_auc = metrics.auc(fpr, tpr)
plt.title('Receiver Operating Characteristic L1')
plt.plot(fpr, tpr, 'b', label = 'AUC = %0.2f' % roc_auc)
plt.legend(loc = 'lower right')
plt.plot([0, 1], [0, 1], 'r--')
plt.ylabel('TPR')
plt.xlabel('FPR')
plt.show()
```



In [22]:

```
preds = probab_l2[:,1]
fpr, tpr, threshold = metrics.roc_curve(y_numpy, preds)
roc_auc = metrics.auc(fpr, tpr)
plt.title('Receiver Operating Characteristic L2')
plt.plot(fpr, tpr, 'b', label = 'AUC = %0.2f' % roc_auc)
plt.legend(loc = 'lower right')
plt.plot([0, 1], [0, 1], 'r--')
plt.ylabel('TPR')
plt.xlabel('FPR')
plt.show()
```





The difference between the inbuilt and implemented versions of the ROC-AUC curve is the resolution. I have implemented the curve for only 11 points of threshold, which is much lesser than the inbuilt version. Increasing the resolution results in the plots being similar.

```
In [1]:
```

```
import numpy as np
import idx2numpy
import random
from matplotlib import pyplot as plt
import cv2
import numpy as np
from sklearn.multiclass import OneVsOneClassifier
from sklearn.svm import SVC
from sklearn.metrics import precision_recall_fscore_support
from sklearn.metrics import accuracy_score
import pandas as pd
from sklearn.linear_model import LogisticRegression
```

In [2]:

```
train_images = idx2numpy.convert_from_file('train-images.idx3-ubyte')
train_labels = idx2numpy.convert_from_file('train-labels.idx1-ubyte')
test_images = idx2numpy.convert_from_file('t10k-images.idx3-ubyte')
test_labels = idx2numpy.convert_from_file('t10k-labels.idx1-ubyte')
train_new=[]
test_new=[]
for i in train_images:
    i=np.array(i)
    train_new.append(i.flatten())
for i in test_images:
    i=np.array(i)
    test_new.append(i.flatten())
train_images=train_new
test_images=test_new
```

In [3]:

```
clf = OneVsOneClassifier(LogisticRegression(random state=0, max iter=1000)).fit(train imag
es, train labels)
# clf = OneVsRestClassifier(SVC()).fit(train images, train labels)
F:\Anaconda3\lib\site-packages\sklearn\linear model\ logistic.py:940: ConvergenceWarning:
lbfgs failed to converge (status=1):
STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.
Increase the number of iterations (max iter) or scale the data as shown in:
    https://scikit-learn.org/stable/modules/preprocessing.html
Please also refer to the documentation for alternative solver options:
    https://scikit-learn.org/stable/modules/linear model.html#logistic-regression
  extra warning msg= LOGISTIC SOLVER CONVERGENCE MSG)
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```

```
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Please also refer to the documentation for alternative solver options:
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  extra warning msg= LOGISTIC SOLVER CONVERGENCE MSG)
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lbfgs failed to converge (status=1):
STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.
```

```
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```

```
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```

```
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```

In [6]:

```
ac=accuracy_score(clf.predict(test_images), test_labels)
print("Accuracy=",ac)
fscore=precision_recall_fscore_support(clf.predict(test_images), test_labels, average='m
icro')
print("Precision=",fscore[0])
print("Recall=",fscore[1])
print("F1-Score=",fscore[2])
```

Accuracy= 0.9249 Precision= 0.9249 Recall= 0.9249 F1-Score= 0.9249

```
In [1]:
```

```
import numpy as np
import idx2numpy
import random
from matplotlib import pyplot as plt
import cv2
import numpy as np
from sklearn.multiclass import OneVsRestClassifier
from sklearn.svm import SVC
from sklearn.metrics import precision_recall_fscore_support
from sklearn.metrics import accuracy_score
import pandas as pd
from sklearn.linear_model import LogisticRegression
```

In [2]:

```
train_images = idx2numpy.convert_from_file('train-images.idx3-ubyte')
train_labels = idx2numpy.convert_from_file('train-labels.idx1-ubyte')
test_images = idx2numpy.convert_from_file('t10k-images.idx3-ubyte')
test_labels = idx2numpy.convert_from_file('t10k-labels.idx1-ubyte')
train_new=[]
test_new=[]
for i in train_images:
    i=np.array(i)
    train_new.append(i.flatten())
for i in test_images:
    i=np.array(i)
    test_new.append(i.flatten())
train_images=train_new
test_images=test_new
```

In [3]:

```
clf = OneVsRestClassifier(LogisticRegression(random state=0, max iter=1000)).fit(train ima
ges, train labels)
# clf = OneVsRestClassifier(SVC()).fit(train images, train labels)
F:\Anaconda3\lib\site-packages\sklearn\linear model\ logistic.py:940: ConvergenceWarning:
lbfgs failed to converge (status=1):
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icro')
print("Precision=",fscore[0])
print("Recall=",fscore[1])
print("F1-Score=",fscore[2])
```

Recall= 0.9168 F1-Score= 0.9168

Λ_{-} \wedge Λ Λ Λ Λ Λ Λ Λ
Hyush Madhan Johini
2019156
Ayush Madhan Sohini 2019156 M2-Assignment 1
M/- Herigmmet. 1

Y -> Vector of predicted values X -> Vector of i/p features E > cross value

The loss function needed to be minimized >

L(O) = I & (y (xi) - xio)^2 -> Mean Squared

2N i=1 (ons

xi ERd -> it sample from date set of size N

In rectar form ->

 $L(\theta) = L \quad (Y - X\theta)^2$ $2N \quad \text{fredicted} \quad \text{bothel vector}$

 $=\frac{1}{2N}\left(Y-XO\right)^{T}\left(Y-XO\right)$

= 1 (YT-(XO)) (Y-XO)

$$= \frac{1}{2N} \left(Y^{\dagger} - 0^{T} X^{\dagger} \right) \left(Y - X \theta \right)$$

$$= \int_{2N} \left(Y^{\dagger} Y - Y^{\dagger} X \theta - \theta^{\dagger} X^{\dagger} Y + \theta^{\dagger} X^{\dagger} X \theta \right)$$

$$\frac{\partial L(0)}{\partial o} = \frac{1}{2N} \left[-X^{T}Y - X^{T}Y + 2X^{T}XO \right]$$

$$= \frac{1}{2N} \left[2x^{T}x \sigma - 2x^{T}y \right]$$

To more in the direction of the optimal solution, re equate the derivative to zero

$$\frac{1}{2N} \left[2x^7xo - 2x^7y \right] = 0$$

$$X^{T} \times O = X^{T} Y$$

$$\Theta = (x^T x)^{-1} x^T Y$$

Whenever the invose of X^TX exists, the closed form solution exists. 21 X^TX is a singular mediax, the closed form solution won't wist.

The closed four solution is a better option ONLY then
the size of the i/p makix X is small on X is
speake. Hen X is a very large (suppose A has 10⁵
entrie), X^T X rould be a 10⁵ × 10⁵ makix ic
A has 10¹⁰ entries, which would be very difficult
to stare. Also performing (X^TX) is also
computationally inefficient on such a large matrix.

Also if X^TX is singular, the imase doesn't east to anyways. In such cases, the iterative methods (the
gradient descent are a better choice.

 $y(x^{i}) = \theta_{0} + x_{1}^{i}\theta_{1} + x_{2}^{i}\theta_{2} - ... x_{n}^{i}\theta_{n} + \varepsilon^{i}$ $y(x^{i}) = \theta_{0} + x_{1}^{i}\theta_{1} + n_{2}^{i}\theta_{2} - ... x_{n}^{i}\theta_{n} + \varepsilon^{i}$

 $y(x^m) = 00 + x_1^m \theta_1 \dots x_n^m \theta_n + \epsilon^m$

m

$$y(x') + \dots y(x^m) = m\theta_0 + \theta_1 \xi_2,$$

$$i=1$$

$$+ \theta_2 \xi_2 + \dots$$

$$i=1$$

$$+ \theta_n \xi_n + \xi_i$$

$$i=1$$

$$i=1$$

$$+ \theta_n \xi_n + \xi_i$$

$$i=1$$

Diricling both sides by m

$$y(n) + \dots y(n) = \theta_0 + \theta_1 \underbrace{\sum_{i=1}^{m} x_i^i}_{ci}$$

$$\dots + \theta_n \underbrace{\sum_{i=1}^{m} + \sum_{i=1}^{m} i}_{m}$$

$$= 0 \text{ as } e \text{ follows } a$$

$$\text{Zero mean gaussian}$$

$$O = \left[\begin{array}{c} O_1 & O_2 & \dots & O_n \end{array}\right]$$

$$\frac{1}{y} = \left[\begin{array}{c} y & Cn' \end{array}\right]$$

$$\vdots & \times 1$$

$$\vdots & m$$

$$y & Cn'' \end{array}$$

Cincar Regression model throws a continuous output. So if we can set a particular threshold is if the off the linear negrenson is about a threshold then it belongs to class A clse class B, it can work as a binary classifier. The thrushold palues can be determined using tools like the ROC-AVC curre. Moreover the data is rarely distributed as a gaussian, so this is not a good classifier as in linear negression the croos in the data is assumed to be a gaussian.