Slip 1

Q1.Write a R program to add, multiply and divide two vectors of integertype. (Vector length should be minimum 4) [10 Marks]

```
vector1 = seq(10,40 , length.out=4)
vector2 = c(20, 10, 40, 40)
print("Original Vectors:")
add= vector1+vector2
cat("Sum of vector is ",add, "\n")
sub_vector= vector1-vector2
cat("Substraction of vector is ",sub_vector, "\n")
mul_vector= vector1 * vector2
cat("Multiplication of vector is ",mul_vector, "\n")
print("Division of two Vectors:")
div_vector = vector1 / vector2
print(div_vector)
```

Q2.Consider the student data set. It can be downloaded from: https://drive.google.com/open?id=1oakZCv7g3mlmCSdv9J8kdSaqO 5_6dIOw . Write a programme in python to apply simple linear regression and find out mean absolute error, mean squared error and root mean squared error. [20 Marks]

```
# Importing the libraries
import numpy as np
import pandas as pd
from sklearn.linear_model import LinearRegression
from sklearn.metrics import mean_absolute_error
from sklearn.metrics import mean_squared_error
from sklearn.metrics import r2_score

# Loading the dataset
data = pd.read_csv('Student_score.csv')

# Separating the features and target variable
X = data.iloc[:, 0:1].values
y = data.iloc[:, 0:1].values

# Fitting the Linear Regression model to the dataset
```

```
regressor = LinearRegression()
regressor.fit(X, y)

# Predicting the results
y_pred = regressor.predict(X)

# Calculating the errors
MAE = mean_absolute_error(y, y_pred)
MSE = mean_squared_error(y, y_pred)
RMSE = np.sqrt(MSE)

# Printing the errors
print("Mean Absolute Error:", MAE)
print("Mean Squared Error:", MSE)
print("Root Mean Squared Error:", RMSE)
```

Slip 2 :

Q1. Write an R program to calculate the multiplication table using afunction. [10 Marks]

```
table<-function(number)
{
for( t in 1:10)
 {
 print ( paste ( number, '*', t, '=', number * t))
 }
}
table(9)
```

Q2. Write a python program to implementt k-means algorithms on a synthetic dataset.

```
# importing necessary libraries

import matplotlib.pyplot as plt
import numpy as np
from sklearn.datasets import make_blobs

# generating random data points
x, y = make_blobs(n_samples=100, centers=3, n_features=2)

# plotting the generated data points
plt.scatter(x[:, 0], x[:, 1], c=y, cmap='gist_rainbow')
plt.show()

# implementing K-Means
k = 3

# assigning random centers
center = 10*np.random.rand(k, 2)
```

```
# computing the distance matrix
dist_matrix = np.zeros((100, 3))

for i in range(k):
    dist_matrix[:, i] = np.sum((x-center[i, :])**2, axis=1)

# assigning labels to each data point
cluster_labels = np.argmin(dist_matrix, axis=1)

# plotting the labeled data points
plt.scatter(x[:, 0], x[:, 1], c=cluster_labels, cmap='gist_rainbow')
plt.show()

# updating the cluster centers
for i in range(k):
    center[i, :] = np.mean(x[cluster_labels == i, :], axis=0)

# recomputing the distance matrix
dist_matrix = np.zeros((100, 3))

for i in range(k):
    dist_matrix[:, i] = np.sum((x-center[i, :])**2, axis=1)

# reassigning labels to each data point
cluster_labels = np.argmin(dist_matrix, axis=1)

# plotting the labeled data points
plt.scatter(x[:, 0], x[:, 1], c=cluster_labels, cmap='gist_rainbow')
plt.show()
```

Slip 3:

Q1. Write a R program to reverse a number and also calculate the sum ofdigits of that number.

```
n=123
Reverse=function(n)
{
sum=0
rev=0
while(n>0)
{
r=n%%10
sum=sum+r
rev=rev*10+r
n=n%/%10
}
```

```
print(sum)
print(rev)
}
Reverse(n)
```

Q2. Consider the following observations/data. And apply simple linear regression and find out estimated coefficients b0 and b1.(use numpypackage) x=[0,1,2,3,4,5,6,7,8,9,11,13] y = ([1, 3, 2, 5, 7, 8, 8, 9, 10, 12,16, 18]

```
import numpy as np
import matplotlib.pyplot as plt

def estimate_coef(x, y):
   # number of observations/points
 n = np.size(x)

 # mean of x and y vector
 m_x = np.mean(x)
 m_y = np.mean(y)

 # calculating cross-deviation and deviation about x
 SS_xy = np.sum(y*x) - n*m_y*m_x
 SS_xx = np.sum(x*x) - n*m_x*m_x

 # calculating regression coefficients
 b_1 = SS_xy / SS_xx
 b_0 = m_y - b_1*m_x

 return (b_0, b_1)

def plot_regression_line(x, y, b):
 # plotting the actual points as scatter plot
 plt.scatter(x, y, color = "m",
 marker = "o", s = 30)

 # predicted response vector
 y_pred = b[0] + b[1]*x

 # plotting the regression line
 plt.plot(x, y_pred, color = "g")

 # putting labels
 plt.xlabel('x')
 plt.ylabel('y')

 # function to show plot
```

```
 plt.show()
def main():
 # observations / data
 x = np.array([0, 1, 2, 3, 4, 5, 6, 7, 8, 9,11,13])
 y = np.array([1, 3, 2, 5, 7, 8, 8, 9, 10, 12,16, 18])

 # estimating coefficients
 b = estimate_coef(x, y)
 print("Estimated coefficients:\nb_0 = {} \ \nb_1 = {}".format(b[0], b[1]))

 # plotting regression line
 plot_regression_line(x, y, b)

if __name__ == "__main__":
    main()
```

Slip 4:

Q1. Write a R program to calculate the sum of two matrices of given size.

```
matrix1<-matrix(c(1,2,3,4,5,6),nrow=2)
print(matrix1)
matrix2<-matrix(c(7,8,9,10,11,12),nrow=2)
print(matrix2)
result<-matrix1+matrix2
cat("Addition : ","\n")
print(result)
```

Q2. Consider following dataset
weather=['Sunny','Sunny','Overcast','Rainy','Rainy','Rainy','Overcast','Sunny','Sunny','Rainy','Sunn y','Overcast','Overcast','Rainy']
temp=['Hot','Hot','Hot','Mild','Cool','Cool','Cool','Mild','Cool','Mild','Mild','Mild','Hot','Mild']
play=['No','No','Yes','Yes','Yes','No','Yes','No','Yes','Yes','Yes','Yes','Yes','No']. Use Naïve Bayes algorithm to predict [0: Overcast, 2: Mild]tuple belongs to which class whether to play the sports or not.

```
# Assigning features and label variables
weather=['Sunny','Sunny','Overcast','Rainy','Rainy','Rainy','Overcast','Sunny','Sunny',
'Rainy','Sunny','Overcast','Overcast','Rainy']
temp=['Hot','Hot','Hot','Mild','Cool','Cool','Cool','Mild','Cool','Mild','Mild','Mild','Hot','Mild']
play=['No','No','Yes','Yes','Yes','No','Yes','No','Yes','Yes','Yes','Yes','Yes','No']
# Import LabelEncoder
from sklearn import preprocessing
#creating labelEncoder
le = preprocessing.LabelEncoder()
# Converting string labels into numbers.
wheather_encoded=le.fit_transform(weather)
print (wheather_encoded)
```

```
# Converting string labels into numbers
temp_encoded=le.fit_transform(temp)
label=le.fit_transform(play)
print ("Temp:",temp_encoded)
print ("Play:",label)
#Combinig weather and temp into single listof tuples
features=zip(wheather_encoded,temp_encoded)
print (features)
#Import Gaussian Naive Bayes model
from sklearn.naive_bayes import GaussianNB
#Create a Gaussian Classifier
model = GaussianNB()
# Train the model using the training sets
model.fit(features,label)
#Predict Output
predicted= model.predict([[0,2]]) # 0:Overcast, 2:Mild
print("Predicted Value:", predicted)
```

SLIP NO 5

Q1. Write a R program to concatenate two given factors.

```
f1 <- factor(sample(LETTERS, size=6, replace=TRUE))
f2 <- factor(sample(LETTERS, size=6, replace=TRUE))
print("Original factors:")
print(f1)
print(f2)
f = factor(c(levels(f1)[f1], levels(f2)[f2]))
print("After concatenate factor becomes:")
print(f)
```

Q2. Write a Python program build Decision Tree Classifier using Scikit- learn package for diabetes data set (download database from https://www.kaggle.com/uciml/pimaindians-diabetes-database

```
import pandas as pd
from sklearn.tree import DecisionTreeClassifier
from sklearn.model_selection import train_test_split
from sklearn import metrics
pima = pd.read_csv("diabetes.csv")
pima.head()
import seaborn as sns
corr = pima.corr()
ax = sns.heatmap(
 corr,
 vmin=-1, vmax=1, center=0,
```

```
 cmap=sns.diverging_palette(20, 220, n=200),
 square=True
)
ax.set_xticklabels(
 ax.get_xticklabels(),
 rotation=45,
 horizontalalignment='right'
);
# feature selection
feature_cols = ['Pregnancies', 'Insulin', 'BMI', 'Age', 'Glucose', 'BloodPressure',
'DiabetesPedigreeFunction']
x = pima[feature_cols]
y = pima.Outcome
# split data
X_train, X_test, Y_train, Y_test = train_test_split(x, y, test_size = 0.3,
random_state=1)
# build model
classifier = DecisionTreeClassifier()
classifier = classifier.fit(X_train, Y_train)
# predict
y_pred = classifier.predict(X_test)
print(y_pred)
from sklearn.metrics import confusion_matrix
confusion_matrix(Y_test, y_pred)
print(confusion_matrix(Y_test, y_pred))
# accuracy
print("Accuracy:", metrics.accuracy_score(Y_test,y_pred))
from six import StringIO
from IPython.display import Image
from sklearn.tree import export_graphviz
import pydotplus
dot_data = StringIO()
export_graphviz(classifier, out_file=dot_data,
 filled=True, rounded=True,
 special_characters=True, feature_names =
feature_cols,class_names=['0','1'])
graph = pydotplus.graph_from_dot_data(dot_data.getvalue())
graph.write_png('diabetes.png')
Image(graph.create_png())
```

SLIP NO 6

Q1. Write a R program to create a data frame using two given vectors and display the duplicate
elements.

```
a = c(10,20,10,10,40,50,20,30)
b = c(10,30,10,20,0,50,30,30)
print("Original data frame:")
ab = data.frame(a,b)
print(ab)
print("Duplicate elements of the said data frame:")
print(duplicated(ab))
```

Q2. Write a python program to implement hierarchical Agglomerative clusteringalgorithm. (Download Customer.csv dataset from github.com).

```
Import pandas as pd
dataset = pd.read_csv('Mall_Customers.csv')
x = dataset.iloc[:, [3, 4]].values
import scipy.cluster.hierarchy as shc
dendro = shc.dendrogram(shc.linkage(x, method="ward"))
mtp.title("Dendrogrma Plot")
mtp.ylabel("Euclidean Distances")
mtp.xlabel("Customers")
mtp.show()
from sklearn.cluster import AgglomerativeClustering
hc= AgglomerativeClustering(n_clusters=5, affinity='euclidean', linkage='ward')
y_pred= hc.fit_predict(x)
mtp.scatter(x[y_pred == 0, 0], x[y_pred == 0, 1], s = 100, c = 'blue', label = 'Cluster 1')
mtp.scatter(x[y_pred == 1, 0], x[y_pred == 1, 1], s = 100, c = 'green', label = 'Cluster 2')
mtp.scatter(x[y_pred== 2, 0], x[y_pred == 2, 1], s = 100, c = 'red', label = 'Cluster 3')
mtp.scatter(x[y_pred == 3, 0], x[y_pred == 3, 1], s = 100, c = 'cyan', label = 'Cluster 4')
mtp.scatter(x[y_pred == 4, 0], x[y_pred == 4, 1], s = 100, c = 'magenta', label = 'Cluster 5')
mtp.title('Clusters of customers')
mtp.xlabel('Annual Income (k$)')
mtp.ylabel('Spending Score (1-100)')
mtp.legend()
mtp.show()
```

Slip 7

Q1. Write a R program to create a sequence of numbers from 20 to 50 and findthe mean of numbers from 20 to 60 and sum of numbers from 51 to 91.

```
print("Sequence of numbers from 20 to 50:")
print(seq(20,50))
print("Mean of numbers from 20 to 60:")
print(mean(20:60))
print("Sum of numbers from 51 to 91:")
print(sum(51:91))
```

Q2. Consider the following observations/data. And apply simple linear regression and find out estimated coefficients b1 and b1 Also analyse theperformance of the model (Use sklearn package) x = np.array([1,2,3,4,5,6,7,8]) y = np.array([7,14,15,18,19,21,26,23])

```
import numpy as np
import matplotlib.pyplot as plt
from scipy import stats
x = np.array([1,2,3,4,5,6,7,8])
y = np.array([7,14,15,18,19,21,26,23])
slope, intercept, r, p, std_err = stats.linregress(x, y)
def myfunc(x):
 return slope * x + intercept
mymodel = list(map(myfunc, x))
plt.scatter(x, y)
plt.plot(x, mymodel)
plt.show()
```

Slip 8

Q1. Write a R program to get the first 10 Fibonacci numbers.

```
Fibonacci <- numeric(20)
Fibonacci[1] <- Fibonacci[2] <- 1
for (i in 3:10) Fibonacci[i] <- Fibonacci[i - 2] + Fibonacci[i - 1]
print("First 10 Fibonacci numbers:")
print(Fibonacci)
```

Q2. Write a python program to implement k-means algorithm to build prediction model (Use Credit Card Dataset CC GENERAL.csv Download from kaggle.com)

```
import numpy as nm
import matplotlib.pyplot as mtp
import pandas as pd
dataset = pd.read_csv('Creditcard.csv')
dataset
x = dataset.iloc[:, [3, 4]].values
print(x)
from sklearn.cluster import KMeans
wcss_list= []
for i in range(1, 11):
 kmeans = KMeans(n_clusters=i, init='k-means++', random_state= 42)
 kmeans.fit(x)
 wcss_list.append(kmeans.inertia_)
mtp.plot(range(1, 11), wcss_list)
mtp.title('The Elobw Method Graph')
mtp.xlabel('Number of clusters(k)')
mtp.ylabel('wcss_list')
```

```
mtp.show()
kmeans = KMeans(n_clusters=3, init='k-means++', random_state= 42)
y_predict= kmeans.fit_predict(x)
mtp.scatter(x[y_predict == 0, 0], x[y_predict == 0, 1], s = 100, c = 'blue', label = 'Cluster 1') #for
first cluster
mtp.scatter(x[y_predict == 1, 0], x[y_predict == 1, 1], s = 100, c = 'green',
label = 'Cluster 2') #for second cluster
mtp.scatter(x[y_predict== 2, 0], x[y_predict == 2, 1], s = 100, c = 'red', label = 'Cluster 3')
#for third cluster
mtp.scatter(kmeans.cluster_centers_[:, 0], kmeans.cluster_centers_[:, 1], s = 300, c = 'yellow',
label = 'Centroid')
mtp.title('Clusters of Credit Card')
mtp.xlabel('V3')
mtp.ylabel('V4')
mtp.legend()
mtp.show()
```

Slip 9

Q1. Write an R program to create a Data frames which contain details of 5 employees and display summary of the data.

Employee contain (empno,empname,gender,age,designation)

```
Employees = data.frame(
                Empno=c(1,2,3,4,5),
                empname=c("Anastasia S","Dima R","Katherine S", "JAMES A","LAURA MARTIN"),
               Gender=c("M","M","F","F","M"),
               Age=c(23,22,25,26,32),
               Designation=c("Clerk","Manager","Exective","CEO","ASSISTANT")
               )
print("Summary of the data:")
print(summary(Employees))
```

Q2. Write a Python program to build an SVM model to Cancer dataset. The dataset is available in the scikit-learn library. Check the accuracyof model with precision and recall

```
from sklearn import datasets
#Load dataset
cancer = datasets.load_breast_cancer()
# print the names of the 13 features
print("Features: ", cancer.feature_names)
# print the label type of cancer('malignant' 'benign')
print("Labels: ", cancer.target_names)
# print data(feature)shape
cancer.data.shape
# print the cancer data features (top 5 records)
```

```
print(cancer.data[0:5])
# print the cancer labels (0:malignant, 1:benign)
print(cancer.target)
# Import train_test_split function
from sklearn.model_selection import train_test_split
# Split dataset into training set and test set
X_train, X_test, y_train, y_test = train_test_split(cancer.data, cancer.target,
test_size=0.3,random_state=109) # 70% training and 30% test
#Import svm model
from sklearn import svm
#Create a svm Classifier
clf = svm.SVC(kernel='linear') # Linear Kernel
#Train the model using the training sets
clf.fit(X_train, y_train)
#Predict the response for test dataset
y_pred = clf.predict(X_test)
#Import scikit-learn metrics module for accuracy calculation
from sklearn import metrics
# Model Accuracy: how often is the classifier correct?
print("Accuracy:",metrics.accuracy_score(y_test, y_pred))
```

Slip 10

Q1. Write a R program to find the maximum and the minimum value of a givenvector [10 Marks]

```
nums = c(10, 20, 30, 40, 50, 60)
print('Original vector:')
print(nums)
print(paste("Maximum value of the said vector:",max(nums)))
print(paste("Minimum value of the said vector:",min(nums)))
```

Q2. Write a Python Programme to read the dataset (“Iris.csv”). dataset download from (https://archive.ics.uci.edu/ml/datasets/iris) and apply Apriori algorithm. [20 Marks]

```
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 "# SET - A\n",
 "\n",
 "### 1) Write a code to read the dataset (“Iris.csv”). dataset download from
(https://archive.ics.uci.edu/ml/datasets/iris) and apply Apriori algorithm."
 ]
 },
 {
```

```
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"import numpy as np\n",
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"import pandas as pd\n",
"from apyori import apriori"
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]
},
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"source": [
"store_data.head()\n"
]
},
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"metadata": {},
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"source": [
"records = []\n",
"for i in range(0,300):\n",
" records.append([str(store_data.values[i,j]) for j in range(0,20)])\n"
]
},
{
"cell_type": "code",
"execution_count": null,
```

```
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 "outputs": [],
 "source": [

"association_rules=apriori(records,min_support=0.0045,min_confidence=0.2,min_lift=3,min
_length=2)\n",
 "association_results=list(association_rules)\n"
 ]
 },
 {
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 "print(len(association_results))\n"
 ]
 },
 {
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 "id": "daa923d5",
 "metadata": {},
 "outputs": [],
 "source": [
 "print(association_results[0])\n"
 ]
 },
 {
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 "id": "4f9ceaad",
 "metadata": {},
 "outputs": [],
 "source": [
 "for item in association_results:\n",
 " pair = item[0]\n",
 " items = [x for x in pair]\n",
 " print(\"Rule:\"+items[0]+\"->\"+items[1])\n",
 " \n",
 " print(\"Support:\"+str(item[1]))\n",
 "\n",
 " print(\"Confidence:\"+str(item[2][0][2]))\n",
 " print(\"Lift:\"+str(item[2][0][3]))\n",
 " print(\"=====================================\")"
```

```
]
}
],
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}
},
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}
```

SLIP NO.11

Q1.Write a R program to find all elements of a given list that are not in another given list.

```
l1 = list("x", "y", "z")
l2 = list("X", "Y", "Z", "x", "y", "z")
print("Original lists:")
print(l1)
print(l2)
print("All elements of l2 that are not in l1:")
setdiff(l2, l1)
```

Q2. Write a python program to implement hierarchical clustering algorithm.(Download Wholesale customers data dataset from github.com).

```
dataset = pd.read_csv('Mall_Customers.csv')
x = dataset.iloc[:, [3, 4]].values
import scipy.cluster.hierarchy as shc
dendro = shc.dendrogram(shc.linkage(x, method="ward"))
mtp.title("Dendrogrma Plot")
mtp.ylabel("Euclidean Distances")
```

```
mtp.xlabel("Customers")
mtp.show()
from sklearn.cluster import AgglomerativeClustering
hc= AgglomerativeClustering(n_clusters=5, affinity='euclidean', linkage='ward')
y_pred= hc.fit_predict(x)
mtp.scatter(x[y_pred == 0, 0], x[y_pred == 0, 1], s = 100, c = 'blue', label = 'Cluster 1')
mtp.scatter(x[y_pred == 1, 0], x[y_pred == 1, 1], s = 100, c = 'green', label = 'Cluster 2')
mtp.scatter(x[y_pred== 2, 0], x[y_pred == 2, 1], s = 100, c = 'red', label = 'Cluster 3')
mtp.scatter(x[y_pred == 3, 0], x[y_pred == 3, 1], s = 100, c = 'cyan', label = 'Cluster 4')
mtp.scatter(x[y_pred == 4, 0], x[y_pred == 4, 1], s = 100, c = 'magenta', label = 'Cluster 5')
mtp.title('Clusters of customers')
mtp.xlabel('Annual Income (k$)')
mtp.ylabel('Spending Score (1-100)')
mtp.legend()
mtp.show()
```

SLIP NO.12

Q1.Write a R program to create a Dataframes which contain details of 5employees and display the details.

Employee contain (empno,empname,gender,age,designation)

```
Employees = data.frame(
                Empno=c(1,2,3,4,5),
                empname=c("Anastasia S","Dima R","Katherine S", "JAMES A","LAURA MARTIN"),
               Gender=c("M","M","F","F","M"),
               Age=c(23,22,25,26,32),
               Designation=c("Clerk","Manager","Exective","CEO","ASSISTANT")
               )
print("Summary of the data:")
print(summary(Employees))
```

Q2. Write a python program to implement multiple Linear Regression modelfor a car dataset. Dataset can be downloaded from:
https://www.w3schools.com/python/python_ml_multiple_regression.asp

```
import pandas
from sklearn import linear_model
```

```
df = pandas.read_csv("car.csv")

X = df[['Weight', 'Volume']]
y = df['CO2']

regr = linear_model.LinearRegression()
regr.fit(X, y)

#predict the CO2 emission of a car where the weight is 2300kg, and the volume is 1300cm3:
predictedCO2 = regr.predict([[2300, 1300]])

print(predictedCO2)
```

SLIP NO 13

Q1. Draw a pie chart using R programming for the following datadistribution:
Digits on
Dice
1 2 3 4 5 6 Frequency of getting each number7 2 6 3 4 8

```
# Create data for the graph.
dice<- c(7, 2, 6, 3, 4, 8)
labels <- c("1", "2", "3", "4","5","6")

# Plot the chart.
pie(dice, labels)
```

Q2. Write a Python program to read "StudentsPerformance.csv" file. Solvefollowing:
- To display the shape of dataset.
- To display the top rows of the dataset with their columns.Note: Download dataset from following link :
(https://www.kaggle.com/spscientist/students-performance-inexams?select=StudentsPerformance.csv)

```
{
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 "### SET-B\n",
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 "1) Write a Python program to read \"StudentsPerformance.csv\" file. solve the
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 "- To display the shape of dataset.\n",
 "- To display the top rows of the dataset with their columns.\n",
 "- To display the number of rows randomly.\n",
 "- To display the number of columns and names of the columns.\n",
 "- Note : Download dataset from following link:\n",
 "(https://www.kaggle.com/spscientist/students-performance-
inexams?select=StudentsPerformance.csv)"
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 "import numpy as np\n",
 "import matplotlib.pyplot as plt\n",
```



```
 "import pandas as pd\n"
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```

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```
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 "store_data.columns()"
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```

SLIP NO 14

Q1. Write a script in R to create a list of employees (name) and perform thefollowing:
a. Display names of employees in the list.
b. Add an employee at the end of the list
c. Remove the third element of the list.

```
list_data <- list("Ram Sharma","Sham Varma","Raj Jadhav", "Ved Sharma")
print(list_data)

#create new employee
new_Emp <-"Kavya Anjali"
#Add new employee at the end
list_data <-append(list_data,new_Emp)
 print(list_data)
#remove 3 employee
list_data[3] <- NULL
print(list_data)
```

Q2. Write a Python Programme to apply Apriori algorithm on Groceries dataset. Dataset can be downloaded from (https://github.com/amankharwal/Websitedata/blob/master/Groceries _dataset.csv).
Also display support and confidence for each rule.

```
import numpy as np
import matplotlib.pyplot as plt
import pandas as pd
from apyori import apriori
store_data = pd.read_csv('Market_Basket_Optimisation.csv',header=None)
store_data.head()
records = []
for i in range(0, 7501):
   records.append([str(store_data.values[i,j]) for j in range(0, 20)])
association_rules = apriori(records, min_support=0.0045, min_confidence=0.2, min_lift=3,
max_length=None)
association_results = list(association_rules)
print(len(association_results))
48
for item in association_results:

   # first index of the inner list
   # Contains base item and add item
```

```
    pair = item[0]
    items = [x for x in pair]

    print("Le:",items)
    print("Rule: " + items[0] + " -> " + items[1])

    #second index of the inner list
    print("Support: " + str(item[1]))

    #third index of the list located at 0th
    #of the third index of the inner list

    print("Confidence: " + str(item[2][0][2]))
    print("Lift: " + str(item[2][0][3]))
    print("=====================================")
```

SLIP NO 15

Q1.Write a R program to add, multiply and divide two vectors of integer type.(vector length should be minimum 4)

```
vector1 = seq(10,40 , length.out=4)
vector2 = c(20, 10, 40, 40)
print("Original Vectors:")
print(vector1)
print(vector2)
add= vector1+vector2
cat("Sum of vector is ",add, "\n")
sub_vector= vector1-vector2
cat("Substraction of vector is ",sub_vector, "\n")
mul_vector= vector1 * vector2
cat("Multiplication of vector is ",mul_vector, "\n")
print("Division of two Vectors:")
div_vector = vector1 / vector2
print(div_vector)
```

Q2. Write a Python program build Decision Tree Classifier forshows.csvfrom pandas and predict class label for show starring a 40 years old American comedian, with 10 years of experience, and a comedy ranking of 7? Create a csv file as shown in https://www.w3schools.com/python/python_ml_decision_tree.asp

```
#Three lines to make our compiler able to draw:
import sys
import matplotlib
matplotlib.use('Agg')
```

```
import pandas
from sklearn import tree
from sklearn.tree import DecisionTreeClassifier
import matplotlib.pyplot as plt

df = pandas.read_csv("data.csv")

d = {'UK': 0, 'USA': 1, 'N': 2}
df['Nationality'] = df['Nationality'].map(d)
d = {'YES': 1, 'NO': 0}
df['Go'] = df['Go'].map(d)

features = ['Age', 'Experience', 'Rank', 'Nationality']

X = df[features]
y = df['Go']

dtree = DecisionTreeClassifier()
dtree = dtree.fit(X, y)

tree.plot_tree(dtree, feature_names=features)

#Two  lines to make our compiler able to draw:
plt.savefig(sys.stdout.buffer)
sys.stdout.flush()
```

SLIP 16

Q1. Write a R program to create a simple bar plot of given data
Year Export Import
2001 26 35
2002 32 40
2003 35 50

```
# Import lattice
library(lattice)

# Create data
gfg <- data.frame(x = c(26,35,32,40,35,50),
                  grp = rep(c("group 1", "group 2",
                              "group 3"),
                            each = 2),
                  subgroup = LETTERS[1:2])

# Create grouped barplot using lattice
barchart(x ~ grp, data = gfg, groups = subgroup)
```

Q2. Write a Python program build Decision Tree Classifier using Scikit-learnpackage for diabetes data set (download database from https://www.kaggle.com/uciml/pima-indiansdiabetes-database

```
import pandas as pd
from sklearn.tree import DecisionTreeClassifier
from sklearn.model_selection import train_test_split
from sklearn import metrics
pima = pd.read_csv("diabetes.csv")
pima.head()
import seaborn as sns
corr = pima.corr()
ax = sns.heatmap(
 corr,
 vmin=-1, vmax=1, center=0,
 cmap=sns.diverging_palette(20, 220, n=200),
 square=True
)
ax.set_xticklabels(
 ax.get_xticklabels(),
 rotation=45,
 horizontalalignment='right'
);
# feature selection
feature_cols = ['Pregnancies', 'Insulin', 'BMI', 'Age', 'Glucose', 'BloodPressure',
'DiabetesPedigreeFunction']
x = pima[feature_cols]
y = pima.Outcome
# split data
X_train, X_test, Y_train, Y_test = train_test_split(x, y, test_size = 0.3,
random_state=1)
# build model
classifier = DecisionTreeClassifier()
classifier = classifier.fit(X_train, Y_train)
# predict
y_pred = classifier.predict(X_test)
print(y_pred)
from sklearn.metrics import confusion_matrix
confusion_matrix(Y_test, y_pred)
print(confusion_matrix(Y_test, y_pred))
# accuracy
print("Accuracy:", metrics.accuracy_score(Y_test,y_pred))
from six import StringIO
from IPython.display import Image
from sklearn.tree import export_graphviz
import pydotplus
dot_data = StringIO()
```

```
export_graphviz(classifier, out_file=dot_data,
 filled=True, rounded=True,
 special_characters=True, feature_names =
feature_cols,class_names=['0','1'])
graph = pydotplus.graph_from_dot_data(dot_data.getvalue())
graph.write_png('diabetes.png')
Image(graph.create_png())
```

SLIP NO 17

Q1. Write a R program to get the first 20 Fibonacci numbers.

```
Fibonacci <- numeric(20)
Fibonacci[1] <- Fibonacci[2] <- 1
for (i in 3:10) Fibonacci[i] <- Fibonacci[i - 2] + Fibonacci[i - 1]
print("First 10 Fibonacci numbers:")
print(Fibonacci)
```

Q2. Write a python programme to implement multiple linear regression modelfor stock market
data frame as follows:
Stock_Market = {'Year':
[2017,2017,2017,2017,2017,2017,2017,2017,2017,2017,2017,2016,2
016,20,16,2016,2016,2016,2016,2016,2016,2016,2016,2016],
'Month': [12, 11,10,9,8,7,6,5,4,3,2,1,12,11,10,9,8,7,6,5,4,3,2,1],
'Interest_Rate': [2.75,2.5,2.5,2.5,2.5,2.5,2.5,2.25,2.25,2.25,2,2,2,1.75,1.75,1.75,1.75,1.75,1
.75,1.75,1.75,1.75,1.75,1.75],
'Unemployment_Rate':
[5.3,5.3,5.3,5.3,5.4,5.6,5.5,5.5,5.5,5.5,5.6,5.7,5.9,6,5.9,5.8,6.1,6.2,6.1,6.1,6.1,5
.9,6.2,6.2,6.1],
'Stock_Index_Price': [1464,1394,1357,1293,1256,1254,1234,1195,1159,1167,1130,1075,1047,
965,943,958,971,949,884,866,876,822,704,719] }
And draw a graph of stock market price verses interest rate.

ANSWER

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

Stock_Market = {'Year':
[2017,2017,2017,2017,2017,2017,2017,2017,2017,2017,2017,2016,2016,2016,2016,2016
,2016,2016,2016,2016,2016,2016,2016,2016],
 'Month': [12, 11,10,9,8,7,6,5,4,3,2,1,12,11,10,9,8,7,6,5,4,3,2,1],
```

```
 'Interest_Rate':
[2.75,2.5,2.5,2.5,2.5,2.5,2.5,2.25,2.25,2.25,2,2,2,1.75,1.75,1.75,1.75,1.75,1.75,1.75,1.75,1.75,1.
75,1.75],
 'Unemployment_Rate':
[5.3,5.3,5.3,5.3,5.4,5.6,5.5,5.5,5.5,5.6,5.7,5.9,6,5.9,5.8,6.1,6.2,6.1,6.1,6.1,5.9,6.2,6.2,6.1],
 'Stock_Index_Price':
[1464,1394,1357,1293,1256,1254,1234,1195,1159,1167,1130,1075,1047,965,943,958,971,949,
884,866,876,822,704,719]
 }

df =
pd.DataFrame(Stock_Market,columns=['Year','Month','Interest_Rate','Unemployment_Rate','St
ock_Index_Price'])

plt.scatter(df['Interest_Rate'], df['Stock_Index_Price'], color='red')
plt.title('Stock Index Price Vs Interest Rate', fontsize=14)
plt.xlabel('Interest Rate', fontsize=14)
plt.ylabel('Stock Index Price', fontsize=14)
plt.grid(True)
plt.show()
```

SLIP NO 18

Q1. Write a R program to find the maximum and the minimum value of a given vector.

```
nums = c(10, 20, 30, 40, 50, 60)
print('Original vector:')
print(nums)
print(paste("Maximum value of the said vector:",max(nums)))
print(paste("Minimum value of the said vector:",min(nums)))
```

Q2. Consider the following observations/data. And apply simple linear regression and find out
estimated coefficients b1 and b1 Also analyse the performance of the model
(Use sklearn package)
x = np.array([1,2,3,4,5,6,7,8])
y = np.array([7,14,15,18,19,21,26,23])

```
import numpy as np
import matplotlib.pyplot as plt

def estimate_coef(x,y):
   #number of observation/point
   n=np.size(x)
   # mean of x and y vector
```

```
    m_x = np.mean(x)
    m_y = np.mean(y)

    # calculating cross-deviation and deviation about x
    SS_xy = np.sum(y*x) - n*m_y*m_x
    SS_xx = np.sum(x*x) - n*m_x*m_x

    # calculating regression coefficients
    b_1 = SS_xy / SS_xx
    b_0 = m_y - b_1*m_x

    return (b_0, b_1)

def plot_regression_line(x, y, b):
    # plotting the actual points as scatter plot
    plt.scatter(x, y, color = "m",
    marker = "o", s = 30)

    # predicted response vector
    y_pred = b[0] + b[1]*x

    # plotting the regression line
    plt.plot(x, y_pred, color = "g")

    # putting labels
    plt.xlabel('x')
    plt.ylabel('y')

    # function to show plot
    plt.show()

def main():
    # observations / data
    x = np.array([0, 1, 2, 3, 4, 5, 6, 7, 8, 9,11,13])
    y = np.array([1, 3, 2, 5, 7, 8, 8, 9, 10, 12,16, 18])

    # estimating coefficients
    b = estimate_coef(x, y)
    print("Estimated coefficients:\nb_0 = {} \ \nb_1 = {}".format(b[0], b[1]))

    # plotting regression line
    plot_regression_line(x, y, b)

if __name__ == "__main__":
    main()
```

SLIP NO 19

Q1. Write aR program to create a Dataframes which contain details of 5 Studentsand display the
details.
Students contain (Rollno,Studname,Address,Marks)

```
Student = data.frame(
            Rollno=c(1,2,3,4,5),
            Stud Name=c("Anastasia S","Dima R","Katherine S", "JAMES A","LAURA MARTIN"),
           Adder=c("pune","mumbai","jadapsar","France","Mp"),
           Marks=c(23,22,25,26,32),
```

Q2. Write a python program to implement multiple Linear Regression modelfor a car dataset. Dataset can be downloaded from:
https://www.w3schools.com/python/python_ml_multiple_regression.asp

```
import pandas
from sklearn import linear_model

df = pandas.read_csv("data.csv")

X = df[['Weight', 'Volume']]
y = df['CO2']

regr = linear_model.LinearRegression()
regr.fit(X, y)

#predict the CO2 emission of a car where the weight is 2300kg, and the volume is 1300cm3:
predictedCO2 = regr.predict([[2300, 1300]])

print(predictedCO2)
```

SLIP NO 20

Q1. Write a R program to create a data frame from four given vectors.

```
name = c('Anastasia', 'Dima', 'Katherine', 'James', 'Emily', 'Michael', 'Matthew', 'Laura', 'Kevin',
'Jonas')
score = c(12.5, 9, 16.5, 12, 9, 20, 14.5, 13.5, 8, 19)
attempts = c(1, 3, 2, 3, 2, 3, 1, 1, 2, 1)
qualify = c('yes', 'no', 'yes', 'no', 'no', 'yes', 'yes', 'no', 'no', 'yes')
print("Original data frame:")
print(name)
print(score)
print(attempts)
print(qualify)
df = data.frame(name, score, attempts, qualify)
print(df)
```

Q2. Write a python program to implement hierarchical Agglomerativeclustering algorithm. (Download Customer.csv dataset from github.com).

```
Import pandas as pd
dataset = pd.read_csv('Mall_Customers.csv')
x = dataset.iloc[:, [3, 4]].values
import scipy.cluster.hierarchy as shc
dendro = shc.dendrogram(shc.linkage(x, method="ward"))
mtp.title("Dendrogrma Plot")
mtp.ylabel("Euclidean Distances")
mtp.xlabel("Customers")
mtp.show()
from sklearn.cluster import AgglomerativeClustering
hc= AgglomerativeClustering(n_clusters=5, affinity='euclidean', linkage='ward')
y_pred= hc.fit_predict(x)
mtp.scatter(x[y_pred == 0, 0], x[y_pred == 0, 1], s = 100, c = 'blue', label = 'Cluster 1')
mtp.scatter(x[y_pred == 1, 0], x[y_pred == 1, 1], s = 100, c = 'green', label = 'Cluster 2')
mtp.scatter(x[y_pred== 2, 0], x[y_pred == 2, 1], s = 100, c = 'red', label = 'Cluster 3')
mtp.scatter(x[y_pred == 3, 0], x[y_pred == 3, 1], s = 100, c = 'cyan', label = 'Cluster 4')
mtp.scatter(x[y_pred == 4, 0], x[y_pred == 4, 1], s = 100, c = 'magenta', label = 'Cluster 5')
mtp.title('Clusters of customers')
mtp.xlabel('Annual Income (k$)')
mtp.ylabel('Spending Score (1-100)')
mtp.legend()
mtp.show()
```