Name: Kshitij V Darwhekar

Roll No: TETB19

Sub: Soft Computitng

Batch: B2

Experiment 1: Experimental Data Analysis: Perform following operations on any open dataset available in Python/Kaggle

import numpy as np
import pandas as pd

from google.colab import drive
drive.mount('/content/drive/')

#data = open('ML/penguins_size','r')

Mounted at /content/drive/

database = pd.read_csv('/content/drive/MyDrive/ML/penguins_size.csv')

database.head()

	species	island	culmen_length_mm	culmen_depth_mm	flipper_length_mm	body_mas
0	Adelie	Torgersen	39.1	18.7	181.0	375
1	Adelie	Torgersen	39.5	17.4	186.0	380
2	Adelie	Torgersen	40.3	18.0	195.0	325
3	Adelie	Torgersen	NaN	NaN	NaN	١
4	Adelie	Torgersen	36.7	19.3	193.0	345

database.head(10)

	species	island	culmen_length_mm	culmen_depth_mm	flipper_length_mm	body_mas
0	Adelie	Torgersen	39.1	18.7	181.0	375
1	Adelie	Torgersen	39.5	17.4	186.0	380
2	Adelie	Torgersen	40.3	18.0	195.0	325
3	Adelie	Torgersen	NaN	NaN	NaN	١
4	Adelie	Torgersen	36.7	19.3	193.0	345
5	Adelie	Torgersen	39.3	20.6	190.0	365
6	Δطهانه	Tornereen	ସହ ପ	17 2	1 91 N	363

database.info()

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 344 entries, 0 to 343
Data columns (total 7 columns):

#	Column	Non-Null Count	Dtype
0	species	344 non-null	object
1	island	344 non-null	object
2	culmen_length_mm	342 non-null	float64
3	culmen_depth_mm	342 non-null	float64
4	flipper_length_mm	342 non-null	float64
5	body_mass_g	342 non-null	float64
6	sex	334 non-null	object

dtypes: float64(4), object(3)

memory usage: 18.9+ KB

print(database.isnull().sum())

dtype: int64

database = database.dropna()
database.head()

	species	island	culmen_length_mm	culmen_depth_mm	flipper_length_mm	body_mas
0	Adelie	Torgersen	39.1	18.7	181.0	375
1	Adelie	Torgersen	39.5	17.4	186.0	380
2	Adelie	Torgersen	40.3	18.0	195.0	325
4	Adelie	Torgersen	36.7	19.3	193.0	345
5	Adelie	Torgersen	39.3	20.6	190.0	365

```
len(database)
```

334

len(database.columns)

7

database.loc[(database['sex'] != 'FEMALE')& (database['sex'] != 'MALE')]

	species	island	culmen_length_mm	<pre>culmen_depth_mm</pre>	flipper_length_mm	body_mas
336	Gentoo	Biscoe	44.5	15.7	217.0	487

database['culmen_depth_mm'].fillna((database['culmen_depth_mm'].mean()), inplace=True)
database['flipper_length_mm'].fillna((database['flipper_length_mm'].mean()), inplace=True)
database['body_mass_g'].fillna((database['body_mass_g'].mean()), inplace=True)
database['culmen_length_mm'].fillna((database['culmen_length_mm'].mean()), inplace=True)
database['sex'].fillna((database['sex'].value_counts().index[0]), inplace=True)

database.reset_index()
database.head()

	species	island	culmen_length_mm	culmen_depth_mm	flipper_length_mm	body_mas
0	Adelie	Torgersen	39.1	18.7	181.0	375
1	Adelie	Torgersen	39.5	17.4	186.0	380
2	Adelie	Torgersen	40.3	18.0	195.0	325
4	Adelie	Torgersen	36.7	19.3	193.0	345
5	Adelie	Torgersen	39.3	20.6	190.0	365

col_new = ['new_species','new_island','new_culmen_length_mm','new_culmen_depth_mm','new_fl
database.columns = col_new
col_new

```
['new_species',
  'new_island',
  'new_culmen_length_mm',
  'new_culmen_depth_mm',
  'new_flipper_length',
  'new_body_mass_g',
  'new_sex']
```

database.head()

new_species new_island new_culmen_length_mm new_culmen_depth_mm new_flipper_l 0 Adelie Torgersen 39.1 18.7 1 Adelie Torgersen 39.5 17.4 Torgersen 2 Adelie 40.3 18.0

database_new = database.drop(['new_island','new_culmen_length_mm','new_flipper_length'],ax
database.head()

	new_species	new_island	new_culmen_length_mm	new_culmen_depth_mm	new_flipper_l
0	Adelie	Torgersen	39.1	18.7	
1	Adelie	Torgersen	39.5	17.4	
2	Adelie	Torgersen	40.3	18.0	
4	Adelie	Torgersen	36.7	19.3	
5	Adelie	Torgersen	39.3	20.6	

database_new.head()

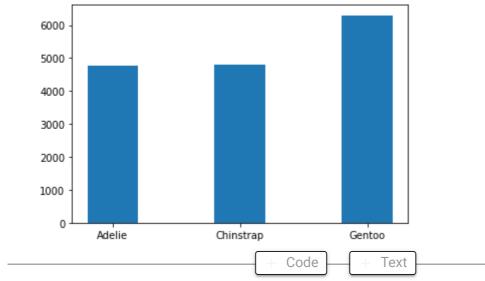
1	new_sex	new_body_mass_g	new_culmen_depth_mm	new_species	
	MALE	3750.0	18.7	Adelie	0
	FEMALE	3800.0	17.4	Adelie	1
	FEMALE	3250.0	18.0	Adelie	2
	FEMALE	3450.0	19.3	Adelie	4
	MALE	3650.0	20.6	Adelie	5

database_new["islands"] = "Torgersen"
database_new.head()

	new_species	new_culmen_depth_mm	new_body_mass_g	new_sex	islands	7
0	Adelie	18.7	3750.0	MALE	Torgersen	
1	Adelie	17.4	3800.0	FEMALE	Torgersen	
2	Adelie	18.0	3250.0	FEMALE	Torgersen	
4	Adelie	19.3	3450.0	FEMALE	Torgersen	
5	Adelie	20.6	3650.0	MALE	Torgersen	

import matplotlib.pyplot as plt
import seaborn as sns

```
X = database['new_species']
Y = database['new_body_mass_g']
plt.bar(X,Y,width = 0.4)
```

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Experiment 2: Liner Regression and Logistic Regression Model Implementation on Given Dataset.

	sbp	tobacco	ldl	adiposity	famhist	typea	obesity	alcohol	age	chd	2
0	160	12.00	5.73	23.11	Present	49	25.30	97.20	52	1	
1	144	0.01	4.41	28.61	Absent	55	28.87	2.06	63	1	
2	118	0.08	3.48	32.28	Present	52	29.14	3.81	46	0	
3	170	7.50	6.41	38.03	Present	51	31.99	24.26	58	1	
4	134	13.60	3.50	27.78	Present	60	25.99	57.34	49	1	

```
history_mapping = {'Absent': 0,'Present': 1}
pf["famhist"] = pf["famhist"].map(history_mapping)
pf.head()
```

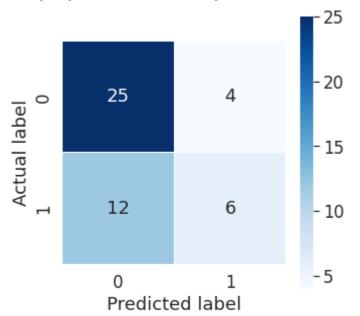
	sbp	tobacco	ldl	adiposity	famhist	typea	obesity	alcohol	age	chd	77:
0	160	12.00	5.73	23.11	1	49	25.30	97.20	52	1	
1	144	0.01	4.41	28.61	0	55	28.87	2.06	63	1	
2	118	0.08	3.48	32.28	1	52	29.14	3.81	46	0	
<pre>2 118 0.08 3.48</pre>									age', 'c		

plt.show()



```
X = pf[['tobacco','ldl','adiposity','famhist','typea','obesity','alcohol','age']].values
y = pf[['chd']].values
from sklearn.model selection import train test split
X_train , X_test , y_train,y_test = train_test_split(X,y,train_size = 0.9)
# Apply logistic regression
from sklearn.linear_model import LogisticRegression
model = LogisticRegression(C=1,penalty='12')
model.fit(X train,y train)
y_pred=model.predict(X_test)
     /usr/local/lib/python3.7/dist-packages/sklearn/utils/validation.py:993: DataConversic
       y = column_or_1d(y, warn=True)
     /usr/local/lib/python3.7/dist-packages/sklearn/linear_model/_logistic.py:818: Convers
     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,
print ('Training Accuracy: %.2f' % model.score(X_train,y_train))
print ('Test Accuracy: %.2f' % model.score(X_test,y_test))
     Training Accuracy: 0.74
     Test Accuracy: 0.66
import seaborn as sns
from sklearn.tree import plot_tree
from sklearn import tree
from sklearn.metrics import confusion matrix
cm = confusion matrix(y test,y pred)
plt.figure(figsize=(5,5))
sns.heatmap(data=cm,linewidths=.5, annot=True,square =True, cmap ='Blues')
plt.vlabel('Actual label')
plt.xlabel('Predicted label')
```

Text(0.5, 37.799999999999, 'Predicted label')



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Experiment 3: Implementation of Decision Tree, Random Forest, KNN, Naïve Bayes with hyperparameter tunning.

▼ 1. DECISION TREE

import pandas as pd
from google.colab import drive
drive.mount('/content/drive')

Mounted at /content/drive

df = pd.read_csv("/content/drive/MyDrive/ML/Titanic-Dataset.csv")
df.head()

	PassengerId	Survived	Pclass	Name	Sex	Age	SibSp	Parch	Ticket	
0	1	0	3	Braund, Mr. Owen Harris	male	22.0	1	0	A/5 21171	7.
1	2	1	1	Cumings, Mrs. John Bradley (Florence Briggs	female	38.0	1	0	PC 17599	71.

df.drop(['PassengerId','Name','SibSp','Parch','Ticket','Cabin','Embarked'],axis='columns',

df.head()

_		Survived	Pclass	Sex	Age	Fare	77
_	0	0	3	male	22.0	7.2500	
	1	1	1	female	38.0	71.2833	
	2	1	3	female	26.0	7.9250	
		<pre>df.drop(' df.Surviv</pre>		d',axis=	'colum	nns')	
	-	J	J	IIIGIO	00.0	0.0000	

inputs.Sex = inputs.Sex.map({'male': 1, 'female': 2})

```
inputs.Age[:10]
```

```
0
    22.0
    38.0
1
2
    26.0
3
    35.0
4
    35.0
5
    NaN
6
     54.0
7
    2.0
8
    27.0
9
    14.0
```

Name: Age, dtype: float64

inputs.Age = inputs.Age.fillna(inputs.Age.mean())

inputs.head()

1	Fare	Age	Sex	Pclass	
	7.2500	22.0	1	3	0
	71.2833	38.0	2	1	1
	7.9250	26.0	2	3	2
	53.1000	35.0	2	1	3
	8.0500	35.0	1	3	4

from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test = train_test_split(inputs,target,test_size=0.2)

len(X_train)

712

len(X_test)

```
from sklearn import tree
model = tree.DecisionTreeClassifier()

model.fit(X_train,y_train)
    DecisionTreeClassifier()

model.score(X_test,y_test)
    0.7877094972067039
```

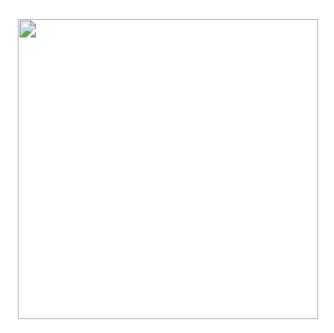
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2. KNN (K Nearest Neighbors) Classification

```
import pandas as pd
from sklearn.datasets import load_iris
iris = load_iris()
```



```
iris.feature_names

['sepal length (cm)',
    'sepal width (cm)',
    'petal length (cm)',
    'petal width (cm)']

iris.target_names

    array(['setosa', 'versicolor', 'virginica'], dtype='<U10')

df = pd.DataFrame(iris.data,columns=iris.feature_names)
df.head()</pre>
```

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)
0	5.1	3.5	1.4	0.2
1	4.9	3.0	1.4	0.2
2	4.7	3.2	1.3	0.2
2	4.6	0.4	4.5	0.0

df['target'] = iris.target
df.head()

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)	target
0	5.1	3.5	1.4	0.2	0
1	4.9	3.0	1.4	0.2	0
2	4.7	3.2	1.3	0.2	0
3	4.6	3.1	1.5	0.2	0
4	5.0	3.6	1.4	0.2	0

df[df.target==1].head()

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)	target
50	7.0	3.2	4.7	1.4	1
51	6.4	3.2	4.5	1.5	1
52	6.9	3.1	4.9	1.5	1
53	5.5	2.3	4.0	1.3	1
54	6.5	2.8	4.6	1.5	1

df[df.target==2].head()

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)	target
100	6.3	3.3	6.0	2.5	2
101	5.8	2.7	5.1	1.9	2
102	7.1	3.0	5.9	2.1	2
103	6.3	2.9	5.6	1.8	2
104	6.5	3.0	5.8	2.2	2

df['flower_name'] =df.target.apply(lambda x: iris.target_names[x])
df.head()

		sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)	target	flower_name
0	0	5.1	3.5	1.4	0.2	0	setosa
1	1	4.9	3.0	1.4	0.2	0	setosa
2	2	4.7	3.2	1.3	0.2	0	setosa
3	3	4.6	3.1	1.5	0.2	0	setosa
df[45:5	55]						
_		sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)	target	flower_name
4	45	4.8	3.0	1.4	0.3	0	setosa
4	46	5.1	3.8	1.6	0.2	0	setosa
4	47	4.6	3.2	1.4	0.2	0	setosa
4	48	5.3	3.7	1.5	0.2	0	setosa
4	49	5.0	3.3	1.4	0.2	0	setosa
5	50	7.0	3.2	4.7	1.4	1	versicolor
5	51	6.4	3.2	4.5	1.5	1	versicolor
5	52	6.9	3.1	4.9	1.5	1	versicolor
5	53	5.5	2.3	4.0	1.3	1	versicolor
5	54	6.5	2.8	4.6	1.5	1	versicolor

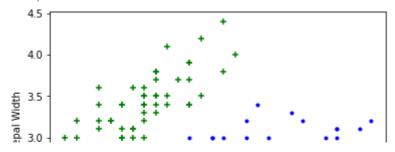
df0 = df[:50]
df1 = df[50:100]
df2 = df[100:]

import matplotlib.pyplot as plt
%matplotlib inline

Sepal length vs Sepal Width (Setosa vs Versicolor)

```
plt.xlabel('Sepal Length')
plt.ylabel('Sepal Width')
plt.scatter(df0['sepal length (cm)'], df0['sepal width (cm)'],color="green",marker='+')
plt.scatter(df1['sepal length (cm)'], df1['sepal width (cm)'],color="blue",marker='.')
```

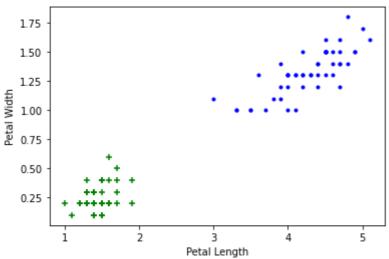
<matplotlib.collections.PathCollection at 0x7f8d07945f50>



Petal length vs Pepal Width (Setosa vs Versicolor)

```
plt.xlabel('Petal Length')
plt.ylabel('Petal Width')
plt.scatter(df0['petal length (cm)'], df0['petal width (cm)'],color="green",marker='+')
plt.scatter(df1['petal length (cm)'], df1['petal width (cm)'],color="blue",marker='.')
```

<matplotlib.collections.PathCollection at 0x7f8d07437910>



Train test split

30

Create KNN (K Neighrest Neighbour Classifier)

```
from sklearn.neighbors import KNeighborsClassifier
knn = KNeighborsClassifier(n_neighbors=10)

knn.fit(X_train, y_train)
    KNeighborsClassifier(n_neighbors=10)

knn.score(X_test, y_test)
    0.96666666666667

knn.predict([[4.8,3.0,1.5,0.3]])
    /usr/local/lib/python3.7/dist-packages/sklearn/base.py:451: UserWarning: X does not h
    "X does not have valid feature names, but"
    array([0])
```

Plot Confusion Matrix

Text(42.0, 0.5, 'Truth')



Print classification report for precesion, recall and f1-score for each classes



from sklearn.metrics import classification_report

print(classification_report(y_test, y_pred))

	precision	recall	f1-score	support
0	1.00	1.00	1.00	11
1	1.00	0.92	0.96	13
2	0.86	1.00	0.92	6
accuracy			0.97	30
macro avg	0.95	0.97	0.96	30
weighted avg	0.97	0.97	0.97	30

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3. RANDOM FOREST

```
import pandas as pd
from sklearn.datasets import load_digits
digits = load_digits()

dir(digits)

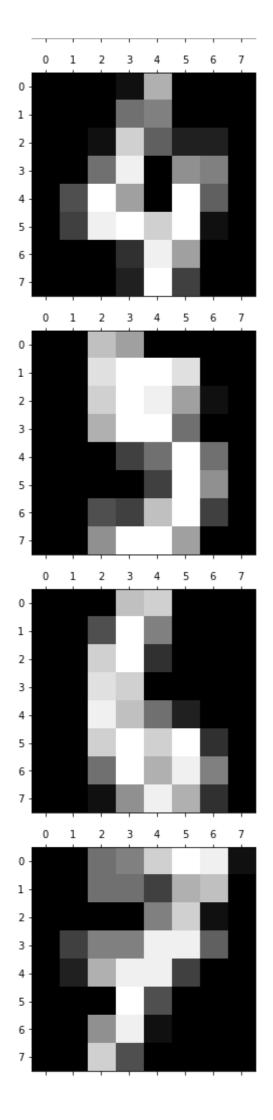
   ['DESCR', 'data', 'feature_names', 'frame', 'images', 'target', 'target_names']

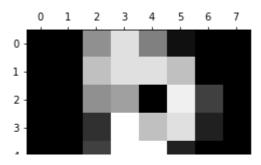
%matplotlib inline
import matplotlib.pyplot as plt

plt.gray()
for i in range(10):
   plt.matshow(digits.images[i])
```

6 -

7 -





df = pd.DataFrame(digits.data)
df.head()

	0	1	2	3	4	5	6	7	8	9	 54	55	56	57	58	59	
0	0.0	0.0	5.0	13.0	9.0	1.0	0.0	0.0	0.0	0.0	 0.0	0.0	0.0	0.0	6.0	13.0	1
1	0.0	0.0	0.0	12.0	13.0	5.0	0.0	0.0	0.0	0.0	 0.0	0.0	0.0	0.0	0.0	11.0	1
2	0.0	0.0	0.0	4.0	15.0	12.0	0.0	0.0	0.0	0.0	 5.0	0.0	0.0	0.0	0.0	3.0	1
3	0.0	0.0	7.0	15.0	13.0	1.0	0.0	0.0	0.0	8.0	 9.0	0.0	0.0	0.0	7.0	13.0	1
4	0.0	0.0	0.0	1.0	11.0	0.0	0.0	0.0	0.0	0.0	 0.0	0.0	0.0	0.0	0.0	2.0	1

5 rows × 64 columns

df['target'] = digits.target

df[0:12]

	0	1	2	3	4	5	6	7	8	9	 55	56	57	58	59	
0	0.0	0.0	5.0	13.0	9.0	1.0	0.0	0.0	0.0	0.0	 0.0	0.0	0.0	6.0	13.0	1
1	0.0	0.0	0.0	12.0	13.0	5.0	0.0	0.0	0.0	0.0	 0.0	0.0	0.0	0.0	11.0	1
2	0.0	0.0	0.0	4.0	15.0	12.0	0.0	0.0	0.0	0.0	 0.0	0.0	0.0	0.0	3.0	1
3	0.0	0.0	7.0	15.0	13.0	1.0	0.0	0.0	0.0	8.0	 0.0	0.0	0.0	7.0	13.0	1
4	0.0	0.0	0.0	1.0	11.0	0.0	0.0	0.0	0.0	0.0	 0.0	0.0	0.0	0.0	2.0	1
5	0.0	0.0	12.0	10.0	0.0	0.0	0.0	0.0	0.0	0.0	 0.0	0.0	0.0	9.0	16.0	1
6	0.0	0.0	0.0	12.0	13.0	0.0	0.0	0.0	0.0	0.0	 0.0	0.0	0.0	1.0	9.0	1
7	0.0	0.0	7.0	8.0	13.0	16.0	15.0	1.0	0.0	0.0	 0.0	0.0	0.0	13.0	5.0	
8	0.0	0.0	9.0	14.0	8.0	1.0	0.0	0.0	0.0	0.0	 0.0	0.0	0.0	11.0	16.0	1
9	0.0	0.0	11.0	12.0	0.0	0.0	0.0	0.0	0.0	2.0	 0.0	0.0	0.0	9.0	12.0	1
10	0.0	0.0	1.0	9.0	15.0	11.0	0.0	0.0	0.0	0.0	 0.0	0.0	0.0	1.0	10.0	1
11	0.0	0.0	0.0	0.0	14.0	13.0	1.0	0.0	0.0	0.0	 0.0	0.0	0.0	0.0	1.0	1

12 rows × 65 columns

```
X = df.drop('target',axis = 'columns')
y = df.target
from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(X,y,test_size=0.1)
from sklearn.ensemble import RandomForestClassifier
model = RandomForestClassifier(n estimators=30)
model.fit(X_train, y_train)
    RandomForestClassifier(n estimators=30)
model.score(X_test, y_test)
    0.95
y_predicted = model.predict(X_test)
from sklearn.datasets import make_classification
## Confusion Matrix
from sklearn.metrics import confusion_matrix
cm = confusion_matrix(y_test, y_predicted)
cm
    array([[18, 0, 0, 0, 0, 0, 0, 0, 0],
           [ 0, 17, 0, 0, 0, 0, 0, 0,
                                              0],
           [ 0, 0, 15, 0, 0, 0, 0, 0, 0,
                                             0],
           [0, 0, 0, 25, 0, 0, 0, 0, 0],
           [0, 0, 0, 0, 15, 0, 0, 0, 0,
                                              01,
           [ 0, 0, 0, 0, 1, 18, 1, 0, 0,
                                             1],
           [ 0, 0, 0, 0, 0, 18, 0, 0,
                                              01,
           [ 0, 0, 0, 0, 0, 0, 15, 0,
           [ 0, 0, 1, 2, 0, 0, 0, 0, 14,
           [0, 1, 0, 0, 0, 1, 0, 0, 1, 16]]
%matplotlib inline
import matplotlib.pyplot as plt
import seaborn as sn
plt.figure(figsize=(10,7))
```

sn.heatmap(cm, annot=True)
plt.xlabel('Predicted')
plt.ylabel('Truth')

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4. NAIVE BAYES

	precision	recall	f1-score	support
0	0.99	0.99 0.85	0.99	178 182
2	0.98	0.64	0.77	177
3	0.94	0.79	0.86	183
4	0.98	0.84	0.90	181
5	0.91	0.93	0.92	182
6	0.96	0.99	0.98	181
7	0.72	0.99	0.83	179
8	0.58	0.86	0.69	174
9	0.94	0.71	0.81	180

	ma	ccura cro a ted a	avg		0.8			0.86 0.86		0.86 0.86 0.86	1797 1797 1797
[[1	76	0	0	0	1	0	0	1	0	0]	
[0	154	0	0	0	0	3	5	14	6]	
[0	13	113	0	0	1	1	0	49	0]	
[0	2	2	145	0	6	0	7	20	1]	
[1	1	0	0	152	1	2	21	3	0]	
[0	0	0	3	0	169	1	6	2	1]	
[0	1	0	0	0	1	179	0	0	0]	
[0	0	0	0	1	1	0	177	0	0]	
[0	8	0	1	0	3	0	12	150	0]	
[1	6	0	5	1	3	0	17	20	127]]	

Multinomial Naive Bayes

from sklearn.naive_bayes import MultinomialNB
model = MultinomialNB()

model.fit(dataset.data, dataset.target)
expected = dataset.target
predicted = model.predict(dataset.data)
print(metrics.classification_report(expected, predicted))
print(metrics.confusion_matrix(expected, predicted))

	preci		cisio	on	red	call	f1	-score	support		
			0		0.9			9.98		0.99	178
			1		0.8			9.75		0.81	182
			2		0.9	90	(0.90		0.90	177
			3		0.9	99	(0.87		0.93	183
			4		0.9	96	(9.96		0.96	181
			5		0.9	97	(3.86		0.91	182
			6		0.9	98	(9.97		0.98	181
			7		0.8	39	(0.99		0.94	179
			8		0.7	78	(3.89		0.83	174
			9		0.7	76	(88.6		0.82	180
	ac	cura	асу							0.91	1797
m	nac	ro a	avg		0.9	91	(0.91		0.91	1797
weig	ght	ed a	avg		0.9	91	(0.91		0.91	1797
[[17	75	0	0	0	3	0	0	0	0	0]	
[0	137	14	0	0	1	2	0	13	15]	
Ī.	0	7	160	0	0	0	0	0	8	2]	
	0	0	2	159	0	2	0	5	8	7]	
[1	0	0	0	173	0	0	4	3	0]	
	0	0	0	0	1	157	1	1	2	20]	
[0	2	0	0	1	1	176	0	1	0]	
	0	0	0	0	0	0	0	178	1	0]	
į	0	11	1	0	1	0	1			5]	
Ĺ	0	1	0	1	1	1	0	11	7		
L	_			_	_	_			,	11	

```
model = BernoulliNB()
model.fit(dataset.data, dataset.target)
expected = dataset.target
predicted = model.predict(dataset.data)
print(metrics.classification_report(expected, predicted))
print(metrics.confusion_matrix(expected, predicted))
                    precision recall f1-score support
                         0.98
                                 0.98
                                             0.98
                                                         178

      0.76
      0.62

      0.86
      0.86

      0.91
      0.86

      0.91
      0.95

      0.93
      0.82

                1
                                              0.68
                                                         182
                2
                                                         177
                                             0.86
                                                        183
                3
                                            0.88
                                           0.93
                4
                                                         181
                5
                                            0.87
                                                         182
                6
                       0.97
                                 0.94
                                            0.96
                                                        181

      0.88
      0.98
      0.93

      0.70
      0.82
      0.75

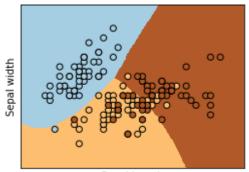
      0.76
      0.81
      0.78

                7
                                                         179
                8
                                                         174
                                                        180
                                             0.86
         accuracy
                                                    1797
                       0.87
                                 0.86
                                             0.86
                                                       1797
        macro avg
                        0.87
                                  0.86
                                              0.86
                                                       1797
     weighted avg
     [[175 1
                    0 2
                                              0]
      [ 0 112 21
                    0 3 1 1 1 32 11]
         0
            6 153 6
                          0 0 0 1 11
                                               0]
        1 1 3 157 0 2 0 3 7
                                             91
      [ 0 1 0 0 172 0 0 7 1
                                             0]
        2 3
                0 2 1 149 2 0 3 20]
        0 5 0 0 2 2 171 0 1 0]
        0 0 0 0 3 0 0 175 1
                                               0]
         0 13 1 4 0 3 2 2 142
                                               71
        0 6 0 3 7 3 0 9
                                           6 146]]
##
import numpy as np
import matplotlib.pyplot as plt
%matplotlib inline
def Naive_bayes(Model_Type):
        # import some data to play with
        iris = datasets.load iris()
        X = iris.data[:, :2] # we only take the first two features.
        Y = iris.target
        h = .02 # step size in the mesh
        # we create an instance of Neighbours Classifier and fit the data.
        if(Model_Type=='Gaussian'):
            model = GaussianNB()
        elif (Model Type=='Multinomial'):
                model = MultinomialNB()
        else:
```

model = BernoulliNB()

from sklearn.naive_bayes import BernoulliNB

```
model.fit(X, Y)
        # Plot the decision boundary. For that, we will assign a color to each
        # point in the mesh [x_min, m_max]x[y_min, y_max].
        x_{min}, x_{max} = X[:, 0].min() - .5, X[:, 0].max() + .5
        y_{min}, y_{max} = X[:, 1].min() - .5, X[:, 1].max() + .5
        xx, yy = np.meshgrid(np.arange(x_min, x_max, h), np.arange(y_min, y_max, h))
        Z = model.predict(np.c_[xx.ravel(), yy.ravel()])
        # Put the result into a color plot
        Z = Z.reshape(xx.shape)
        plt.figure(1, figsize=(4, 3))
        plt.pcolormesh(xx, yy, Z, cmap=plt.cm.Paired)
        # Plot also the training points
        plt.scatter(X[:, 0], X[:, 1], c=Y, edgecolors='k', cmap=plt.cm.Paired)
        plt.xlabel('Sepal length')
        plt.ylabel('Sepal width')
        plt.xlim(xx.min(), xx.max())
        plt.ylim(yy.min(), yy.max())
        plt.xticks(())
        plt.yticks(())
        plt.show()
        model.fit(dataset.data, dataset.target)
        expected = dataset.target
        predicted = model.predict(dataset.data)
        print(metrics.classification_report(expected, predicted))
        print(metrics.confusion_matrix(expected, predicted))
from IPython.html import widgets
from IPython.html.widgets import interact
from IPython.display import display
import warnings
warnings.filterwarnings('ignore')
i = interact(Naive bayes, Model Type=['Gaussian','Multinomial','Bernoulli'])
```



Senal	LI GO	na	th

					Sepal length					
support	score	f1-	call	red	on	cisi	pred			
178	0.99		0.99	(99	0.9		0		
182	0.84		0.85	(33	0.8		1		
177	0.77		0.64	(98	0.9		2		
183	0.86		3.79	(94	0.9		3		
181	0.90		84	(98	0.9		4		
182	0.92		9.93	(91	0.9		5		
181	0.98		9.99	(96	0.9		6		
179	0.83		3.99	(72	0.7		7		
174	0.69		3.86	(58	0.5		8		
1/9/ 1797 1797	0.86 0.86		0.86 0.86		0.88 0.89		accuracy macro avg weighted avg			
	0] 6] 0] 1] 0] 0] 0] 0]	0 14 49 20 3 2 0 0 150 20	1 5 0 7 21 6 0 177 12 17	0 3 1 0 2 1 179 0 0	0 0 1 6 1 169 1 3 3	1 0 0 152 0 0 1 0	0 0 145 0 3 0 0 1 5	0 0 113 2 0 0 0 0	0 154 13 2 1 0 1 0 8 6	[[176 [0 [0 [1 [0 [0 [0 [0

Name: Kshitij V Darwhekar

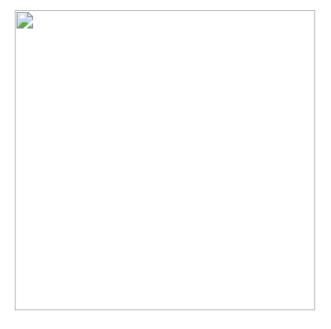
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Batch: B2

Experiment 4: Machine Learning for Image Classification (Support Vector Machine Tutorial Using Python Sklearn)

```
import pandas as pd
from sklearn.datasets import load_iris
iris = load_iris()
```



```
iris.feature_names

['sepal length (cm)',
    'sepal width (cm)',
    'petal length (cm)',
    'petal width (cm)']

iris.target_names
    array(['setosa', 'versicolor', 'virginica'], dtype='<U10')

df = pd.DataFrame(iris.data,columns=iris.feature_names)</pre>
```

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)
0	5.1	3.5	1.4	0.2
1	4.9	3.0	1.4	0.2
2	4.7	3.2	1.3	0.2
3	4.6	3.1	1.5	0.2
4	5.0	3.6	1.4	0.2

df['target'] = iris.target
df.head()

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)	target
0	5.1	3.5	1.4	0.2	0
1	4.9	3.0	1.4	0.2	0
2	4.7	3.2	1.3	0.2	0
3	4.6	3.1	1.5	0.2	0
4	5.0	3.6	1.4	0.2	0

from google.colab import drive
drive.mount('/content/drive')

Mounted at /content/drive

df[df.target==1].head()

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)	target
50	7.0	3.2	4.7	1.4	1
51	6.4	3.2	4.5	1.5	1
52	6.9	3.1	4.9	1.5	1
53	5.5	2.3	4.0	1.3	1
54	6.5	2.8	4.6	1.5	1

df[df.target==2].head()

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)	target
100	6.3	3.3	6.0	2.5	2

df['flower_name'] =df.target.apply(lambda x: iris.target_names[x])
df.head()

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)	target	flower_name
0	5.1	3.5	1.4	0.2	0	setosa
1	4.9	3.0	1.4	0.2	0	setosa
2	4.7	3.2	1.3	0.2	0	setosa
3	4.6	3.1	1.5	0.2	0	setosa
4	5.0	3.6	1.4	0.2	0	setosa

df[45:55]

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)	target	flower_name
45	4.8	3.0	1.4	0.3	0	setosa
46	5.1	3.8	1.6	0.2	0	setosa
47	4.6	3.2	1.4	0.2	0	setosa
48	5.3	3.7	1.5	0.2	0	setosa
49	5.0	3.3	1.4	0.2	0	setosa
50	7.0	3.2	4.7	1.4	1	versicolor
51	6.4	3.2	4.5	1.5	1	versicolor
52	6.9	3.1	4.9	1.5	1	versicolor
53	5.5	2.3	4.0	1.3	1	versicolor
54	6.5	2.8	4.6	1.5	1	versicolor

df0 = df[:50]
df1 = df[50:100]
df2 = df[100:]

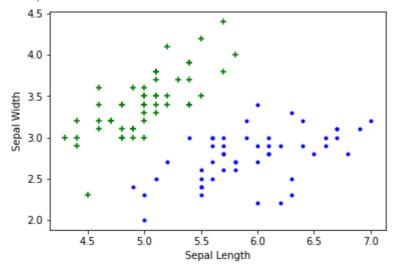
import matplotlib.pyplot as plt
%matplotlib inline

Sepal length vs Sepal Width (Setosa vs Versicolor)

plt.xlabel('Sepal Length')
plt.ylabel('Sepal Width')

```
plt.scatter(df0['sepal length (cm)'], df0['sepal width (cm)'],color="green",marker='+')
plt.scatter(df1['sepal length (cm)'], df1['sepal width (cm)'],color="blue",marker='.')
```

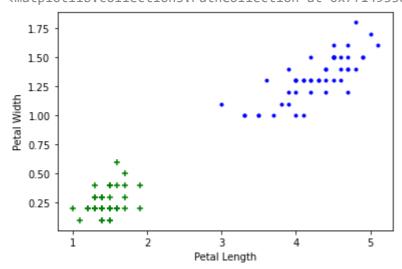
<matplotlib.collections.PathCollection at 0x7f14938e3a10>



Petal length vs Pepal Width (Setosa vs Versicolor)

```
plt.xlabel('Petal Length')
plt.ylabel('Petal Width')
plt.scatter(df0['petal length (cm)'], df0['petal width (cm)'],color="green",marker='+')
plt.scatter(df1['petal length (cm)'], df1['petal width (cm)'],color="blue",marker='.')
```

<matplotlib.collections.PathCollection at 0x7f14933d5b10>



Train Using Support Vector Machine (SVM)

```
from sklearn.model_selection import train_test_split

X = df.drop(['target','flower_name'], axis='columns')
y = df.target
```

```
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2)
len(X train)
     120
len(X_test)
     30
from sklearn.svm import SVC
model = SVC()
model.fit(X_train, y_train)
     SVC()
model.score(X_test, y_test)
     0.9666666666666667
model.predict([[4.8,3.0,1.5,0.3]])
     /usr/local/lib/python3.7/dist-packages/sklearn/base.py:451: UserWarning: X does not h
       "X does not have valid feature names, but"
     array([0])
```

Tune parameters

1. Regularization (C)

The Regularization parameter (often termed as C parameter in python's sklearn library) tells the SVM optimization how much you want to avoid misclassifying each training example.

2. Gamma

Gamma parameter: gamma determines the distance a single data sample exerts influence. That is, the gamma parameter can be said to adjust the curvature of the decision boundary.

```
model_g = SVC(gamma=10)
model_g.fit(X_train, y_train)
model_g.score(X_test, y_test)

0.96666666666666667
```

3. Kernel

A kernel is a specialized kind of similarity function. It takes two points as input, and returns their similarity as output, just as a similarity metric does. A mathematical result from linear algebra known as Mercer's theorem has the implication that a broad class of functions (e.g. similarity metrics) may be expressed in terms of a dot product in some (possibly very and even infinitely) high dimensional space. This means that calculations performed on points in high-dimensional spaces may be restated in terms of dot products

Exercise

Train SVM classifier using sklearn digits dataset (i.e. from sklearn.datasets import load_digits) and then,

- 1. Measure accuracy of your model using different kernels such as rbf and linear.
- 2. Tune your model further using regularization and gamma parameters and try to come up with highest accurancy score
- 3. Use 80% of samples as training data size

```
import pandas as pd
import numpy as np
import sklearn
import matplotlib.pyplot as plt
```

```
import seaborn as sns
digits=load_digits()
print(digits)
    {'data': array([[ 0., 0., 5., ..., 0., 0., 0.],
           [0., 0., 0., \dots, 10., 0., 0.],
           [0., 0., 0., ..., 16., 9.,
                  0., 1., ..., 6., 0., 0.],
           [ 0.,
           [ 0.,
                  0., 2., ..., 12., 0., 0.],
           [ 0., 0., 10., ..., 12., 1., 0.]]), 'target': array([0, 1, 2, ..., 8, 9, 8]
           [ 0., 0., 13., ..., 15., 5., 0.],
            Γ0.,
                  3., 15., ..., 11., 8.,
            [ 0.,
                  4., 11., ..., 12., 7.,
            [ 0., 2., 14., ..., 12., 0.,
            [0., 0., 6., ..., 0., 0.,
                                          0.]],
                  0., 0., ..., 5., 0.,
           [[ 0.,
                                          0.],
                  0., 0., ..., 9., 0.,
           [ 0.,
                                          0.],
                  0.,
                       3., ..., 6.,
            [ 0.,
                  0., 1., ..., 6., 0.,
            [ 0.,
                  0., 1., ..., 6., 0.,
                                          0.],
            [ 0.,
                  0., 0., ..., 10., 0.,
                                          0.]],
                  0., 0., ..., 12., 0.,
           [[ 0.,
           [ 0., 0., 3., ..., 14., 0.,
                                          0.],
                  0., 8., ..., 16., 0.,
           [ 0.,
                                          0.],
            [ 0.,
                  9., 16., ..., 0., 0.,
                                          0.],
                  3., 13., ..., 11., 5., 0.],
            [ 0.,
                  0., 0., ..., 16., 9., 0.]],
            [ 0.,
           . . . ,
           [[ 0.,
                  0., 1., ..., 1., 0.,
                                         0.],
           [0., 0., 13., \ldots, 2., 1., 0.],
                  0., 16., ..., 16.,
                                    5.,
            [ 0.,
            [0., 0., 16., \ldots, 15., 0.,
            [ 0., 0., 15., ..., 16., 0.,
                                          0.],
            Γ0.,
                  0., 2., ..., 6., 0.,
                                          0.]],
                  0., 2., ..., 0., 0.,
           [[ 0.,
           [ 0.,
                  0., 14., ..., 15., 1.,
                                          0.],
                  4., 16., ..., 16.,
                                    7.,
           [ 0.,
                                          0.],
            . . . ,
                  0., 0., ..., 16., 2.,
                                          0.],
            [ 0.,
                  0., 4., ..., 16., 2.,
            [ 0.,
                                          0.],
                  0., 5., ..., 12., 0.,
            [ 0.,
                                          0.]],
           [[0., 0., 10., ..., 1., 0., 0.],
```

from sklearn.datasets import load_digits

```
[0., 2., 16., \ldots, 1., 0., 0.],
            [ 0.,
                  0., 15., ..., 15., 0., 0.],
                  4., 16., ..., 16., 6., 0.],
            [ 0.,
            [0., 8., 16., ..., 16., 8., 0.],
                  1., 8., ..., 12., 1., 0.]]]), 'DESCR': ".. _digits_dataset:\n\nOpti
digits.keys()
    dict_keys(['data', 'target', 'frame', 'feature_names', 'target_names', 'images', 'DES
df=pd.DataFrame(digits.data)
print(df.head())
print(df.shape)
                  2
                        3
                             4
                                   5
                                        6
                                             7
                                                  8
                                                       9
                                                           . . .
                                                                54
                                                                     55
                                                                          56 \
    0.0
            0.0
                5.0
                     13.0
                            9.0
                                  1.0
                                       0.0
                                            0.0
                                                 0.0
                                                     0.0
                                                          . . .
                                                               0.0
                                                                    0.0
                                                                         0.0
    1 0.0 0.0 0.0
                     12.0 13.0
                                  5.0
                                       0.0
                                            0.0
                                                 0.0
                                                      0.0
                                                               0.0
                                                                    0.0
                                                                         0.0
    2 0.0 0.0 0.0
                      4.0 15.0 12.0
                                       0.0
                                                               5.0
                                            0.0 0.0
                                                     0.0
                                                                    0.0 0.0
    3 0.0 0.0 7.0
                     15.0 13.0
                                  1.0
                                       0.0
                                            0.0
                                                0.0
                                                     8.0
                                                               9.0
                                                                    0.0 0.0
    4 0.0 0.0
                0.0
                      1.0 11.0
                                  0.0
                                            0.0 0.0 0.0
                                       0.0
                                                               0.0
                                                                    0.0 0.0
                                                          . . .
             58
                   59
                         60
                               61
                                   62
                                  0.0
    0 0.0 6.0 13.0 10.0
                              0.0
                                       0.0
    1 0.0 0.0 11.0 16.0
                            10.0
                                  0.0
                                       0.0
    2 0.0 0.0
                 3.0
                       11.0
                            16.0
                                  9.0
    3 0.0 7.0 13.0 13.0
                             9.0 0.0 0.0
    4 0.0 0.0
                  2.0 16.0
                             4.0 0.0 0.0
    [5 rows x 64 columns]
     (1797, 64)
df.columns
     RangeIndex(start=0, stop=64, step=1)
df.isnull().sum()
    0
          0
    1
          0
     2
          0
     3
          0
    4
          0
    59
          0
    60
          0
    61
          0
    62
          0
    63
          0
    Length: 64, dtype: int64
```

```
df['target']=digits.target
```

df.head()

	0	1	2	3	4	5	6	7	8	9	• • •	55	56	57	58	59	60
0	0.0	0.0	5.0	13.0	9.0	1.0	0.0	0.0	0.0	0.0		0.0	0.0	0.0	6.0	13.0	10.0
1	0.0	0.0	0.0	12.0	13.0	5.0	0.0	0.0	0.0	0.0		0.0	0.0	0.0	0.0	11.0	16.0
2	0.0	0.0	0.0	4.0	15.0	12.0	0.0	0.0	0.0	0.0		0.0	0.0	0.0	0.0	3.0	11.0
3	0.0	0.0	7.0	15.0	13.0	1.0	0.0	0.0	0.0	8.0		0.0	0.0	0.0	7.0	13.0	13.0
4	0.0	0.0	0.0	1.0	11.0	0.0	0.0	0.0	0.0	0.0		0.0	0.0	0.0	0.0	2.0	16.0
5 rows × 65 columns ◀													•				

df.target

```
0
       0
1
       1
2
       2
3
      3
4
      4
1792
      9
1793
     0
1794
1795
     9
1796
Name: target, Length: 1797, dtype: int64
```

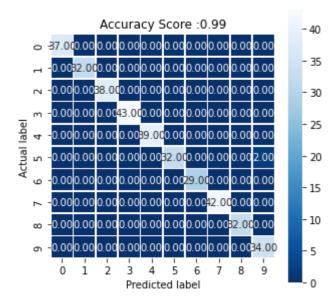
df.values

```
from sklearn.model_selection import train_test_split
x=df.drop(['target'],axis='columns')
```

```
y=df.target
x_train,x_test,y_train,y_test= train_test_split(x,y,test_size=0.2,random_state=12)
print(len(x_train))
print(len(x_test))
     1437
     360
from sklearn.metrics import accuracy_score
from sklearn.svm import SVC
model1=SVC(kernel='rbf',random_state=0, probability=True)
model1.fit(x_train,y_train)
y_pred_1=model1.predict(x_test)
print("Model Score of Kernal(rbf) :", model1.score(x_test,y_test))
     Model Score of Kernal(rbf) : 0.9916666666666667
model2=SVC(kernel='linear',random_state=0, probability=True)
model2.fit(x_train,y_train)
y_pred_2=model2.predict(x_test)
print("Model Score of Kernal(linear) :", model2.score(x_test,y_test))
     Model Score of Kernal(linear): 0.975
model3=SVC(kernel='poly',random state=0, probability=True)
model3.fit(x_train,y_train)
y pred 3=model3.predict(x test)
print("Model Score of Kernal(poly) :", model3.score(x_test,y_test))
     Model Score of Kernal(poly) : 0.99444444444445
accuracy=accuracy_score(y_test,y_pred_3)
```

```
print('ACCURACY is',accuracy)
    ACCURACY is 0.994444444444445
from sklearn.metrics import confusion_matrix
cm=np.array(confusion_matrix(y_test,y_pred_3))
cm
    array([[37, 0, 0, 0, 0, 0, 0, 0, 0],
                                        0,
                                             0],
           [0, 32, 0, 0, 0, 0, 0, 0,
           [0, 0, 38, 0, 0, 0, 0, 0, 0,
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           [0, 0, 0, 43, 0, 0, 0, 0,
                                             0],
           [ 0, 0,
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                                     0,
                                             0],
           [0, 0, 0, 0, 32, 0, 0, 0,
                                             2],
           [ 0, 0, 0, 0, 0, 29, 0,
                                         0,
                                             0],
           [ 0, 0, 0, 0, 0, 0, 42, 0,
                                            0],
           [0, 0, 0, 0, 0, 0, 0, 32,
                                             0],
           [0, 0, 0, 0, 0, 0, 0, 0, 34]])
from sklearn.metrics import mean_squared_error
mse=mean_squared_error(y_test,y_pred_3)
mse
    0.088888888888889
model1_C=SVC(C=3)
model1_C.fit(x_train,y_train)
model1_C.score(x_test,y_test)
    0.99444444444445
model2 C=SVC(C=3)
model2_C.fit(x_train,y_train)
model2_C.score(x_test,y_test)
    0.99444444444445
model3_C=SVC(C=3)
model3_C.fit(x_train,y_train)
model3_C.score(x_test,y_test)
    0.99444444444445
```

```
plt.figure(figsize=(5,5))
sns.heatmap(cm, annot=True, fmt=".2f", linewidths=.5, square = True, cmap = 'Blues_r')
plt.ylabel('Actual label')
plt.xlabel('Predicted label')
A=f'Accuracy Score :{accuracy:.2f}'
plt.title(A)
plt.show()
```



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Sub: Soft Computitng

Batch: B2

Experiment 5: To implement both the k-means algorithm and the Hierarchical Agglomerative Clustering (HAC) algorithm

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
%matplotlib inline
```

Implementation of hierarchial clustering

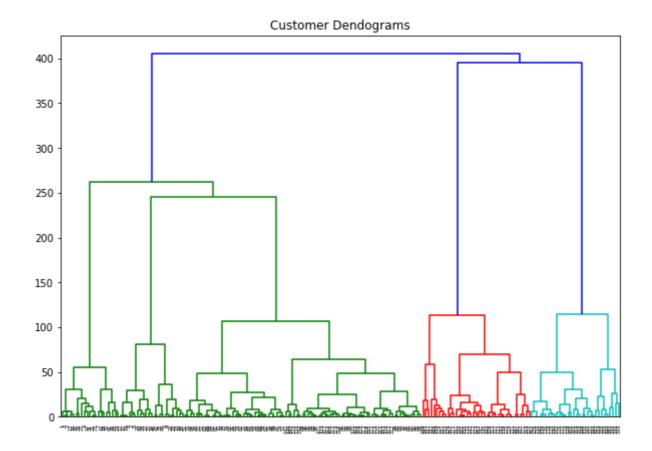
```
df1.shape
(200, 5)
```

df1.head()

	CustomerID	Genre	Age	Annual Income (k\$)	Spending Score (1-100)	17
0	1	Male	19	15	39	-
1	2	Male	21	15	81	
2	3	Female	20	16	6	
3	4	Female	23	16	77	
4	5	Female	31	17	40	

data = df1.iloc[:, 3:5].values

```
plt.figure(figsize=(10, 7))
plt.title("Customer Dendograms")
dend = shc.dendrogram(shc.linkage(data, method='ward'))
```

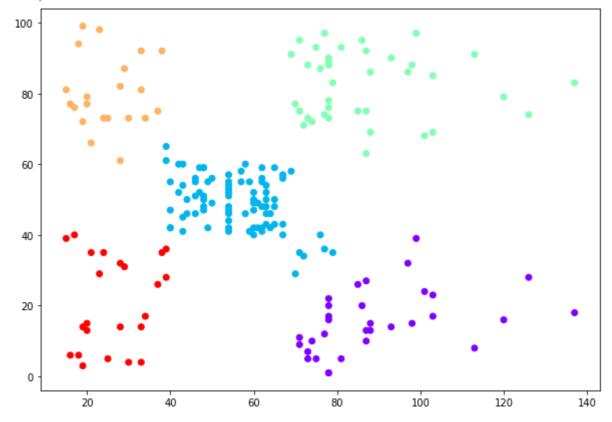


from sklearn.cluster import AgglomerativeClustering

cluster = AgglomerativeClustering(n_clusters=5, affinity='euclidean', linkage='ward')
cluster.fit_predict(data)

```
plt.figure(figsize=(10, 7))
plt.scatter(data[:,0], data[:,1], c=cluster.labels_, cmap='rainbow')
```

<matplotlib.collections.PathCollection at 0x7f658ce67f90>



Name: Kshitij V Darwhekar

Roll No: TETB19

Sub: Soft Computitng

Batch: B2

- Experiment 6: Implementation of IOT Solution using Machine Learning
- Importing the libraries

```
import sklearn
import numpy as np
import pandas as pd
```

Importing the dataset

```
from google.colab import drive
drive.mount('/content/drive/')
```

Drive already mounted at /content/drive/; to attempt to forcibly remount, call drive

dataset = pd.read_csv("/content/drive/MyDrive/ML/Crop_recommendation.csv")

```
X = dataset.iloc[:, :-1].values
y = dataset.iloc[:, -1].values
dataset.head()
```

ph



Data Preprocessing

2 60 55 44 23.004459 82.320763 7.840207 263.964248 rice

Taking care of missing data

```
from sklearn.impute import SimpleImputer
imputer = SimpleImputer(missing_values=np.nan, strategy='mean')
imputer.fit(X[:,:])
X[:,:] = imputer.transform(X[:,:])
```

Encoding categorical data

```
from sklearn.preprocessing import LabelEncoder
le = LabelEncoder()
y = le.fit_transform(y)
```

Splitting the dataset into the Training set and Test set

```
from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size = 0.25, random_state =
print(X_train)
                                                                   6.6912681
     [[134.
                     56.
                                   18.
                                               ... 83.91902605
        70.97358303]
      [ 29.
                    122.
                                  196.
                                                ... 81.15595212
                                                                   5.63832848
        73.06862952]
                                   19.
                                               ... 64.25510719
                                                                   7.10845012
      [ 25.
                     68.
        67.47677295]
      [ 35.
                     64.
                                    15.
                                                ... 63.53604453
                                                                   6.50014496
        69.5274407
      [ 39.
                     65.
                                    23.
                                                ... 69.12613376
                                                                   7.6859593
        41.02682925]
                     22.
                                    9.
                                               ... 91.13772765
                                                                   6.54319181
      [ 14.
       112.5090516 ]]
print(y_train)
```

print(X_test)

[6 7 2 ... 2 10 16]

```
[[105.
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                                                            6.41905219
  59.65590798]
                                         ... 85.49938185
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                             46.
                                                            6.34394252
               12.
  48.31219031]
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                                         ... 83.74765639
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              121.
                                                            6.15868941
  74.46411148]
 [ 84.
                27.
                            29.
                                          ... 53.00366334
                                                            7.16709259
 168.2644287
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 148.3003692
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                                                           6.29790758
 104.6735454 ]]
```

print(y_test)

```
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 6 5 10 16 13 9 19 20 11 15 4 6 12 12 21 13 11
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 6 14 13 2 0 15 18 1 17 12 10 6 16 14 21 20 15 0
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 6 20 17 6 17 3 0 19 13 20 2 12 16 8 1 17 5 6 12
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```

Feature Scaling

```
-0.65155552]
     [-0.42217223 0.31710702 -0.65031993 ... -0.35830141 0.03492274
     [-0.31405735 0.34730072 -0.4932292 ... -0.10613716 1.5951916
     -1.12940532]
     [-0.98977536 -0.95102828 -0.76813798 ... 0.88678747 0.09156286
      0.16200634]]
print(X_test)
    [ 1.46983819 -1.19257786 0.03695202 ... 0.73119109 -0.07177737
     -0.79284878]
     [ 1.09143611 -1.25296526 -0.04159334 ... 0.63244639 -0.17060505
     -0.99778658]
     [-0.98977536 2.03814779 3.0413123 ... 0.55342752 -0.41435709
     -0.52532091]
     1.16929376]
     [-0.53028711 -1.22277156 -0.29686578 ... 1.07058549 -0.17255083
      0.8086192
     [-1.23303384 -0.89064089 -0.15941139 ... 1.01005156 -0.23117683
      0.02044856]]
```

Random Forest

Training the Random Forest Classification model on the Training set

```
from sklearn.ensemble import RandomForestClassifier
classifier = RandomForestClassifier(n_estimators = 10, criterion = 'entropy', random_state
classifier.fit(X_train, y_train)
```

RandomForestClassifier(criterion='entropy', n estimators=10, random state=0)

Predicting the Test set results

Making the Confusion Matrix

```
from sklearn.metrics import confusion_matrix, accuracy_score
cm = confusion_matrix(y_test, y_pred_RF)
%matplotlib inline
import matplotlib.pyplot as plt
import seaborn as sn
plt.figure(figsize=(10,7))
sn.heatmap(cm, annot=True)
plt.xlabel('Predicted')
plt.ylabel('Truth')
     Text(69.0, 0.5, 'Truth')
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```

Predicted

10 11 12 13 14 15 16 17 18 19 20 21

accuracy_score(y_test, y_pred_RF)

0.9927272727272727

Naive Bayes

Training the Naive Bayes model on the Training set

```
from sklearn.naive_bayes import GaussianNB
classifier = GaussianNB()
classifier.fit(X_train, y_train)
     GaussianNB()
y pred NV = classifier.predict(X test)
print(np.concatenate((y_pred_NV.reshape(len(y_pred_NV),1), y_test.reshape(len(y_test),1)),
      [[21 21]
       [21 21]
       [77]
       [55]
       [ 4 4]
      [19 19]]
from sklearn.metrics import confusion matrix, accuracy score
cm = confusion_matrix(y_test, y_pred_NV)
%matplotlib inline
import matplotlib.pyplot as plt
import seaborn as sn
plt.figure(figsize=(10,7))
sn.heatmap(cm, annot=True)
plt.xlabel('Predicted')
plt.ylabel('Truth')
     Text(69.0, 0.5, 'Truth')
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Predicted

9 10 11 12 13 14 15 16 17 18 19 20 21

□ Os completed at 2:22 PM

Poster:

Crop Recommendation on Analyzing Soil Using Machine Learning

Electronics & Telecommunication Engineering Department SOFT COMPUTING (ET363) TYBTECH-Term-II (2021-22)

INTRODUCTION

- The main challenge faced in agriculture sector is the lack of knowledge about the changing variations in climate. Each crop has its own suitable climatic features. This can be handled with the help of precise farming techniques.
- The precision farming not only maintains the productivity of crops but also increases the yield rate of production. These disadvantages can be overcome with the help of precision farming. With the use of IOT and prediction system, precision farming makes decision.
- The proposed system helps in overcoming the drawbacks found in the existing system. The methods in the proposed system includes increasing the yield of crops, real-time analysis of crops using IOT, selecting efficient parameters, making smarter decisions and getting better yield.

PROBLEM STATEMENT

Recommend most suitable crop by analysing various soil parameters using machine learning and IoT

OBJECTIVES

- To use emerging technologies in improving productivity of the crops by using precision farming
- To solve the issue of cultivating crops with helps of Machine Learning model
- To collect and display data about soil parameters such as soil moisture, humidity, temperature using IoT sensors to better understand the soil
- To develop a crop recommendation system to help farmers in taking valuable decisions.

BLOCK DIAGRAM

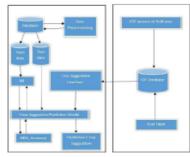
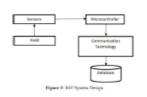


Figure 1: Block diagram of Crop Suggestion System



DATA COLLECTION & DATA PREPROCESSING

Data Collectio

- Collecting data from the field area.
- · The collected data is then stored and given as input to ML model.
- · DHT22 humidity sensor is used to measure air temperature and moisture.
- Arduino microcontroller is used which is responsible for collecting information from the sensors.
- The collected information from the sensor is stored in the excel sheet using Wi-Fi.

Data Preprocessing

- Data cleaning, data integration, data transformation and data reduction are the basic steps involved in data preprocessing.
- The data is cleaned since noisy data cant be input to ML model.
- Any difference in features is referred as incompatible or inconsistent data. The lack features or attributes in the dataset is referred to as incomplete data.

IMPLEMENTATION OF ML MODEL

- Use of supervised learning algorithm Random forest and Naïve Bayes.
- It combines multiple classifiers to solve a complex problem and to improve the performance of the model.
- Random forest takes prediction from multiple trees
- · Naïve Bayes is based on Bayes theorem.
- It is one of the simple and most effective algorithm for classification..

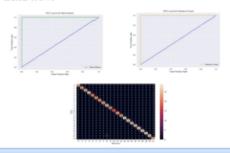
MLMODEL DEPLOYMENT USING IOT

In our proposed system we collect data from sensors and them implement ML model on the dataset for predicting the crop.

Then we use the training data to train the model and test dataset to test the model.

RESULTS AND CONCLUSION

The model then predicts and suggests the crops to be shown with an accuracy of about 99.97%



MIT Academy of Engineering

```
Name: Kshitij V Darwhekar
Roll No: TETB19
Sub: Soft Computitng
Batch: B2
import tensorflow
from tensorflow import keras
import tensorflow as tf
from tensorflow import keras
import matplotlib.pyplot as plt
%matplotlib inline
import numpy as np
(X_train, y_train) , (X_test, y_test) = keras.datasets.mnist.load_data()
    Downloading data from <a href="https://storage.googleapis.com/tensorflow/tf-keras-datasets/mni">https://storage.googleapis.com/tensorflow/tf-keras-datasets/mni</a>
    11493376/11490434 [===========] - Os Ous/step
    11501568/11490434 [============= ] - Os Ous/step
len(X_train)
    60000
len(X_test)
 X_train[0].shape
     (28, 28)
X_train[0]
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              0,
                                                0, 80, 156, 107, 253, 253,
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```

plt.matshow(X train[0])

```
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y_train[0]

5

```
# Scaling Technique
```

```
X_train = X_train / 255
X_test = X_test / 255
```

X_train[0]

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X_train_flattened = X_train.reshape(len(X_train), 28*28)
X_test_flattened = X_test.reshape(len(X_test), 28*28)
X_train_flattened.shape
       (60000, 784)
X_test_flattened.shape
       (10000, 784)
X_train_flattened[0]
                 0.97647059, 0.25098039, 0. , 0. , 0.
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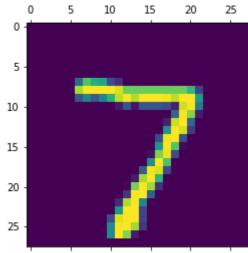
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10 Output neuron and 784 in input neuron

```
To undo cell deletion use Ctrl+M Z or the Undo option in the Edit menu \,\,	imes\, oid')
```

```
model.evaluate(X_test_flattened, y_test)
    [0.2695145010948181, 0.9254000186920166]
y_predicted = model.predict(X_test_flattened)
y_predicted[0]
    array([1.5027165e-02, 3.9325224e-07, 6.3410342e-02, 9.5975685e-01,
          3.2548308e-03, 1.0176152e-01, 1.0720740e-06, 9.9978119e-01,
          7.0441395e-02, 6.0749489e-01], dtype=float32)
plt.matshow(X_test[0])
    <matplotlib.image.AxesImage at 0x7efc920031d0>
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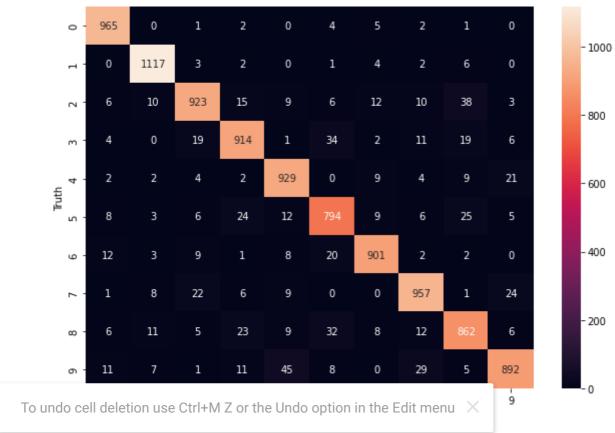
nn argmax finds a maximum element from an array and returns the index of it To undo cell deletion use Ctrl+M Z or the Undo option in the Edit menu X

```
y_predicted_labels = [np.argmax(i) for i in y_predicted]
y_predicted_labels[:5]
    [7, 2, 1, 0, 4]
cm = tf.math.confusion_matrix(labels=y_test,predictions=y_predicted_labels)
\mathsf{cm}
    <tf.Tensor: shape=(10, 10), dtype=int32, numpy=
    array([[ 965,
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dtype=int32)>
```

```
import seaborn as sn
plt.figure(figsize = (10,7))
sn.heatmap(cm, annot=True, fmt='d')
plt.xlabel('Predicted')
plt.ylabel('Truth')
```

Text(69.0, 0.5, 'Truth')



Using hidden layer

model = keras.Sequential([

```
Epoch 2/5
   Epoch 3/5
   Epoch 4/5
   Epoch 5/5
   <keras.callbacks.History at 0x7efc8ed501d0>
model.evaluate(X_test_flattened,y_test)
   313/313 [============== ] - 0s 1ms/step - loss: 0.0786 - accuracy: 0.9
   [0.07863084971904755, 0.9763000011444092]
y_predicted = model.predict(X_test_flattened)
y_predicted_labels = [np.argmax(i) for i in y_predicted]
cm = tf.math.confusion_matrix(labels=y_test,predictions=y_predicted_labels)
plt.figure(figsize = (10,7))
sn.heatmap(cm, annot=True, fmt='d')
plt.xlabel('Predicted')
plt.ylabel('Truth')
   Text(69.0, 0.5, 'Truth')
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Predicted

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Using Flatten layer so that we don't have to call .reshape on input dataset

```
model = keras.Sequential([
 keras.layers.Flatten(input_shape=(28, 28)),
 keras.layers.Dense(100, activation='relu'),
 keras.layers.Dense(10, activation='sigmoid')
])
model.compile(optimizer='adam',
     loss='sparse_categorical_crossentropy',
     metrics=['accuracy'])
model.fit(X_train, y_train, epochs=10)
Epoch 1/10
  Epoch 2/10
  Epoch 3/10
  Epoch 4/10
  Epoch 5/10
  Epoch 6/10
  Epoch 7/10
  Epoch 8/10
  Epoch 9/10
  Epoch 10/10
  <keras.callbacks.History at 0x7efc92f47c90>
To undo cell deletion use Ctrl+M Z or the Undo option in the Edit menu X
  313/313 [============= ] - 0s 1ms/step - loss: 0.0913 - accuracy: 0.9
  [0.0912703275680542, 0.9740999937057495]
```

Name: Kshitij V Darwhekar

Roll No: TETB19

Sub: Soft Computitng

Batch: B2

Experiment 8: Open CV for Computer Vision

```
In [ ]:
        import cv2
In [ ]:
        scale_percent = 20 # percent of original size
        width = int(img.shape[1] * scale_percent / 100)
        height = int(img.shape[0] * scale_percent / 100)
         dim = (width,height)
In [ ]:
        faceHar = cv2.CascadeClassifier('haarcascade_frontalface_default.xml')
         img = cv2.imread('./Kshit.JPG')
         imgGray = cv2.cvtColor(img, cv2.COLOR_BGR2GRAY)
         faces = faceHar.detectMultiScale(imgGray, 1.1, 4)
In [ ]:
        for (x, y ,w, h) in faces:
             cv2.rectangle(img, (x, y), (x+h, y+w), (0, 255, 0), 2)
         resized = cv2.resize(img, dim, interpolation = cv2.INTER_AREA)
In [ ]:
        cv2.imshow("Result", resized)
         cv2.waitKey(0)
Out[]:
```

Output :

