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1. Study of Machine learning Libraries In python.

* The libraries used in python for machine learning :-

* Numpy

Numpy - short for Numerical python. Is a fundamental package for numerical computing in python. It provides support for large multi-dimensional arrays and matrices, along with a collection of mathematical functions to operate on these arrays efficiently.

Numpy is widely used in scientific computing, data analysis, and machine learning due to its speed and versatility.

* Pandas

pandas is a powerful open-source data analysis and manipulation library built on top of Numpy.

It provides easy to use data structure and functions for working with structured data, such as time series tabular data and heterogeneous data. pandas is widely used for data cleaning, transformation, exploration, and analysis in data science and machine learning projects.

* Scikit-learn :

Scikit-learn, often abbreviated as sklearn, is a popular python library for machine learning tasks. It provides a wide array of tools for building and applying various machine learning algorithms.

Some of the key libraries and modules within Scikit-learn.

- * **dataset** : This module includes utilities for loading datasets, both standard datasets included with Scikit-learn and functions to load external datasets. It provides easy access to sample datasets for practice and experimentation.
- * **Preprocessing** : This module contains functions for preprocessing data before feeding it into machine learning models. It includes tools for scaling, centering, normalizing, binarizing and imputing missing values in datasets.
- * **Feature_extraction** : This module is for feature extraction from raw data. It includes methods such as Bag of words, TF-IDF, and feature hashing, which are commonly used in natural language processing (NLP) tasks.
- * **Feature_selection** : This module provides utilities for selecting relevant features or reducing the dimensionality of the dataset. Techniques include univariate feature selection, recursive feature elimination and feature importances.
- * **Model_selection** : This module includes tools for model selection and evaluation. It provides

functions for splitting datasets into training and test sets for cross-validation, hyperparameter tuning using grid search and performance metrics evaluation

- * **Metric** : This module contains various metrics for evaluating the performance of machine learning models. It includes metrics for classification, regression clustering, and ranking tasks, such as accuracy, precision, recall, f1-score, mean squared error and ROC AUC.
- * **Classification** : This module includes algorithms and tools for classification tasks. It provides implementations for popular classifiers such as Support Vector Machine (SVM), Random Forest, k-Nearest Neighbors (k-NN), and Naive Bayes.
- * **Regression** : Similar to the classification module, this contains algorithms and tools for regression, Lasso regression, and support vector regression, among others.
- * **Clustering** : This module provides algorithms and tools for clustering data into groups based on similarity. It includes implementations of k-means, hierarchical clustering, DBSCAN, and spectral clustering algorithms.

* **Dimensionality reduction** : This module contains techniques for reducing the dimensionality of data while preserving its structure and important features. Method includes Principal Component Analysis (PCA), t-distributed Stochastic Neighbor Embedding (t-SNE), and manifold learning techniques.

* **Ensemble** : This module includes ensemble methods for combining multiple base estimators to improve predictive performance. It provides implementation of techniques such as bagging, boosting and random forests.

* **neighbors** : This module provides functionalities for working with nearest neighbor algorithms, including k-nearest neighbors (kNN) for classification and regression tasks.

The `sklearn.compose` module in scikit-learn provides tools for combining and transforming datasets. It includes classes and functions that allow you to create composite estimators and pipelines, perform feature unions, and apply column transformations.

* Few classes or functions within the scikit-learn library.

1. **SimpleImputer** : This is a class within the `scikit-learn` `imputer` module. It is used for imputing missing

Values in datasets. The SimpleImputer class provides several strategies for imputation, such as replacing missing values with the mean, median, most frequent value or a constant value.

2. LabelEncoder :- This is a class within the sklearn.preprocessing module. It is used for encoding categorical labels as numerical values. The LabelEncoder class assigns a unique integer to each unique category in the categorical variable.

3. OneHotEncoder :- Also a class within the sklearn.preprocessing module. It is used for converting categorical variables into one-hot encoded format. Onehot encoding is a process where categorical variables are converted into binary vectors, with each binary vector representing a unique category. This is useful for algorithms that require numerical input.

4. StandardScaler :- This is another class within the sklearn.preprocessing module. It is used for standardizing features by removing the mean and scaling to unit variance. Standardization is important for many machine learning algorithms as it helps in bringing features onto the same scale, which can prevent some features from dominating others in the model training process.

5.

Column Transformer : key component of ~~the~~ the sklearn. ~~Compose~~. This class allows you to apply different transformations to different columns of a dataset. It's particularly useful when dealing with datasets containing a mix of numerical and categorical features. You can specify the transformations to be applied to each column using a list of tuples, where each tuple consists of a transformer and a list of column indices or names.

*

Seaborn is a python data visualization library based on matplotlib. It provides a high-level interface for drawing attractive and informative statistical graphics. Seaborn is particularly useful for visualizing statistical relationships and patterns in datasets.

Seaborn will generate a grid of plots showing the pairwise relationships between the variables in the dataset. The diagonal plots will display histograms or kernel density estimates of each variables distribution, while the off-diagonal plots will show scatterplots of the relationships between pairs of variables.

There are commonly used libraries for machine learning workflows. They provide convenient functions and methods to perform various tasks in machine learning.

2. Take 10 blocks and go running in nature
Morning.

1000 meters (a night) in 20

1000 meters in 18

1000 meters in 22

20 - 25 m/s. (at 1000 m)

20 sec.

20 sec.

20 sec.

20 sec.)

output	country	age	salary	purchased
0	France	NaN	7244	No
1	Spain	27.0	4900	Yes
2	Germany	36.0	5400	Yes
3	UK	49.0	98000	No

*	country	age	salary	purchased
0	France	NaN	7200	No
1	Spain	27.0	4500	Yes
2	Germany	36.0	5400	Yes
3	UK	49.0	98000	No

*	#	column	Non-null count	dtype
0	0	country	4 non-null	object
1	1	age	3 non-null	float64
2	2	salary	4 non-null	int64
3	3	purchased	4 non-null	object

*	age	salary
Count	3.460800	4.500000
mean	35.333333	28230.000000
std	10.930353	46111.218441
min	27.000000	4800.000000
25%	28.800000	5256.000000
50%	30.000000	6360.000000
75%	39.500000	24900.000000
max	49.000000	98000.000000

Country 0
age 1
salary 0
purchased 0
dtype : int64

country 0
age 0
salary 0
purchased 0
dtype: int64

array ([['France', 35.3333333336, 7200.0],
['Spain', 27.0, 4800.0],
['Germany', 30.0, 5400.0],
['UK', 49.0, 9800.0]], dtype = object)

array ([['No'],
['Yes'],
['Yes'],
['No']], dtype = object)

df. isnull(). sum()

df['age'].fillna(df['age'].mean(), inplace = True)

df['salary'].fillna(df['salary'].mean(), inplace = True)

df. is null. sum()

from sklearn. impute import SimpleImputer

x = df. iloc[:, :-1]. values

y = df. iloc[:, 3:]. values

imp = SimpleImputer (missing_values = np. nan, strategy

x[:, 1:3] = imp. fit_transform (x[:, 1:3])

x

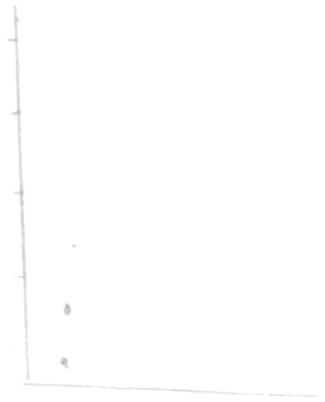
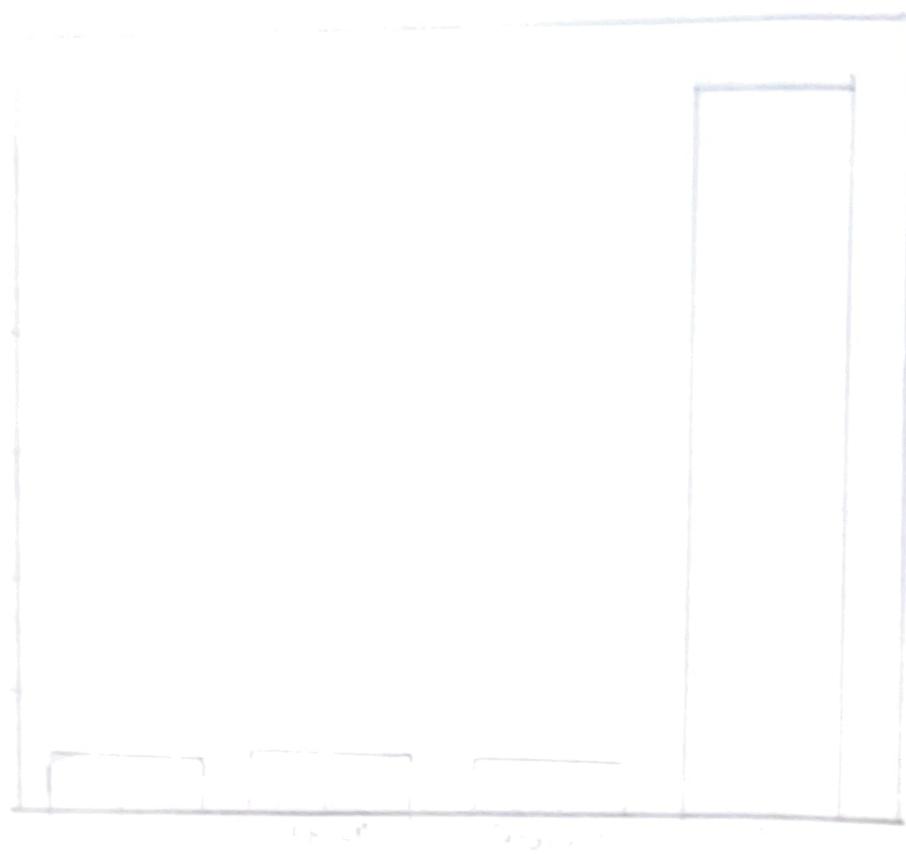
y

from sklearn. preprocessing import LabelEncoder

le = LabelEncoder()

h = le. fit_transform (x[:, 0])

y = le. fit_transform (y)



Output : Logistic Regression (max_iter, 1000)

3. Evaluate the classifier using various performance measures.

```
from sklearn.datasets import load_iris
```

```
from sklearn.model_selection import train_test_split
```

```
from sklearn.linear_model import LogisticRegression
```

```
from sklearn.metrics import accuracy_score, precision_score,  
    f1_score, roc_auc, classification_report,  
    confusion_matrix, recall_score
```

```
iris = load_iris()
```

```
X = iris.data
```

```
y = iris.target
```

```
X_train, X_test, y_train, y_test = train_test_split(X, y,  
    test_size=0.2, random_state=42)
```

```
log_reg = LogisticRegression(max_iter=1000)
```

```
log_reg.fit(X_train, y_train)
```

```
y_pred = log_reg.predict(X_test)
```

```
accuracy = round(accuracy_score(y_test, y_pred) * 100, 2)
```

```
print("Accuracy:", accuracy)
```

* Accuracy : 100.0

* Classification Report

	precision	recall	f1-score	support
0	1.00	1.00	1.00	10
1	1.00	1.00	1.00	9
2	1.00	1.00	1.00	11
accuracy			1.00	30
macro avg	1.00	1.00	1.00	30
weighted avg	1.00	1.00	1.00	30

* precision : 1.0

* Recall : 1.0

* f1 score : 1.0

* ROC AUC score: 1.0

* Confusion matrix:

$$\begin{bmatrix} [10 & 0 & 0] \\ [0 & 9 & 0] \\ [0 & 0 & 11] \end{bmatrix}$$

```
result = classification_report(y_test, y_pred)
print("Classification Report: ", )
print(result)
```

```
precision = precision_score(y_test, y_pred, average = 'weighted')
print("Precision: ", precision)
```

```
recall = recall_score(y_test, y_pred, average = 'weighted')
print("Recall: ", recall)
```

```
f1 = f1_score(y_test, y_pred, average = 'weighted')
print("F1 Score: ", f1)
```

```
roc_auc = roc_auc_score(y_test, logreg.predict_proba(x_test), multi_class = 'ovr')
print("ROC AUC Score: ", roc_auc)
```

```
confusion_matrix = confusion_matrix(y_test, y_pred)
print("Confusion Matrix: \n", confusion_matrix)
```

Output : confusion Matrix

$$\begin{bmatrix} 11 & 0 & 0 \\ 0 & 12 & 1 \\ 0 & 0 & 6 \end{bmatrix}$$

4. Implement k-nearest neighbor classification using python

```
import pandas as pd
```

```
import numpy as np
```

```
from sklearn.datasets import load_iris
```

```
iris = load_iris()
```

```
X = iris.data
```

```
y = iris.target
```

```
from sklearn.model_selection import train_test_split  
X_train, X_test, y_train, y_test = train_test_split  
(X, y, test_size=0.2, random_state=0)
```

```
from sklearn.neighbors import KNeighborsClassifier
```

```
lr = KNeighborsClassifier(n_neighbors=5)
```

```
lr.fit(X_train, y_train)
```

```
y_pred = lr.predict(X_test)
```

```
from sklearn.metrics import classification_report, confusion  
matrix, accuracy_score
```

```
result = confusion_matrix(y_test, y_pred)
```

```
print("Confusion matrix :")
```

```
print(result)
```

Classification Report

	precision	recall	f1-score	support
0	1.00	1.00	1.00	11
1	1.00	0.92	0.96	13
2	0.26	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

Accuracy: 96.67

result1 = classification_report(y_test, y_pred)
print("Classification Report: ")
print(result1)

result2 = round(accuracy_score(y_test, y_pred)*100, 2)
print("Accuracy : ", result2)

```
result1 = classification_report(y_test, y_pred)
print ("Classification Report: ")
print (result1)
```

```
result2 = round(accuracy_score(y_test, y_pred)*100,2)
print ("Accuracy : ", result2)
```

Output: $\begin{bmatrix} [5.1 \ 3.5 \ 1.4 \ 0.2] \\ \vdots \ \vdots \ \vdots \ \vdots \\ [4.8 \ 3.4 \ 1.6 \ 0.2] \end{bmatrix}$

(150, 2)

Output: $\begin{bmatrix} 5.1 & 3.5 & 1.4 & 0.2 \end{bmatrix}$

$\begin{bmatrix} 4.8 & 3.4 & 1.6 & 0.2 \end{bmatrix}$

(150, 2)

5. Dimensionality reduction using PCA

```
import numpy as np
```

```
import pandas as pd
```

```
from sklearn.datasets import load_iris
```

```
from sklearn.preprocessing import StandardScaler
```

```
from sklearn.decomposition import PCA
```

```
import matplotlib.pyplot as plt
```

```
iris = load_iris()
```

```
X = iris.data
```

```
y = iris.target
```

```
print(X)
```

```
target_names = iris.target_names
```

```
scaler = StandardScaler()
```

```
X_scaled = scaler.fit_transform(X)
```

```
pca = PCA(n_components=2)
```

```
X_pca = pca.fit_transform(X_scaled)
```

```
print(X_pca.shape)
```

DATE.....

```
plt.scatter(x_pca[:,0], x_pca[:,1], c=y, cmap='  
plasma')  
plt.xlabel('First principal component')  
plt.ylabel('Second principal component')  
plt.title('PCA of IRIS dataset')  
plt.show()
```

- Direct Expression
- Indirect Expression
- Indirect Indirect Expression
- Indirect Indirect Indirect Expression

Output : * Linear Regression.

* 0.35170909~~4~~23059894

* 0.95332824 88134365

* 0.1368291774258005

6.

Regression Analysis using Linear Regression

```
import numpy as np
```

```
import pandas as pd
```

```
from sklearn.model_selection import train_test_split
```

```
from sklearn.metrics import mean_squared_error,  
mean_absolute_error, r2_score
```

```
from sklearn.datasets import load_iris
```

```
iris = load_iris()
```

```
X = iris.data
```

```
y = iris.target
```

```
x_train, *test, y_train, y_test = train_test_split
```

```
(X, y, train_size=0.7, test_size=0.3, random_state=100)
```

```
from sklearn.linear_model import LinearRegression
```

```
lr = LinearRegression()
```

```
lr.fit(X_train, y_train)
```

```
y_pred = lr.predict(X_test)
```

```
mse = mean_squared_error(y_test, y_pred)
```

```
print(mse)
```

```
r2 = r2_score(y_test, y_pred)
```

```
print(r2)
```

```
mae = mean_absolute_error(y_test, y_pred)
```

```
print(mae)
```

Output ± 100.0

Classification Report

	precision	recall	F1-score	support
0	1.00	1.00	1.00	11
1	1.00	1.00	1.00	13
2	1.00	1.00	1.00	6
accuracy			1.00	30
macro avg	1.00	1.00	1.00	30
weighted avg	1.00	1.00	1.00	30

7. Classification using Logistic Regression

```
import pandas as pd
```

```
import numpy as np
```

```
from sklearn.datasets import load_iris
```

```
iris = load_iris()
```

```
x = iris.data
```

```
y = iris.target
```

```
from sklearn.model_selection import train_test_split
x_train, x_test, y_train, y_test = train_test_split
(x, y, test_size=0.2, random_state=0)
from sklearn.linear_model import LogisticRegression
from sklearn.metrics import classification_report,
confusion_matrix, accuracy_score
```

```
logreg = LogisticRegression(max_iter=1000)
```

```
logreg.fit(x_train, y_train)
```

```
y_pred = logreg.predict(x_test)
```

```
acc_logreg = round(accuracy_score(y_pred, y_test)*
100, 2)
```

```
print(acc_logreg)
```

```
result1 = classification_report(y_test, y_pred)
```

```
print("Classification Report :")
```

```
print(result1)
```

Confusion matrix:

$$\begin{bmatrix} 11 & 0 & 0 \\ 0 & 13 & 0 \\ 0 & 0 & 6 \end{bmatrix}$$

Confusion matrix:

$$\begin{bmatrix} [11 & 0 & 0] \\ [0 & 13 & 0] \\ [0 & 0 & 6] \end{bmatrix}$$

```
result = confusion_matrix(y_test, y_pred)
print ("Confusion matrix:")
print (result)
```

Output :- Decision Tree classifier
Decision Tree classifier()

* array(['setosa', 'versicolor', 'virginica'], dtype='|O')

8. Classification using Decision Tree

```
import pandas as pd  
from sklearn.tree import DecisionTreeClassifier, plot_tree  
from sklearn.model_selection import train_test_split  
from sklearn import metrics  
  
from sklearn.datasets import load_iris  
iris = load_iris()  
X = iris.data  
y = iris.target  
  
dtc = DecisionTreeClassifier()  
  
X_train, X_test, y_train, y_test = train_test_split(iris.data, iris.target, test_size=0.3, random_state=1)  
dtc.fit(X_train, y_train)  
  
iris.target_names
```

```
import matplotlib.pyplot as plt
plt.figure(figsize=(12,8))
plot_tree(dtcl, feature_names = bus.feature_names,
         class_names = bus.target_names, filled = True)
plt.show()
```

Output : * SVC
SVC()

Confusion matrix
[[16 60]
 [0 110]
 [0 117]]

q.

Classification using Support Vector machine

```
import numpy as np
import pandas as pd
from sklearn import model_selection import train_test_split
from sklearn import datasets import load_iris
iris = load_iris()
X = iris.data
y = iris.target

from sklearn import svm
lr = SVM().fit(X_train, y_train)

y_pred = lr.predict(X_test)
from sklearn import classification_report,
from sklearn import metrics import classification_report,
confusion_matrix, accuracy_score
result = confusion_matrix(y_test, y_pred)
print ("confusion matrix")
print(result)
```

Performance Metrics

	precision	recall	f1 score	accuracy
0	1.00	1.00	1.00	1.00
1	0.94	1.00	0.96	0.97
2	1.00	0.94	0.97	0.97

Average : 91.9133333

result = classification_report(y_test, y_pred)

print ("Classification report :")

print (result)

result = accuracy_score(y_true, y_pred)*100.2

print ("Accuracy %", result)

* Output : Gaussian NB
Gaussian NBC

Accuracy : 96.67

10. Classification Techniques - Naïve Bayes

```

import pandas as pd
import numpy as np
from sklearn.model_selection import train_test_split
from sklearn.naive_bayes import GaussianNB
from sklearn.metrics import accuracy_score
from sklearn.datasets import load_iris

iris = load_iris()
X = iris.data
y = iris.target

from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=0)

from sklearn.naive_bayes import GaussianNB
gnb = GaussianNB()
gnb.fit(X_train, y_train)

y_pred = gnb.predict(X_test)

from sklearn.metrics import classification_report, confusion_matrix
accuracy = accuracy_score(y_test, y_pred)
print("Accuracy : ", accuracy)
result2 = round(accuracy_score(y_test, y_pred) * 100)
print("Accuracy : ", result2)

```

Classification Report

	Precision	Recall	F1 Score	Support
0	1.00	1.00	1.00	11
1	0.93	1.00	0.96	13
2	1.00	0.93	0.96	6
All	0.94	0.94	0.94	30

Accuracy

	Macro Avg	Weighted Avg
Accuracy	0.98	0.97

3000 2000 1000 0

1000 800 600 400 200 0

1000 800 600 400 200 0

1000 800 600 400 200 0

1000 800 600 400 200 0

1000 800 600 400 200 0

1000 800 600 400 200 0

1000 800 600 400 200 0

1000 800 600 400 200 0

1000 800 600 400 200 0

1000 800 600 400 200 0

1000 800 600 400 200 0

1000 800 600 400 200 0

1000 800 600 400 200 0

1000 800 600 400 200 0

1000 800 600 400 200 0

1000 800 600 400 200 0

1000 800 600 400 200 0

1000 800 600 400 200 0

1000 800 600 400 200 0

1000 800 600 400 200 0

```
result = classification_report(y_true,y_pred)
print ("Classification Report:")
print(result)
```

output
size of X : (500, 2)
size of Y : (500, 1)

* random

n-clusters = 4, random_state = 42)

array([5, 6, 2, 0, 3, 2, - - - - - ,
- - - - - - - - - - - , 1, 3, 2, 2, 0])

Clustering using K-means Clustering

```
import numpy as np  
import pandas as pd  
import matplotlib.pyplot as plt  
from sklearn import datasets  
import sklearn.cluster as cluster  
from sklearn import datasets  
from sklearn.cluster import KMeans
```

```
x, y = make_blobs(n_samples=500, centers=4,  
cluster_std=3, random_state=42)
```

```
print("Shape of x: ", x.shape)
```

```
print("Shape of y: ", y.shape)
```

```
K-means = KMeans(n_clusters=4, random_state=42)  
K-means.fit(x)
```

```
labels = K-means.labels
```

```
plt.figure(figsize=(8,6))
plt.scatter(x=[0,0], y=[0,1], c='blue', cmap='viridis')
plt.scatter(means.cluster_centers_[:,0], means.cluster_centers_[:,1], s=100, c="red", label="centers")
```

= 'centers')

plt.title('K means clustering')

```
plt.xlabel('x')
plt.ylabel('y')
plt.show()
```

Output: Random Forest classifier
Random Forest classifier (criterion = 'entropy',
n_estimators = 10)

confusion matrix

| | | | | |
|---|---|----|----|----|
| [| [| 19 | 0 | 0] |
| [| 0 | 13 | 0 | 3] |
| [| 0 | 0 | 13 |] |

12. Ensemble Method Bagging

```
import pandas as pd
import numpy as np
from sklearn. datasets import load_iris
from sklearn. model_selection import train_test_split
from sklearn. metrics import accuracy_score
```

```
iris = load_iris()
```

```
X = iris. data
```

```
y = iris. target
```

```
X_train, X_test, y_train, y_test = train_test_split(X,
y, test_size=0.3, random_state=42)
```

```
from sklearn. ensemble import RandomForestClassifier
dt = RandomForestClassifier(n_estimators=10,
                            criterion="entropy")
```

```
dt. fit(X_train, y_train)
```

```
y_pred = dt. predict(X_test)
```

```
from sklearn. metrics import classification_report,
confusion_matrix, accuracy_score
result1 = confusion_matrix(y_true, y_pred)
print ("Confusion matrix")
print(result1)
```

Implementation Report

| Iteration | Wall | fl-solo | Support |
|-----------|------|---------|---------|
| 1 | 1.00 | 1.00 | 1.9 |
| 2 | 1.00 | 1.00 | 1.3 |
| 3 | 1.00 | 1.00 | 1.3 |
| 4 | 1.00 | 1.00 | 4.5 |
| 5 | 1.00 | 1.00 | 4.5 |
| 6 | 1.00 | 1.00 | 4.5 |

Accuracy = 1.0

Result 2 - Classification based on test & pred
Based on "Classification Report":
Actual (positive) vs Predicted (positive)

Output : * Gradient Boosting classifier
* Gradient Boosting

* Confusion Matrix
 $\begin{bmatrix} 19 & 0 & 0 \\ 0 & 13 & 0 \\ 0 & 0 & 13 \end{bmatrix}$

* Classification Report

| | Precision | Recall | f1-score | Support |
|---|-----------|--------|----------|---------|
| 0 | 1.00 | 1.00 | 1.00 | 19 |
| 1 | 1.00 | 1.00 | 1.00 | 13 |
| 2 | 1.00 | 1.00 | 1.00 | 13 |

accuracy

macro avg

weighted avg

1.00
1.00
1.00
1.00
1.00

45
45
45

13 Ensemble method - Boosting

```
from sklearn.ensemble import AdaBoostClassifier
from sklearn.ensemble import GradientBoostingClassifier
from sklearn.model_selection import train_test_split
from sklearn.metrics import accuracy_score, classification_report, confusion_matrix
```

```
iris = load_iris()
```

```
X = iris.data
```

```
y = iris.target
```

```
(X_train, X_test, y_train, y_test) = train_test_split(X, y, test_size=0.3, random_state=42)
```

```
from sklearn.ensemble import GradientBoostingClassifier
```

```
gb = GradientBoostingClassifier()
```

```
gb.fit(X_train, y_train)
```

```
y_pred = gb.predict(X_test)
```

```
result = confusion_matrix(y_test, y_pred)
```

```
print ("Confusion Matrix")
```

```
print(result)
```

```
result1 = classification_report(y_test, y_pred)
```

```
print ("Classification Report")
```

```
print(result1)
```

140 Ensemble Method - Stacking

```

from sklearn.datasets import load_iris
from sklearn.ensemble import StackingClassifier
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LogisticRegression
from sklearn.svm import SVC
from sklearn.metrics import accuracy_score
from sklearn.metrics import confusion_matrix

```

```

iris = load_iris()
X = iris.data
y = iris.target
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=42)
base_learners = [DecisionTreeClassifier(max_depth=1),
                 LogisticRegression()]
stack = StackingClassifier(base_learners=base_learners,
                           final_estimator=SVC())
stack.fit(X_train, y_train)
print("Accuracy: ", accuracy_score(y_test, stack.predict(X_test)))

```

```
y_pred = stack_y4 . predict(x_test)
```

```
print ("Stacking model accuracy : ", accuracy_score(y_test, y_pred))
```

```
result = confusion_matrix(y_test, y_pred)
print ("Confusion matrix : ")
print (result)
```

Output:

Stacking model accuracy : 1.0

Confusion matrix:

$$\begin{bmatrix} 0 & 19 & 0 & 0 \\ 1 & 0 & 13 & 0 \\ 2 & 0 & 0 & 13 \end{bmatrix}$$

Classification Report

| | precision | recall | f1-score | support |
|---|-----------|--------|----------|---------|
| 0 | 1.00 | 1.00 | 1.00 | 19 |
| 1 | 1.00 | 1.00 | 1.00 | 13 |
| 2 | 1.00 | 1.00 | 1.00 | 13 |

| | accuracy | | | |
|--------------|----------|------|------|----|
| macro avg | 1.00 | 1.00 | 1.00 | 45 |
| weighted avg | 1.00 | 1.00 | 1.00 | 45 |

y_pred = stackclf.predict(x_test)

print ("Stacking model accuracy:", accuracy_score(y_test, y_pred))

result = confusion_matrix(y_test, y_pred)
print ("Confusion matrix :")
print (result)