

```
# -*- coding: utf-8 -*-  
"""ML_Assessment4.ipynb
```

Automatically generated by Colab.

Original file is located at

<https://colab.research.google.com/drive/1C4vgZs4XV7F6RRUaxHWuGQzbxBByfui1>

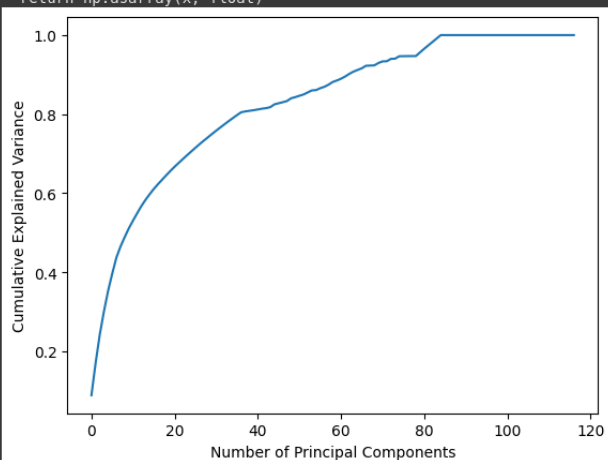
```
import numpy as np  
import pandas as pd  
import matplotlib.pyplot as plt  
import seaborn as sns  
from sklearn.cluster import AgglomerativeClustering  
from sklearn.decomposition import PCA  
from sklearn.ensemble import RandomForestClassifier, AdaBoostClassifier  
from sklearn.model_selection import train_test_split  
from sklearn.metrics import accuracy_score, classification_report,  
confusion_matrix  
from sklearn.preprocessing import StandardScaler  
from scipy.cluster.hierarchy import dendrogram, linkage  
  
from sklearn.datasets import load_iris
```

```
from sklearn.tree import DecisionTreeClassifier
```

1. Hierarchical Clustering on Iris Dataset

```
df_iris = pd.DataFrame(load_iris().data, columns=load_iris().feature_names)  
linkage_matrix = linkage(df_iris, method='ward')  
plt.figure(figsize=(10, 5))  
dendrogram(linkage_matrix)  
plt.title("Dendrogram for Hierarchical Clustering")  
plt.show()
```

```
/usr/local/lib/python3.11/dist-packages/matplotlib/ctbook.py:1709: ComplexWarning: Casting complex values to real discards the imaginary part  
return math.isfinite(val)  
/usr/local/lib/python3.11/dist-packages/matplotlib/ctbook.py:1345: ComplexWarning: Casting complex values to real discards the imaginary part  
return np.asarray(x, float)
```



2. PCA on Mushroom Dataset

```

try:
dataset_url =
"https://archive.ics.uci.edu/ml/machine-learning-databases/mushroom/
agaricus-lepiota.data"
column_names = ["class", "cap-shape", "cap-surface", "cap-color", "bruises",
"odor", "gill-attachment", "gill-spacing", "gill-size", "gill-color",
"stalk-shape", "stalk-root", "stalk-surface-above-ring", "stalk-surface-
below-ring", "stalk-color-above-ring", "stalk-color-below-ring", "veil-
type", "veil-color", "ring-number", "ring-type", "spore-print-color",
"population", "habitat"]
df_mushroom = pd.read_csv(dataset_url, names=column_names)
X_mushroom = pd.get_dummies(df_mushroom.drop('class', axis=1))
scaler = StandardScaler()
X_scaled = scaler.fit_transform(X_mushroom)
cov_matrix = np.cov(X_scaled.T)
eigenvalues, eigenvectors = np.linalg.eig(cov_matrix)
explained_variance = eigenvalues / np.sum(eigenvalues)
cumulative_variance = np.cumsum(explained_variance)
k = np.argmax(cumulative_variance >= 0.95) + 1 # Retaining 95% variance
pca = PCA(n_components=k)
X_pca = pca.fit_transform(X_scaled)
plt.plot(cumulative_variance)
plt.xlabel("Number of Principal Components")
plt.ylabel("Cumulative Explained Variance")
plt.show()
except Exception as e:
print(f"Error loading mushroom dataset: {e}")

```

```

Random Forest Accuracy: 0.9927272727272727
      precision    recall  f1-score   support

     0       0.99      1.00      0.99        148
     1       1.00      0.98      0.99        127

 accuracy          0.99          0.99          0.99        275
 macro avg         0.99          0.99          0.99        275
 weighted avg      0.99          0.99          0.99        275

```

3. Random Forest on Bill Authentication Dataset

```

Authentication Dataset
try:
df_bill = pd.read_csv("https://archive.ics.uci.edu/ml/machine-learning-
databases/00267/data_banknote_authentication.txt", names=["Variance",
"Skewness", "Curtosis", "Entropy", "Class"])
X_bill = df_bill.drop('Class', axis=1)
y_bill = df_bill['Class']
X_train, X_test, y_train, y_test = train_test_split(X_bill, y_bill,
test_size=0.2, random_state=42)

rf_model = RandomForestClassifier(n_estimators=100, random_state=42)
rf_model.fit(X_train, y_train)

```

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

```
print("Random Forest Accuracy:", accuracy_score(y_test, y_pred))
```

```
print(classification_report(y_test, y_pred))
```

```
except FileNotFoundError:
```

```
print("Error: bill_authentication.csv not found. Please provide the dataset.")
```

```
AdaBoost Accuracy: 0.9666666666666667
```

	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

4. AdaBoost for Flower Species Classification

```
X_train, X_test, y_train, y_test = train_test_split(load_iris().data,  
load_iris().target, test_size=0.2, random_state=1)
```

```
adaboost = AdaBoostClassifier(estimator=DecisionTreeClassifier(max_depth=2),  
n_estimators=50, random_state=1)
```

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

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

```
print("AdaBoost Accuracy:", accuracy_score(y_test, y_pred))
```

```
print(classification_report(y_test, y_pred))
```

```
Decision Tree Accuracy: 0.9667  
Random Forest Accuracy: 0.9667  
AdaBoost Accuracy: 0.9667
```

5. Comparison of Classification Algorithms

```
models = {
```

```
"Decision Tree": DecisionTreeClassifier(),
```

```
"Random Forest": RandomForestClassifier(n_estimators=100),
```

```
"AdaBoost":
```

```
AdaBoostClassifier(estimator=DecisionTreeClassifier(max_depth=1),  
n_estimators=50)
```

```
}
```

```
for name, model in models.items():
```

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

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

```
print(f"{name} Accuracy: {accuracy_score(y_test, y_pred):.4f}")
```