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ASSIGNMENT NAME: KNN & PCA

DRIVE LINK: <u>link</u>

Assignment Code: DA-AG-016

KNN & PCA Assignment

Instructions:Carefully readeach question. UseGoogleDocs, Microsoft Word, or a similar tool to create a document where you type out each question along with its answer. Save the document as a PDF, and then upload it to the LMS. Please do not zip or archive the files before uploading them. Each question carries 20 marks.

Total Marks: 200

Question 1: What is K-Nearest Neighbors (KNN) and how does it work in both classification and regression problems?

Answer:

K-Nearest Neighbors is a supervised machine learning algorithm used for classification and regression tasks. Its called lazy learning because it doesn't explicitly build a model during training it just stores the data and makes predictions at query time.

*In classification, prediction is based on the majority class of neighbors.

*In regression, prediction is based on the average value of neighbors.

Question 2: What is the Curse of Dimensionality and how does it affect KNN performance?

Answer:

The curse of dimensionality refers to the problems that arise when working with high-dimensional data.

- *Nearest neighbor becomes less meaningful
- *Increased risk of overfitting
- *Higher computational cost



Question 3: What is Principal Component Analysis (PCA)? How is it different from feature selection?

Answer:

Principal Component Analysis(PCA) is a dimensionally reduction technique that transforms the original features into a new set of features called principal components.

Feature selection- Selecting a subset of the original feature that are most relevant. Keeps some of the original feature.

Remove irrelevant/redundant features.

Easy to interpret.

Question 4: What are eigenvalues and eigenvectors in PCA, and why are they important?

Answer:

 $Av = \lambda v$

Here v = eigenvector

 λ = eigenvalue (a scalar)

Importance:

- *Eigenvectors define principal components
- *Eigenvalues tell us the importance
- *Dimensionally reduction

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Question5: How do KNN and PCA complement each other when applied in a single pipeline?

Answer:

- *PCA transforms high-dimensional data into a smaller set of informative features.
- *KNN then classifies or predicts based on distance in this reduced space.
- *Both complement each other because PCA reduces dimensionally and makes distance more meaningful, while KNN use those distance for prediction.
- *The combo reduces curse of dimensionality, improves efficiency, and often increases accuracy.



Dataset:

Use the **Wine Dataset** from sklearn.datasets.load_wine().

Question 6: Train a KNN Classifier on the Wine dataset with and without feature scaling. Compare model accuracy in both cases.

(Include your Python code and output in the code box below.

Answer:

```
from sklearn.datasets import load_wine
from sklearn.model_selection import train_test_split
from sklearn.melghbors import KleighborsClassifier
from sklearn.preprocessing import StandardScaler
from sklearn.priperior tstandardScaler
from sklearn.priperior tstandardScaler
from sklearn.priperior tstandardScaler
from sklearn.priperior tstandardScaler
x, y = load_wine(return_X_y=True)
X_tr, X_te, y_tr, y_te = train_test_split(
X, y, test_size=0.30, random_state=42, stratify=y
)
knn_plain = KNeighborsClassifier(n_neighbors=5)
knn_plain.fit(X_tr, y_tr)
acc_no = knn_plain.score(X_te, y_te)
knn_scaled = make_plain(standardScaler(),
KNeighborsClassifier(n_neighbors=5))
knn_scaled = make_plain(standardScaler(),
KNeighborsClassifier(n_neighbors=5))
knn_scaled.fit(X_tr, y_tr)
acc_yes = knn_scaled.score(X_te, y_te)
print(f"Accuracy without scaling: {acc_no:.3f}")
print(f"Accuracy with scaling: {acc_no:.3f}")

Accuracy without scaling: 0.722
Accuracy without scaling: 0.944

Activate Windows
```

Question 7:Train a PCA model on the Wine dataset and print the explained variance

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ratio of each principal component.

(Include your Python code and output in the code box below.)

Answer:

```
from sklearn.datasets import load_wine from sklearn.preprocessing import StandardScaler from sklearn.decomposition import PCA wine = load_wine()

X, y = wine.data, wine.target

Y coalad = StandardScaler().fit_transform(X)
                          X_scaled = StandardScaler().fit_transform(X)
pca = PCA()
 ⊙⊽
                          pca.fit(X scaled)
                         print("Explained Variance Ratio for each Principal Component:")
for i, ratio in enumerate(pca.explained_variance_ratio_):
print(f"PC{i+1}: {ratio:.4f}")
print("\nCumulative Explained Variance:")
                          print(pca.explained_variance_ratio_.cumsum())
               Explained Variance Ratio for each Principal Component:
PC1: 0.3620
PC2: 0.1921
PC3: 0.1921
PC4: 0.0707
PC5: 0.0656
PC6: 0.0494
PC7: 0.0424
PC8: 0.0222
PC10: 0.0193
                Explained Variance Ratio for each Principal Component:
 Q
                          Explained Var
PC1: 0.3620
PC2: 0.1921
PC3: 0.1912
PC3: 0.0556
PC5: 0.0656
PC6: 0.0494
PC7: 0.0424
PC8: 0.0226
PC9: 0.0222
PC10: 0.0139
PC11: 0.0134
PC12: 0.0130
 <>
 0
PC13: 0.0080
                          Cumulative Explained Variance:

[0.36198848 0.55406338 0.66529969 0.73598999 0.80162293 0.85098116

0.89336795 0.92017544 0.94239698 0.96169717 0.97906553 0.99204785
```

Question 8: Train a KNN Classifier on the PCA-transformed dataset (retain top 2 components). Compare the accuracy with the original dataset. (*Include your Python code and output in the code box below.*)

Answer:



```
from sklearn.datasets import load_wine
from sklearn.model_selection import train_test_split
from sklearn.neighbors import KNeighborsClassifier
from sklearn.preprocessing import StandardScaler
from sklearn.decomposition import PCA
from sklearn.pipeline import make_pipeline
X, y = load_wine(return_X_y=True)
Y train_Y = Train_test_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_select_s
 <>
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X_train, X_test, y_train, y_test = train_test_split(
    X, y, test_size=0.3, random_state=42,
                                                                      stratify=y
                                                      scaler = StandardScaler()
                                                      X_train_scaled = scaler.fit_transform(X_train)
X_test_scaled = scaler.transform(X_test)
                                                      knn = KNeighborsClassifier(n neighbors=5)
                                                      knn.fit(X_train_scaled, y_train)
y_pred_no_pca = knn.predict(X_test_scaled)
                                                      acc_no_pca = accuracy_score(y_test, y_pred_no_pca)
pca = PCA(n_components=2)
                                                      X_train_pca = pca.fit_transform(X_train_scaled)
X_test_pca = pca.transform(X_test_scaled)
                                                     knn_pca = KNeighborsClassifier(n_neighbors=5)
                                                     /scaler = StandardScaler()
X_train_scaled = scaler.fit_transform(X_train)
X_test_scaled = scaler.transform(X_test)
knn = KNeighborsClassifier(n_neighbors=5)
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                                                    knn.fit(X_train_scaled, y_train)
y_pred_no_pca = knn.predict(X_test_scaled)
                                                    y_preu_no_pca = xim.preuzct(z_test_scaled)
acc_no_pca = accuracy_score(y_test, y_pred_no_pca)
pca = PCA(n_components=2)
X_train_pca = pca.fit_transform(X_train_scaled)
X_test_pca = pca.transform(X_test_scaled)
knn_pca = KNeighborsClassifier(n_neighbors=5)
knn_pca.fit(X_train_pca, y_train)
y_pred_pca = knn_pca.predict(X_test_pca)
                                                      acc_pca = accuracy_score(y_test, y_pred_pca)
print(f"Accuracy with original scaled data:{acc_no_pca:.3f}")
                                                      print(f"Accuracy with PCA (2 components):{acc_pca:.3f}")
                                 Accuracy with original scaled data:0.944
Accuracy with PCA (2 components):0.944
```

Question 9: Train a KNN Classifier with different distance metrics (euclidean, manhattan) on the scaled Wine dataset and compare the results. (*Include your Python codeandoutputinthecodeboxbelow.*)

Answer:

```
| Staler = StandardScaler()
| X_train_scaled = scaler.fit_transform(X_train)
| X_test_scaled = scaler.transform(X_test)
| metrics = ['euclidean', 'manhattan']
| result = ()
| for metric in metrics:
| knn = KNeighborsClassifier(n_neighbors=5,
| metric=metric)
| knn.fit(X_train_scaled, y_train)
| y_pred = knn.predict(X_test_scaled)
| acc = accuracy_score(y_test, y_pred)
| result[metric] = acc
| for metric, acc in result.items():
| print(f'KNN with (metric) distance Accuracy: (acc:.3f)")
| This with euclidean distance Accuracy: 0.944
| KNN with manhattan distance Accuracy: 0.981
```

Question 10: You are working with a high-dimensional gene expression dataset to classify patients with different types of cancer.

Due to the large number of features and a small number of samples, traditional models overfit.

Explain how you would:

- Use PCA to reduce dimensionality
- Decide how many components to keep
- Use KNN for classification post-dimensionality reduction
- Evaluate the model
- Justify this pipeline to your stakeholders as a robust solution for real-world biomedical data

(Include your Python code and output in the code box below.

Answer:



Accuracy: 0.19

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

| [34] import numpy as np | from sklearn.datasets import make_classification | from sklearn.decomposition import PCA | from sklearn.mode_selection import extandardScaler | from sklearn.mode_selection import train_test_split | from sklearn.meds_selection import train_test_split | from sklearn.meds_selection import train_test_split | from sklearn.metrics import accuracy_score | X, y = make_classification(n_samples=500, n_features=2000, n_informative=50, n_classes=5, random_state=42) | X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42) | x_train_scaled = scaler.fit_transform(X_train) | X_test_scaled = scaler.transform(X_test) | pca = PCA(n_components=50) | X_train_pca = pca.fit_transform(X_train_scaled) | X_test_pca = pca.transform(X_train_scaled) | x_test_pca = pca.transform(X_train_

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