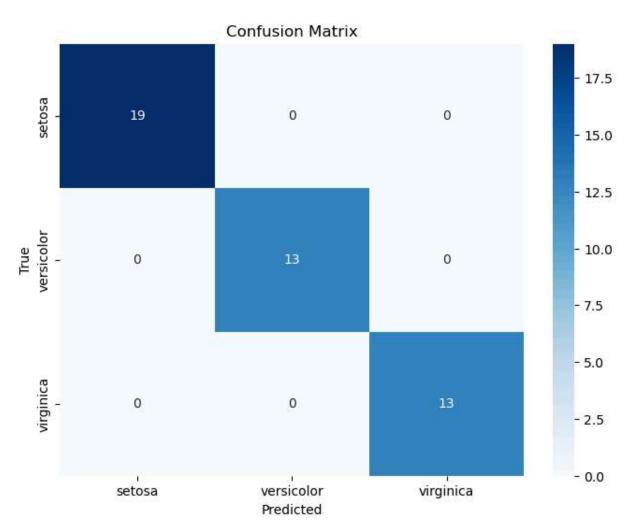
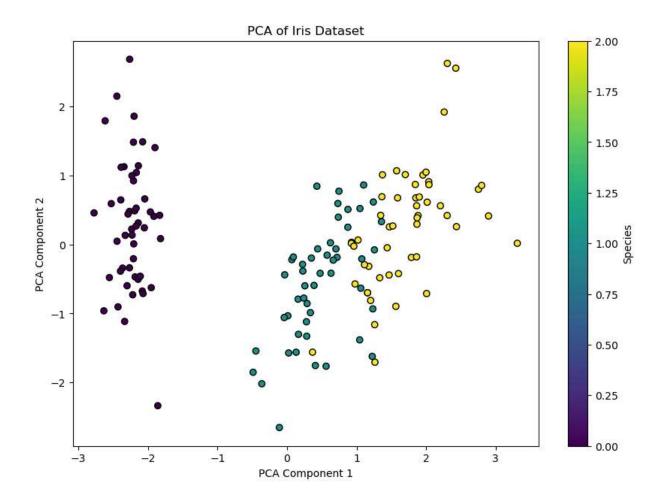
```
In [1]: import pandas as pd
        import numpy as np
        import seaborn as sns
        import matplotlib.pyplot as plt
        from sklearn.datasets import load iris
        from sklearn.model selection import train test split
        from sklearn.preprocessing import StandardScaler
        from sklearn.decomposition import PCA
        from sklearn.linear model import LogisticRegression
        from sklearn.metrics import classification_report, confusion_matrix
        # Load Iris dataset
        iris = load iris()
        X = iris.data
        y = iris.target
        feature_names = iris.feature_names
        target names = iris.target names
        # Create DataFrame
        df = pd.DataFrame(X, columns=feature names)
        df['species'] = y
        # Preprocess Data
        scaler = StandardScaler()
        X_scaled = scaler.fit_transform(X)
        # Split Data
        X_train, X_test, y_train, y_test = train_test_split(X_scaled, y, test_size=0.3, ran
        # Train Model
        model = LogisticRegression(max_iter=200)
        model.fit(X_train, y_train)
        # Evaluate Model
        y_pred = model.predict(X_test)
        print("Classification Report:")
        print(classification report(y test, y pred, target names=target names))
        # Confusion Matrix
        cm = confusion_matrix(y_test, y_pred)
        plt.figure(figsize=(8, 6))
        sns.heatmap(cm, annot=True, fmt='d', cmap='Blues', xticklabels=target names, ytickl
        plt.xlabel('Predicted')
        plt.ylabel('True')
        plt.title('Confusion Matrix')
        plt.show()
        # PCA for Visualization
        pca = PCA(n components=2)
        X_pca = pca.fit_transform(X_scaled)
        plt.figure(figsize=(10, 7))
```

```
scatter = plt.scatter(X_pca[:, 0], X_pca[:, 1], c=y, cmap='viridis', edgecolor='k')
plt.xlabel('PCA Component 1')
plt.ylabel('PCA Component 2')
plt.title('PCA of Iris Dataset')
plt.colorbar(scatter, label='Species')
plt.show()
```

Classification Report:

	precision	recall	f1-score	support
setosa	1.00	1.00	1.00	19
versicolor	1.00	1.00	1.00	13
virginica	1.00	1.00	1.00	13
2661112614			1.00	45
accuracy			1.00	45
macro avg	1.00	1.00	1.00	45
weighted avg	1.00	1.00	1.00	45





wine quality classification

In []:

```
In [1]: import pandas as pd
        import numpy as np
        import seaborn as sns
        import matplotlib.pyplot as plt
        from sklearn.model_selection import train_test_split
        from sklearn.preprocessing import StandardScaler
        from sklearn.decomposition import PCA
        from sklearn.ensemble import RandomForestClassifier
        from sklearn.metrics import classification report, confusion matrix
        # Load Wine Quality dataset
        url = "https://archive.ics.uci.edu/ml/machine-learning-databases/wine-quality/wineq
        df = pd.read csv(url, sep=';')
        # Preview the dataset
        print("First 5 rows of the dataset:")
        print(df.head(10))
        # Binary Classification: Good (quality >= 6) or Bad (quality < 6)
        df['quality'] = df['quality'].apply(lambda x: 1 if x >= 6 else 0)
        # Features and Target
        X = df.drop('quality', axis=1)
```

```
y = df['quality']
# Preprocess Data
scaler = StandardScaler()
X_scaled = scaler.fit_transform(X)
# Split Data
X_train, X_test, y_train, y_test = train_test_split(X_scaled, y, test_size=0.3, ran
# Train Model
model = RandomForestClassifier(n_estimators=100, random_state=42)
model.fit(X train, y train)
# Evaluate Model
y pred = model.predict(X test)
print("\nClassification Report:")
print(classification_report(y_test, y_pred))
# Confusion Matrix
cm = confusion_matrix(y_test, y_pred)
plt.figure(figsize=(8, 6))
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues', xticklabels=['Bad', 'Good'], yti
plt.xlabel('Predicted')
plt.ylabel('True')
plt.title('Confusion Matrix')
plt.show()
# PCA for Visualization
pca = PCA(n_components=2)
X_pca = pca.fit_transform(X_scaled)
plt.figure(figsize=(10, 7))
scatter = plt.scatter(X_pca[:, 0], X_pca[:, 1], c=y, cmap='viridis', edgecolor='k')
plt.xlabel('PCA Component 1')
plt.ylabel('PCA Component 2')
plt.title('PCA of Wine Quality Dataset')
plt.colorbar(scatter, label='Quality')
plt.show()
```

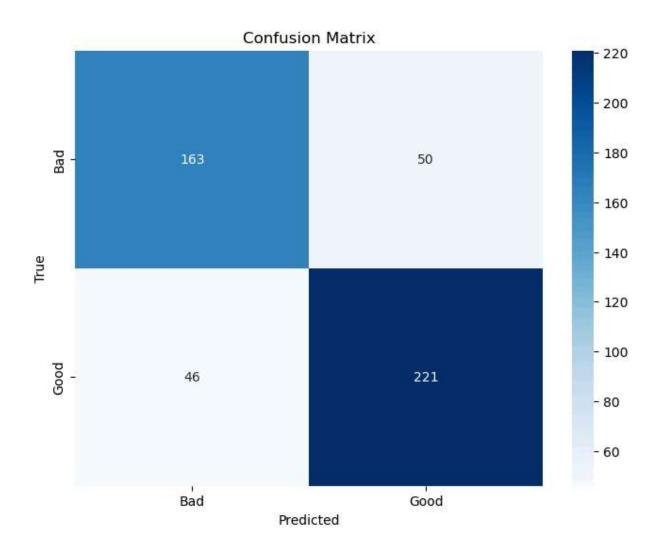
First 5 rows of the dataset:						
	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	\
0	7.4	0.70	0.00	1.9	0.076	
1	7.8	0.88	0.00	2.6	0.098	
2	7.8	0.76	0.04	2.3	0.092	
3	11.2	0.28	0.56	1.9	0.075	
4	7.4	0.70	0.00	1.9	0.076	
5	7.4	0.66	0.00	1.8	0.075	
6	7.9	0.60	0.06	1.6	0.069	
7	7.3	0.65	0.00	1.2	0.065	
8	7.8	0.58	0.02	2.0	0.073	
9	7.5	0.50	0.36	6.1	0.071	

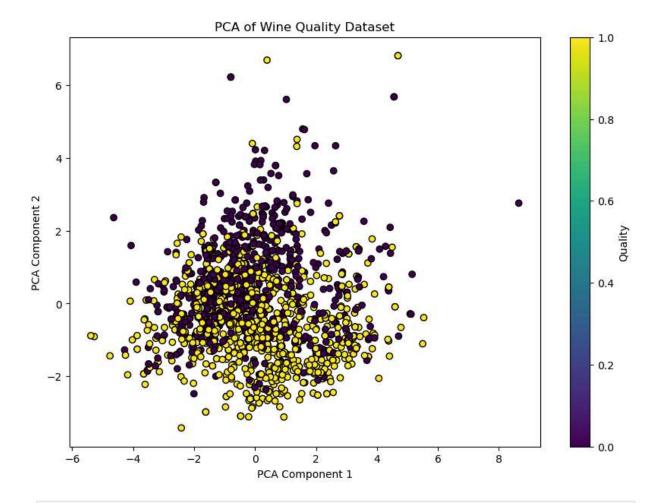
0 11 0	34.0 0.9978	3.51	0 50
0 11.0			0.56
1 25.0	57.0 0.9968	3.20	0.68
2 15.0	54.0 0.9970	3.26	0.65
3 17.0	50.0 0.9980	3.16	0.58
4 11.0	34.0 0.9978	3.51	0.56
5 13.0	10.0 0.9978	3.51	0.56
6 15.0	59.0 0.9964	3.30	0.46
7 15.0	21.0 0.9946	3.39	0.47
9.0	18.0 0.9968	3.36	0.57
9 17.0 1	0.9978	3.35	0.80

	alcohol	quality
0	9.4	5
1	9.8	5
2	9.8	5
3	9.8	6
4	9.4	5
5	9.4	5
6	9.4	5
7	10.0	7
8	9.5	7
9	10.5	5

Classification Report:

	precision	recall	f1-score	support
0	0.78	0.77	0.77	213
1	0.82	0.83	0.82	267
accuracy			0.80	480
macro avg	0.80	0.80	0.80	480
weighted avg	0.80	0.80	0.80	480





In []: