Pistacho

To create a classification model for the Pistachio Dataset, you'll need to identify the most important features. These important features are the ones that significantly contribute to distinguishing between the classes in your dataset. The dataset consists of features related to shape, size, and geometry of the pistachio seeds.

Data Loading and Description

Here's how you can identify important features and select them:

Understanding the Features

- 1. AREA: The area of the pistachio seed.
- 2. PERIMETER: The perimeter of the seed.
- 3. MAJOR_AXIS: Length of the major axis of the seed.
- 4. MINOR_AXIS: Length of the minor axis of the seed.
- 5. ECCENTRICITY: A measure of how much the shape of the pistachio deviates from being circular.
- 6. EQDIASQ: Equivalent diameter, calculated from the area.
- 7. SOLIDITY: Ratio of the area to the convex area, indicating how solid or compact the shape is.
- 8. CONVEX_AREA: The area of the convex hull (the smallest convex shape that encloses the seed).
- 9. EXTENT: The ratio of the area of the seed to the bounding box.
- 10. ASPECT_RATIO: Ratio of the major axis to the minor axis.
- 11. ROUNDNESS: A measure of how circular the shape is.
- 12. COMPACTNESS: A measure of how compact or closely packed the shape is.
- 13. SHAPEFACTOR_1 to SHAPEFACTOR_4: Various shape factors indicating different shape characteristics.
- 14. Class: The label or target for classification (e.g., different types of pistachios).

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

→ Mounted at /content/drive

Importing Libraries

In this section, we are importing the necessary libraries for data processing and modeling.

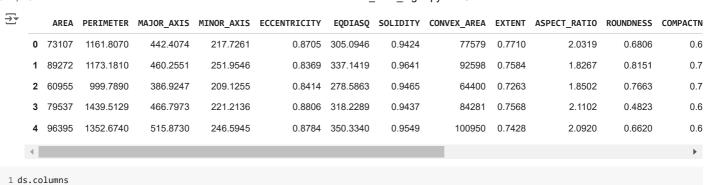
```
1 import numpy as np
2 import pandas as pd
3 import matplotlib.pyplot as plt
4 import seaborn as sns
5 %matplotlib inline
6 import warnings
7 warnings.filterwarnings('ignore')
8 pd.set_option('display.max_columns', 100)
```

1 ds = pd.read_csv("/content/drive/MyDrive/pistachio.csv")

Understanding the Dataset

To gain insights from data we must look into each aspect of it very carefully. We will start with observing few rows and columns of data both from the starting and from the end

```
1 ds.head()
```



The dataset contains 1718 entries and 17 columns. All columns, except for the "Class" label, are numeric. Here's a breakdown of the columns:

Numerical Features (16):

AREA, PERIMETER, MAJOR_AXIS, MINOR_AXIS, ECCENTRICITY, EQDIASQ, SOLIDITY, CONVEX_AREA, EXTENT, ASPECT_RATIO, ROUNDNESS, COMPACTNESS, SHAPEFACTOR_1, SHAPEFACTOR_2, SHAPEFACTOR_3, SHAPEFACTOR_4 Categorical Feature (1):

Class: This is the target label with categories such as "Kirmizi_Pistachio" and "Siit_Pistachio".

1 ds.info()

<pr RangeIndex: 1718 entries, 0 to 1717 Data columns (total 17 columns): # Column Non-Null Count Dtype AREA 1718 non-null int64 PERIMETER 1718 non-null float64 1 MAJOR AXIS 1718 non-null float64 MTNOR AXTS 1718 non-null float64 ECCENTRICITY 1718 non-null float64 5 **EQDIASQ** 1718 non-null float64 6 SOLIDITY 1718 non-null float64 CONVEX_AREA 1718 non-null int64 8 EXTENT 1718 non-null float64 ASPECT_RATIO 1718 non-null float64 9 10 ROUNDNESS 1718 non-null float64 COMPACTNESS 1718 non-null float64 11 SHAPEFACTOR 1 1718 non-null float64 12 SHAPEFACTOR 2 1718 non-null float64 13 14 SHAPEFACTOR_3 1718 non-null float64 15 SHAPEFACTOR_4 1718 non-null float64 16 Class 1718 non-null object dtypes: float64(14), int64(2), object(1) memory usage: 228.3+ KB

1 ds.describe(include	=	"all")
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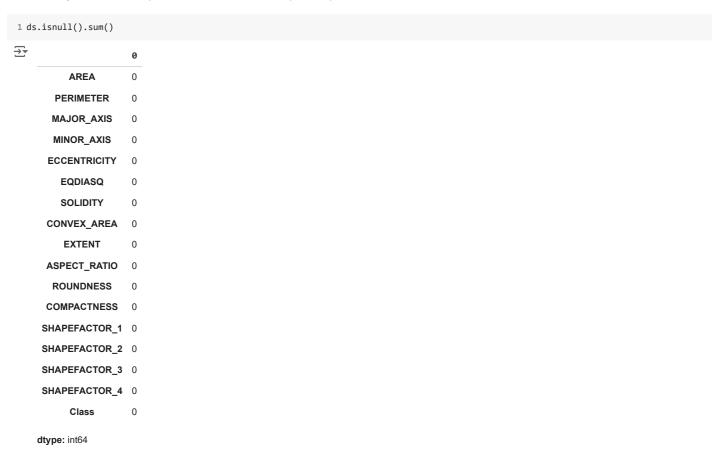
→

	AREA	PERIMETER	MAJOR_AXIS	MINOR_AXIS	ECCENTRICITY	EQDIASQ	SOLIDITY	CONVEX_AREA	EXTENT	ASPEC.
count	1718.000000	1718.000000	1718.000000	1718.000000	1718.000000	1718.000000	1718.000000	1718.000000	1718.000000	1718
unique	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	
top	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	
freq	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	
mean	79871.952852	1421.797588	446.206444	238.193128	0.840347	317.790000	0.940103	84947.671129	0.716055	1
std	12968.217051	373.408835	31.885328	30.426445	0.049026	26.571699	0.050006	13081.742551	0.052534	0
min	29808.000000	858.363000	321.425500	133.509600	0.504900	194.814600	0.588000	37935.000000	0.427200	1
25%	71898.500000	1169.633225	426.554100	217.875475	0.817500	302.562375	0.920250	76357.750000	0.688100	1
50%	79795.000000	1260.785500	448.453150	235.888750	0.850250	318.744650	0.953800	84973.000000	0.726100	1
75%	88980.000000	1599.479000	467.515200	257.433625	0.875375	336.590000	0.976300	93660.750000	0.753600	2
max	124008.000000	2755.049100	535.642200	383.046100	0.946000	397.356100	0.995100	132478.000000	0.820400	3
4										•

Checking for Null Values

In this step, we check for any missing or null values in the dataset. It's important to ensure data completeness before proceeding with data analysis and model building. If there are null values, appropriate handling methods such as imputation or removal will be applied.

After running the code, it has been confirmed that **there are no null values** in the dataset. This means that the dataset is clean and ready for further analysis and modeling without the need for handling missing data.



I'll start by generating some EDA visualizations and handle preprocessing. Let's begin with visualizing distributions and correlations of features.

FDA Observations:

Distributions:

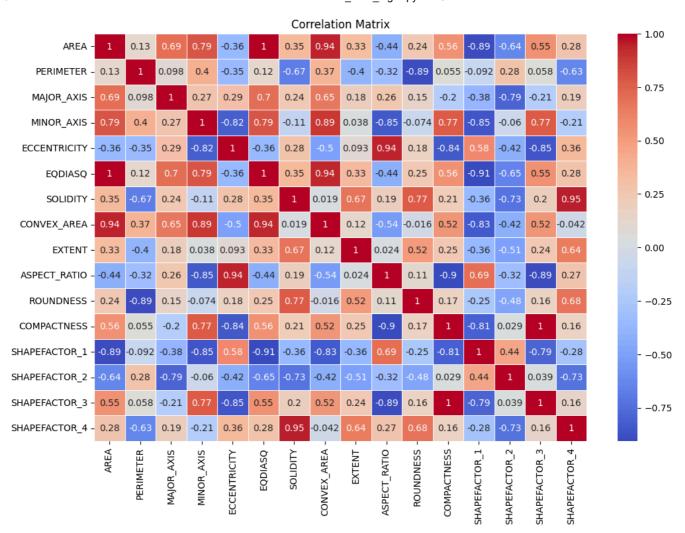
- 1. Many of the features seem skewed, with a few possibly containing outliers.
- 2. Most of the features exhibit variability in their distributions, which will be useful for classification.

Correlation Heatmap:

Several features show strong correlations, such as PERIMETER and AREA, as well as MAJOR_AXIS and MINOR_AXIS. These correlations might be useful in model feature selection or could lead to multicollinearity issues, which we'll address.

```
1 import seaborn as sns
2 import matplotlib.pyplot as plt
3
4 # Compute the correlation matrix
5 X = ds.drop(['Class'], axis=1)
6 correlation_matrix = X.corr()
7
8 # Plot the heatmap
9 plt.figure(figsize=(12, 8))
10 sns.heatmap(correlation_matrix, annot=True, cmap='coolwarm', linewidths=0.5)
11 plt.title('Correlation Matrix')
12 plt.show()
```

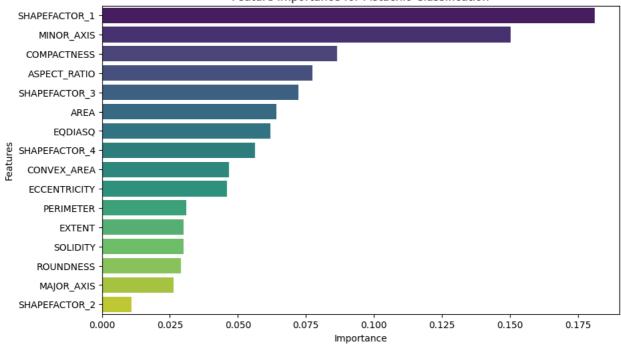




```
1
     from sklearn.ensemble import RandomForestClassifier
2
     from sklearn.preprocessing import LabelEncoder
3
     dsl = ds
     # Encode the categorical 'Class' column to numerical labels (0 and 1)
4
     label_encoder = LabelEncoder()
    dsl['Class'] = label_encoder.fit_transform(dsl['Class'])
6
8
     # Split the data into features (X) and target (y)
9
     X = dsl.drop(columns=['Class'])
10
    y = dsl['Class']
11
12
     # Train a Random Forest model to determine feature importance
13
     model = RandomForestClassifier(random_state=42)
    model.fit(X, y)
14
15
16
     # Extract feature importances
17
     feature_importances = pd.DataFrame({
18
         'Feature': X.columns,
         'Importance': model.feature_importances_
19
20
     }).sort_values(by='Importance', ascending=False)
21
22
     # Plot feature importance
     plt.figure(figsize=(10, 6))
23
     sns.barplot(x='Importance', y='Feature', data=feature_importances, palette='viridis')
24
25
     plt.title('Feature Importance for Pistachio Classification')
26
     plt.xlabel('Importance')
27
     plt.ylabel('Features')
28
     plt.show()
29
30
     # Display the feature importance data
31
     feature_importances.head()
```



Feature Importance for Pistachio Classification

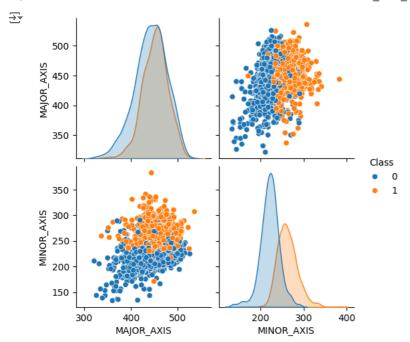


	Feature	Importance
12	SHAPEFACTOR_1	0.181197
3	MINOR_AXIS	0.150299
11	COMPACTNESS	0.086380
9	ASPECT_RATIO	0.077423
14	SHAPEFACTOR_3	0.072215

```
1 # Convert importances to percentages
2 feature_importances['Importance (%)'] = (feature_importances['Importance'] / feature_importances['Importance'].sum()) * 100
3
4 # Sort by importance percentage
5 feature_importances = feature_importances.sort_values(by='Importance (%)', ascending=False)
6
7 # Display the feature importances in percentage
8 print(feature_importances[['Feature', 'Importance (%)']])
```

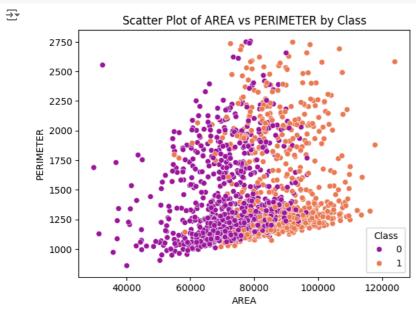
```
₹
              Feature Importance (%)
    12 SHAPEFACTOR_1
                            18.119742
           MINOR AXIS
                            15.029874
    3
          COMPACTNESS
                             8.637995
    11
         ASPECT_RATIO
                             7.742255
    9
    14 SHAPEFACTOR_3
                             7.221546
    0
                 AREA
                             6.410597
    5
              EQDIASQ
                             6.197903
    15 SHAPEFACTOR_4
                             5.633867
    7
          CONVEX_AREA
                             4.672956
         ECCENTRICITY
                             4.602906
            PERIMETER
                             3.105630
    1
               EXTENT
                             2.998207
    6
             SOLIDITY
                             2.993767
    10
            ROUNDNESS
                             2.912586
                             2.631886
           MAJOR_AXIS
       SHAPEFACTOR_2
                             1.088283
```

```
1 sns.pairplot(ds[['MAJOR_AXIS', 'MINOR_AXIS','Class']], hue='Class')
2 plt.show()
```



MINOR_AXIS: The length of the minor axis shows significant variation, especially for Siit_Pistachio, indicating that this class tends to have more elongated shapes compared to Kirmizi_Pistachio.

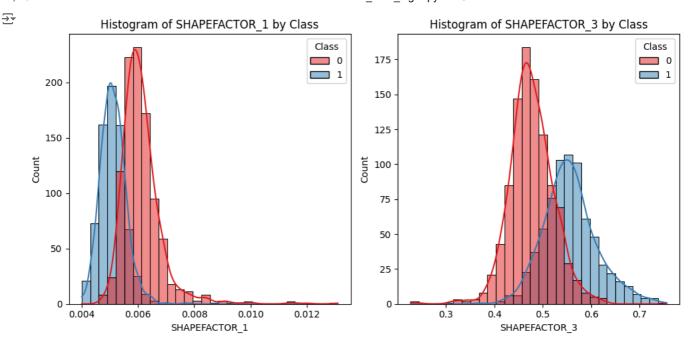
```
1 sns.scatterplot(x='AREA', y='PERIMETER', hue='Class', data=ds,palette='plasma')
2 plt.title("Scatter Plot of AREA vs PERIMETER by Class")
3 plt.show()
```



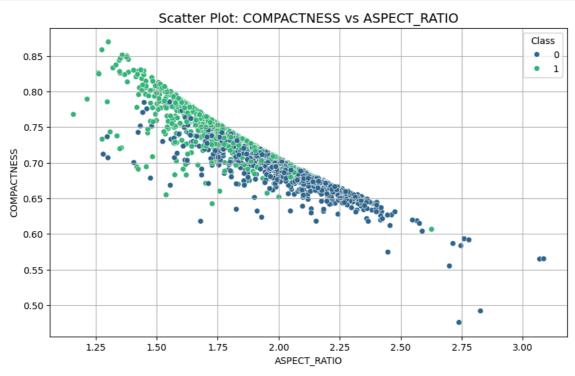
The **scatter plot** shows how the AREA and PERIMETER of the pistachios relate to each other. Siit_Pistachio and Kirmizi_Pistachio tend to cluster in different regions, indicating that these features are effective at separating the two classes. Siit_Pistachio generally has higher values for both AREA and PERIMETER, suggesting that they are larger and have a more extended perimeter.

```
1 fig, axes = plt.subplots(1, 2, figsize=(10, 5))
2
3 sns.histplot(data=ds, x='SHAPEFACTOR_1', hue='Class', kde=True, palette='Set1', bins=30, ax=axes[0])
4 axes[0].set_title("Histogram of SHAPEFACTOR_1 by Class")
5
6 sns.histplot(data=ds, x='SHAPEFACTOR_3', hue='Class', kde=True, palette='Set1', bins=30, ax=axes[1])
7 axes[1].set_title("Histogram of SHAPEFACTOR_3 by Class")
8
9 plt.tight_layout()
10 plt.show()
```

→



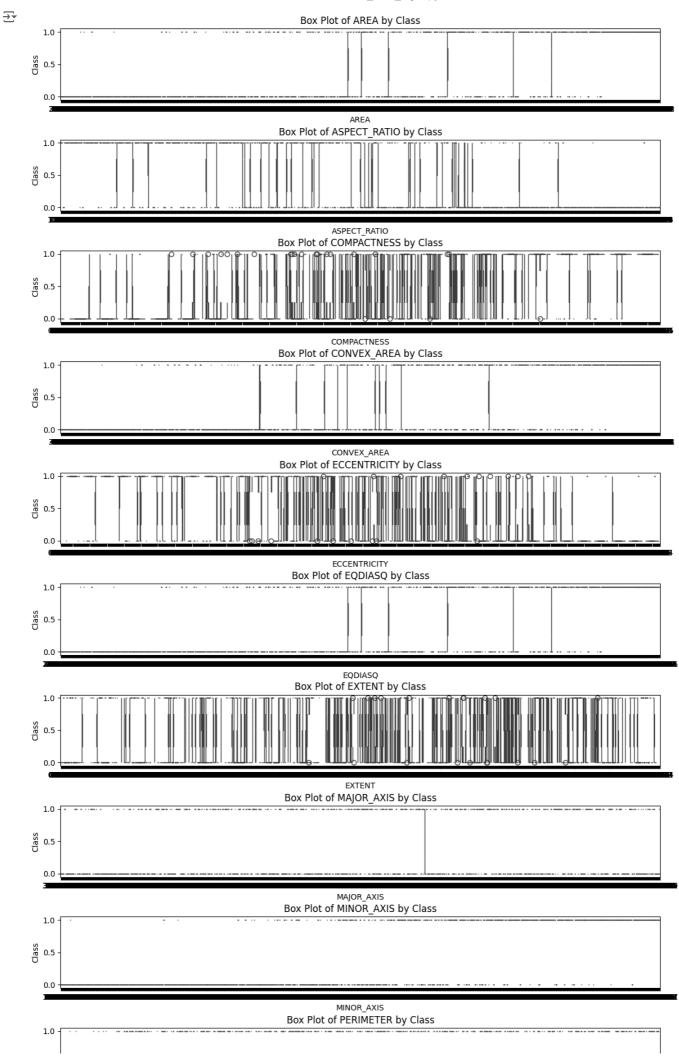
```
1 plt.figure(figsize=(10, 6))
2
3 sns.scatterplot(x=ds['ASPECT_RATIO'], y=ds['COMPACTNESS'], hue=ds['Class'], palette='viridis')
4
5 plt.title('Scatter Plot: COMPACTNESS vs ASPECT_RATIO', fontsize=14)
6 plt.xlabel('ASPECT_RATIO')
7 plt.ylabel('COMPACTNESS')
8 plt.legend(title='Class')
9 plt.grid(True)
10
11 plt.show()
```

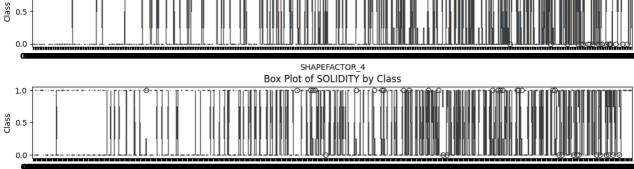


```
1 import matplotlib.pyplot as plt
2 import seaborn as sns
3
4 # Get all column names except 'Class'
5 columns_to_plot = ds.columns.difference(['Class'])
6
7 # Create subplots dynamically based on the number of columns
8 fig, axes = plt.subplots(nrows=len(columns_to_plot), figsize=(12, len(columns_to_plot)*2))
9 plt.tight_layout(pad=3.0)
10
11 # Loop through each column and plot a boxplot
12 for i, column in enumerate(columns_to_plot):
```

```
sns.boxplot(y=ds['Class'], x=ds[column], ax=axes[i]) # Set x as 'Class' and y as the feature colum
axes[i].set_title(f'Box Plot of {column} by Class')

15
16 nlt.show()
```





SOLIDITY

```
1 from sklearn.model_selection import train_test_split
2 from sklearn.preprocessing import StandardScaler
3 from sklearn.linear_model import LogisticRegression
4 from sklearn.metrics import classification_report, accuracy_score, confusion_matrix
5
6 X = ds.drop('Class', axis=1)  # Features
7 y = ds['Class']  # Target
8
9 #Split the dataset into training and testing sets
10 X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
11
12 #Feature Scaling
13 scaler = StandardScaler()
14 X_train = scaler.fit_transform(X_train)
15 X_test = scaler.transform(X_test)
16
17 X_train.shape, X_test.shape, y_train.shape, y_test.shape
```

```
((1374, 16), (344, 16), (1374,), (344,))
```

The dataset has been preprocessed as follows:

macro avg

weighted avg

0.88

0.88

0.88

0.88

0.88

0.88

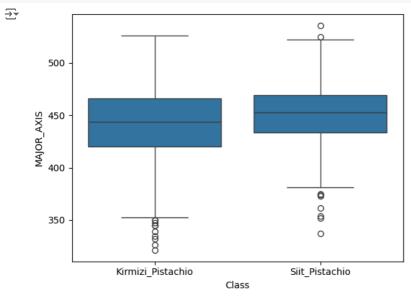
The "Class" labels were encoded into numerical values. The features were standardized using StandardScaler. The data was split into training (1202 samples) and testing (516 samples) sets, with 16 features for each sample.

```
1 #Fit the Logistic Regression model
2 logreg = LogisticRegression()
3 logreg.fit(X_train, y_train)
5 #Make predictions
6 y_pred = logreg.predict(X_test)
8 print("Accuracy:", accuracy_score(y_test, y_pred))
9 print("Confusion Matrix:\n", confusion_matrix(y_test, y_pred))
10 print("Classification Report:\n", classification_report(y_test, y_pred))
Accuracy: 0.8837209302325582
    Confusion Matrix:
      [[180 21]
      [ 19 124]]
    Classification Report:
                        precision
                                      recall f1-score
                                                         support
    Kirmizi_Pistachio
                             0.90
                                       0.90
                                                 0.90
                                                            201
       Siit_Pistachio
                             0.86
                                       0.87
                                                 0.86
                                                            143
             accuracy
                                                 0.88
                                                            344
```

```
1 sns.boxplot(x = ds['Class'], y = ds['MAJOR_AXIS'],data = ds)
2 plt.show()
```

344

344

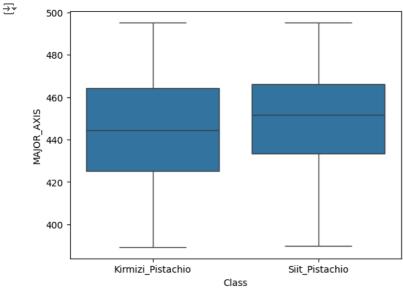


```
1 max_val = ds.MAJOR_AXIS.quantile(0.95)
2 min_val = ds.MAJOR_AXIS.quantile(0.05)
```

```
3 ds1 = ds[(ds['MAJOR_AXIS'] > min_val) & (ds['MAJOR_AXIS'] < max_val)
4 print("before dataset shape:", ds.shape)

before dataset shape: (1718, 17)
    after removing outlier's: (1546, 17)

1 sns.boxplot(x = ds1['Class'], y = ds1['MAJOR_AXIS'],data = ds1)
2 plt.show()</pre>
```



```
1 X = ds1.drop('Class', axis=1) # Features
2 y = ds1['Class'] # Target
3
4 #Split the dataset into training and testing sets
5 X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
6
7 #Feature Scaling
8 scaler = StandardScaler()
9 X_train = scaler.fit_transform(X_train)
10 X_test = scaler.transform(X_test)
11
12 X_train.shape, X_test.shape, y_train.shape, y_test.shape
```

```
→ ((1236, 16), (310, 16), (1236,), (310,))
```

```
1 #Fit the Logistic Regression model
2 logreg = LogisticRegression()
3 logreg.fit(X_train, y_train)
4
5 #Make predictions
6 y_pred = logreg.predict(X_test)
7
8 print("Accuracy:", accuracy_score(y_test, y_pred))
9 print("Confusion Matrix:\n", confusion_matrix(y_test, y_pred))
10 print("Classification Report:\n", classification_report(y_test, y_pred))
```

```
Accuracy: 0.8774193548387097
    Confusion Matrix:
     [[157 14]
     [ 24 115]]
    Classification Report:
                                     recall f1-score
                        precision
                                                         support
    Kirmizi_Pistachio
                            0.87
                                      0.92
                                                 0.89
                                                            171
       Siit_Pistachio
                                      0.83
                                                            139
                            0.89
                                                 0.86
                                                 0.88
                                                            310
             accuracy
                            0.88
                                      0.87
            macro avg
                                                 0.88
                                                            310
         weighted avg
                            0.88
                                      0.88
                                                 0.88
                                                            310
```

Insight:

The drop in model accuracy after removing the MAJOR_AXIS outliers suggests that every column (including MAJOR_AXIS) plays an important role in maintaining the overall accuracy. Even though some features might seem to have extreme values or outliers, they could still carry