

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn import metrics
from sklearn.model_selection import train_test_split
from imblearn.under_sampling import RandomUnderSampler
```

```
from collections import Counter
from sklearn import preprocessing, svm
from sklearn.linear_model import LogisticRegression
from sklearn.model_selection import RandomizedSearchCV
from sklearn.model_selection import train_test_split
from sklearn.model_selection import cross_val_score
from sklearn.model_selection import KFold
from sklearn.decomposition import PCA
from sklearn.ensemble import RandomForestClassifier
from sklearn.neighbors import KNeighborsClassifier
from xgboost import XGBClassifier
import sklearn.metrics as metrics
import pandas as pd
import matplotlib.pyplot as plt
import numpy as np
import seaborn as sns
from bioinfokit.visuz import cluster
from imblearn.over_sampling import SMOTE
from collections import Counter
from scipy.stats import randint, loguniform
```

```
df=pd.read_csv("/content/data.csv")
df.head()
```

	Area	Perimeter	MajorAxisLength	MinorAxisLength	AspectRatio	Eccentricity
0	28395	610.291	208.178117	173.888747	1.197191	0.549812
1	28734	638.018	200.524796	182.734419	1.097356	0.411785
2	29380	624.110	212.826130	175.931143	1.209713	0.562727
3	30008	645.884	210.557999	182.516516	1.153638	0.498616
4	30110	620.134	201.847882	190.279279	1.060798	0.333680

```
df.shape
```

```
(13611, 17)
```

```
df.nunique()
```

```
Area          12011
Perimeter     13351
MajorAxisLength 13543
MinorAxisLength 13543
AspectRatio    13543
Eccentricity   13543
ConvexArea    12066
EquivDiameter 12011
Extent        13535
Solidity      13522
roundness     13540
Compactness   13543
ShapeFactor1  13521
ShapeFactor2  13506
ShapeFactor3  13543
ShapeFactor4  13532
Class         7
dtype: int64
```

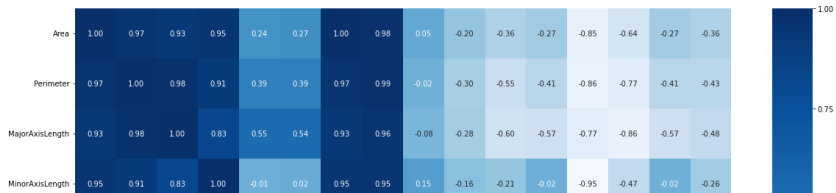
```
df["Class"].value_counts()
```

```
DERMASON    3546
SIRA        2636
SEKER       2027
HOROZ       1928
CALI        1630
BARBUNYA    1322
BOMBAY      522
Name: Class, dtype: int64
```

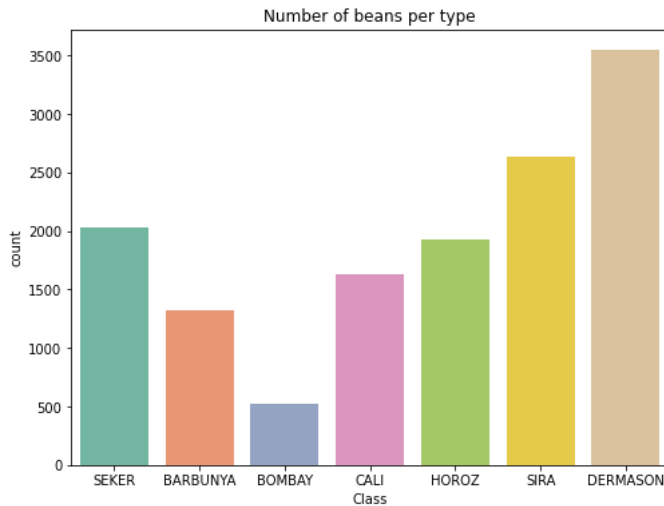
```
#check null value  
df.isnull().sum()
```

```
Area          0  
Perimeter     0  
MajorAxisLength  0  
MinorAxisLength  0  
AspectRatio    0  
Eccentricity   0  
ConvexArea     0  
EquivDiameter  0  
Extent         0  
Solidity       0  
roundness      0  
Compactness    0  
ShapeFactor1   0  
ShapeFactor2   0  
ShapeFactor3   0  
ShapeFactor4   0  
Class          0  
dtype: int64
```

```
#check correlation between features  
plt.figure(figsize=(20,20))  
sns.heatmap(df.corr(),annot=True,cmap="Blues",fmt=".2f",vmin=-1.00,vmax=1.00)  
plt.show()
```



```
#display beans per type
plt.figure(figsize=(8,6))
sns.countplot(x=df["Class"],palette="Set2")
plt.title("Number of beans per type")
plt.show()
```

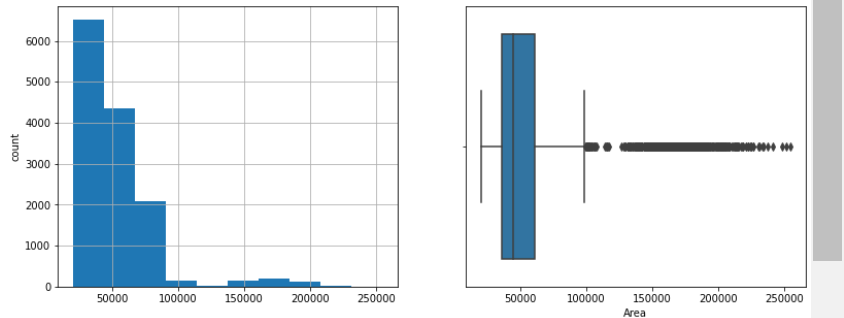


```
#display distributions and outlier
numerical_features=df.select_dtypes(include=[np.number]).columns
print("numerical features:",numerical_features)
numerical_features=numerical_features.tolist()

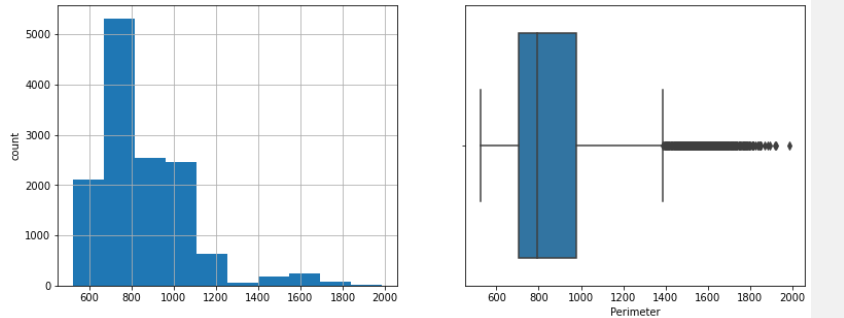
numerical features: Index(['Area', 'Perimeter', 'MajorAxisLength', 'MinorAxisLength',
    'AspectRatio', 'Eccentricity', 'ConvexArea', 'EquivDiameter', 'Extent',
    'Solidity', 'roundness', 'Compactness', 'ShapeFactor1', 'ShapeFactor2',
    'ShapeFactor3', 'ShapeFactor4'],
    dtype='object')

for i in numerical_features:
    print(i)
    print('Skewness : ', round(df[i].skew(),3))
    plt.figure(figsize = (13,5))
    plt.subplot(1,2,1)
    df[i].hist()
    plt.ylabel('count')
    plt.subplot(1,2,2)
    sns.boxplot(x = df[i])
    plt.show()
```

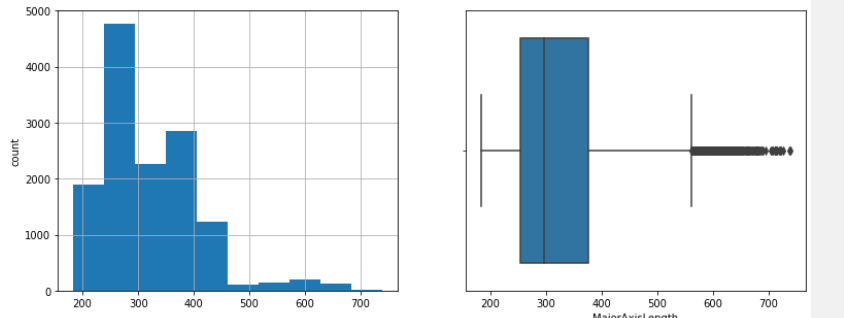
Area  
Skewness : 2.953



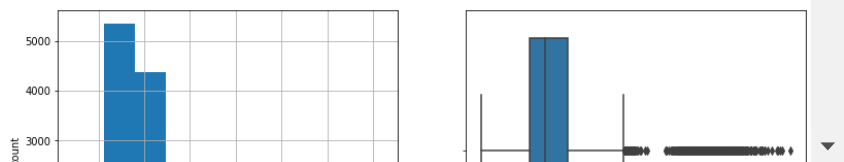
Perimeter  
Skewness : 1.626



MajorAxisLength  
Skewness : 1.358



MinorAxisLength  
Skewness : 2.238







```
X = df.drop("Class", axis=1)
Y = df['Class']

print(X.shape)
print(Y.shape)

(13611, 16)
(13611,)

# Detect outliers in the dataset

def detect_outliers(df, features):
    outlier_indices = []

    for c in features:
        Q1 = np.percentile(df[c], 25)
        Q3 = np.percentile(df[c], 75)
        IQR = Q3 - Q1
        outlier_step = IQR * 1.5
        outlier_list_col = df[(df[c] < Q1 - outlier_step) | (df[c] > Q3 + outlier_step)].index
        outlier_indices.extend(outlier_list_col)

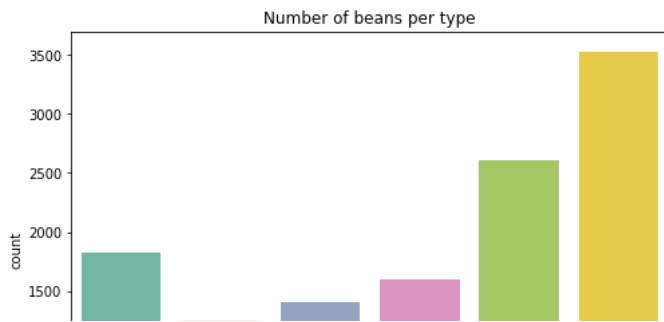
    outlier_indices = Counter(outlier_indices)
    multiple_outliers = list(i for i, v in outlier_indices.items() if v > 1)

    return multiple_outliers

data = df.drop(detect_outliers(df,['Area', 'Perimeter', 'MajorAxisLength', 'MinorAxisLength', 'AspectRatio', 'Eccentricity',
                                'ConvexArea', 'EquivDiameter', 'Extent', 'Solidity', 'roundness', 'Compactness', 'ShapeFactor1',
                                'ShapeFactor2', 'ShapeFactor3', 'ShapeFactor4']), axis=0).reset_index(drop=True)
print('Number of of samples in the dataset after removing outliers: %d' % len(data))

Number of of samples in the dataset after removing outliers: 12218

# Bar Chart to visualize the labels in the output variable
plt.figure(figsize=(8,6))
sns.countplot(x=data["Class"],palette="Set2")
plt.title("Number of beans per type")
plt.show()
```



```
# Convert Class String labels into Integers
from sklearn.preprocessing import LabelEncoder
lab_enc = LabelEncoder()
label_Y = lab_enc.fit_transform(Y)
```

SEKER BARBUNYA CALI HOROZ SIRA DERMASON

```
# Normalize the input features of the dataset
from sklearn.preprocessing import StandardScaler
normalizer = StandardScaler()
norm_X = normalizer.fit_transform(X)
```

```
##### Feature Extraction #####
#####
from sklearn.decomposition import PCA
```

```
# Visualizing the Principal Components in the feature space
pca = PCA()
pca.fit(norm_X)
loadings = pca.components_
num_pc = pca.n_features_
pc_list = ["PC" + str(i) for i in list(range(1, num_pc + 1))]
loadings_df = pd.DataFrame.from_dict(dict(zip(pc_list, loadings)))
loadings_df['variable'] = X.columns.values
loadings_df = loadings_df.set_index('variable')
```

/usr/local/lib/python3.9/dist-packages/sklearn/utils/deprecation.py:101: FutureWarning: Attribute `n\_features\_` was deprecated in v

```
# Screeplot of Principal Components
from bioinfokit.visuz import cluster
cluster.screepLOT(obj=[pc_list, pca.explained_variance_ratio_])
```

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```

```
# 2D Bi-plot of Principal Components
pca_scores = PCA().fit_transform(norm_X)
cluster.biplot(cscore=pca_scores, loadings=loadings, labels=X.columns.values, var1=round(pca.explained_variance_ratio_[0]*100, 2), var2=r
```

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```



```

# Cumulative Explained Variance Plot
plt.plot(np.cumsum(pca.explained_variance_ratio_)); plt.title('CUMULATIVE EXPLAINED VARIANCE OF THE PRINCIPAL COMPONENTS')
plt.xlabel('Number of Components'); plt.ylabel('Cumulative Explained Variance')
plt.show()

#machine learning
def training_model_metrics(model, X, Y):
    train_x, test_x, train_y, test_y = train_test_split(X, Y, test_size=0.2, random_state=12, shuffle=True)
    model.fit(train_x, train_y)
    y_pred = model.predict(test_x)
    model_acc = metrics.accuracy_score(test_y, y_pred)
    f1_measure = metrics.f1_score(test_y, y_pred, average='macro')
    model_precision = metrics.precision_score(test_y, y_pred, average='macro')
    model_recall = metrics.recall_score(test_y, y_pred, average='macro')
    print('Accuracy: %.3f, f1 measure: %.3f, precision: %.3f, recall: %.3f' % (model_acc, f1_measure, model_precision, model_recall))
    metrics.plot_confusion_matrix(model, test_x, test_y);plt.show()

def optimize_param(model, param, X_optim, Y_optim):
    rf_grid = RandomizedSearchCV(estimator=model, n_iter=30, param_distributions=param, scoring='f1_macro', n_jobs=-1,
                                cv=5, verbose=2, random_state=12)
    print('Performance Metrics for ML Model of Dataset using optimized hyper-parameters')
    print('-----')
    training_model_metrics(rf_grid, X_optim, Y_optim)
    print('The hyper-parameters with the best f1_macro performance:')
    print('-----')
    print(rf_grid.best_params_)

def evaluate_PC(model, user_input, user_output):
    train_x, test_x, train_y, test_y = train_test_split(user_input, user_output, test_size=0.2, random_state=12, shuffle=True)
    acc, comp = list(), list()

    for n in range(1, 16):
        pca = PCA(n_components=n)
        pca.fit(train_x)
        pca_transform = pca.fit_transform(train_x)
        cv = KFold(n_splits=5, shuffle=True, random_state=12)
        scores = cross_val_score(model, pca_transform, train_y, scoring='f1_macro', cv=cv, n_jobs=-1)
        acc.append(np.mean(scores))
        comp.append(n)
        print('> No of Components=%d, Accuracy=%.3f' % (n, np.mean(scores)))

    return acc, comp

def display_perf_plot(acc, comp):
    plt.plot(comp, acc)
    plt.title('PRINCIPAL COMPONENT ANALYSIS PERFORMANCE PLOT USING CROSS-VALIDATION')
    plt.axhline(y=max(acc), color='r', linestyle='--')
    plt.xlabel('NUMBER OF COMPONENTS')
    plt.ylabel('F1-MEASURE')
    plt.show()

def KFold_evaluation(model, X, Y):
    means, mins, maxs = list(), list(), list()
    folds = range(2, 13)
    for k in folds:
        cv = KFold(n_splits=k, shuffle=True, random_state=12)
        scores = cross_val_score(model, X, Y, scoring='f1_macro', cv=cv, n_jobs=-1)
        means.append(np.mean(scores))

```