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Module: Machine Learning
Module Leader: Mr. Prasan Yapa

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Machine Learning Coursework Report

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Introduction

This is the report on the coursework for machine learning. The primary goal of this assignment is to conduct a classification to label emails as spam or non-spam based on the word content. Two algorithms KNN and Decision Tree Classification were applied to accomplish this categorization. UCI Machine Learning repository was used to get the dataset.

Dataset

The spam-non spam dataset, which has over 4601 rows and 57 characteristics, was used to train the model.

Source of the DatasetUCI Machine learning repository.

Number of instances4601Number of attributes55-57Missing valuesYesNumber of classes02

Relatable Tasks Classification

Corpus Preparation

Pre – processing techniques

Data Cleaning

Data cleaning is achieved by removing null duplicates from the Spam base dataset.

No of rows in dataset before removing duplicates	No of rows in dataset after removing duplicates
4601	4210

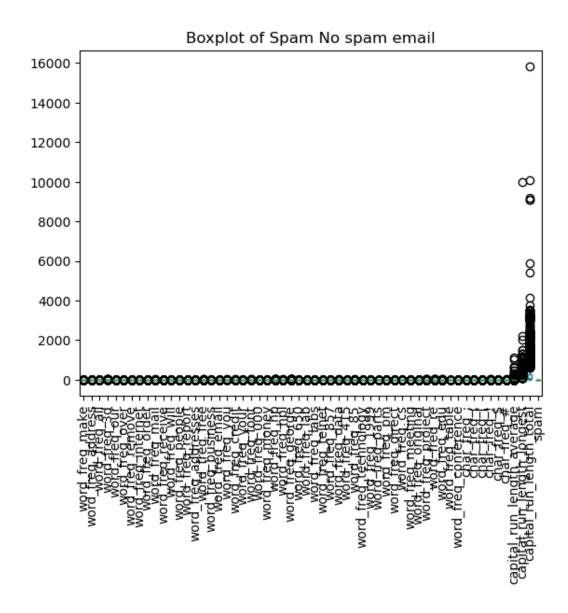
Data transformation

Data transformation is achieved by removing outliers from the dataset and by performing Standard Scaling using "sklearn.preprocessing import StandardScaler". The detected outliers were converted to null values and removed from the dataset.

No of rows in dataset before removing outliers	No of rows in dataset after removing outliers
4210	3446





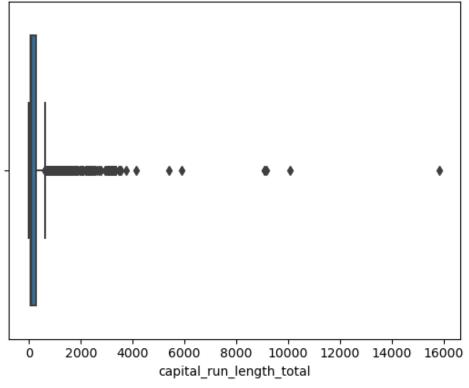


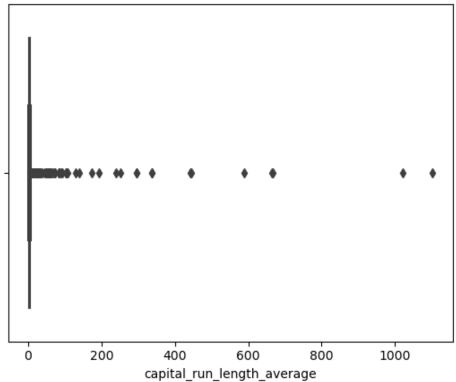
Above box plot shows that outliers are present in the columns; capital_run_length_total, capital_run_length_average and capital_run_length_longest.





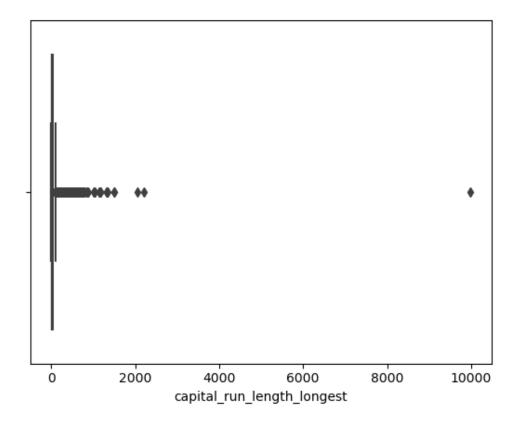
Below are the individual box plots of the above columns before removing the outliers.







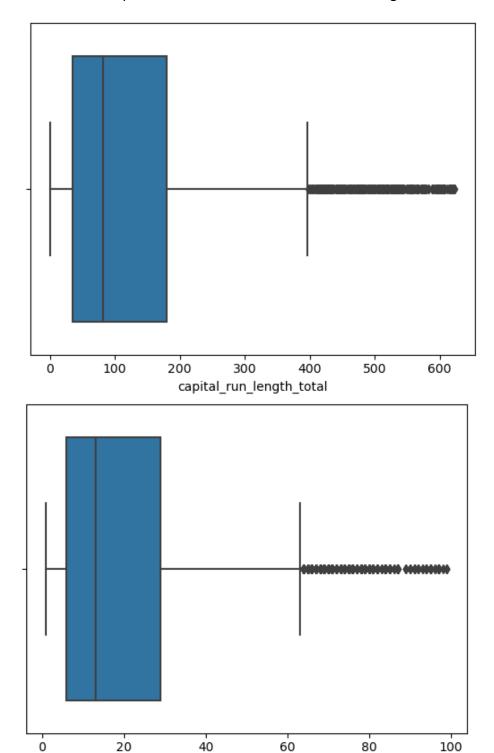








Below are the individual box plots of the above columns after removing the outliers.

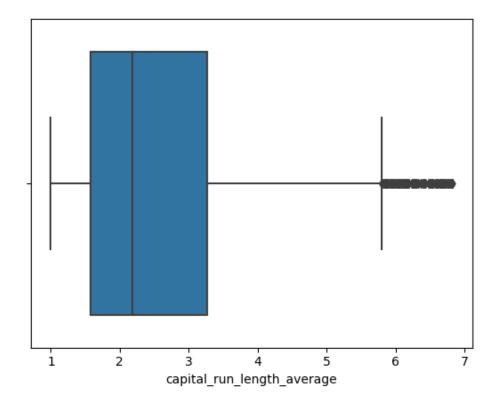


60

 $capital_run_length_longest$







Standard Scaling and relevant Visualizations

A KDE (Kernel Density Plot) plot can be used to display the distribution of a feature within the context of the Spam Base dataset.

Each feature in the Spam Base dataset has a range of values that can vary dramatically before conventional scaling. This can make it difficult to compare the distribution of different features using a KDE plot.

After performing standard scaling, however, the range of values of each feature is normalized to have zero mean and unit variance.

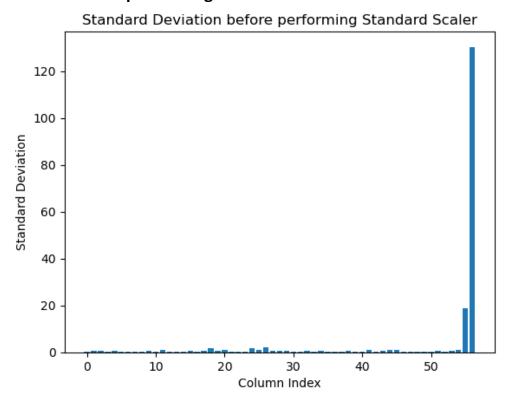
As a result, the scales of the features are now comparable, which makes it simpler to compare the distributions of the features using a KDE plot.

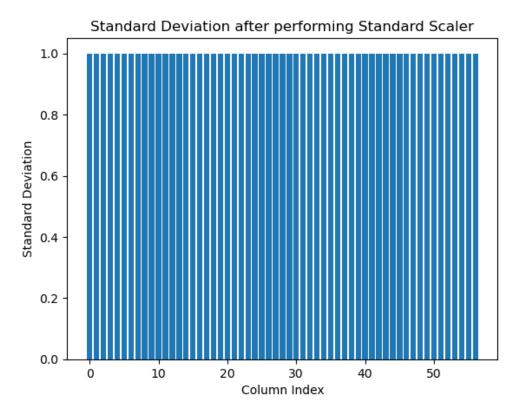
The x-axis represents the values of the feature of the Spam Base dataset, and the y-axis represents the estimated density of those values.





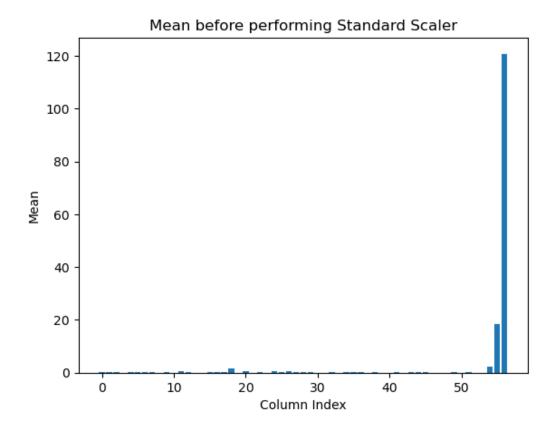
Changes achieved after performing Standard Scaler.

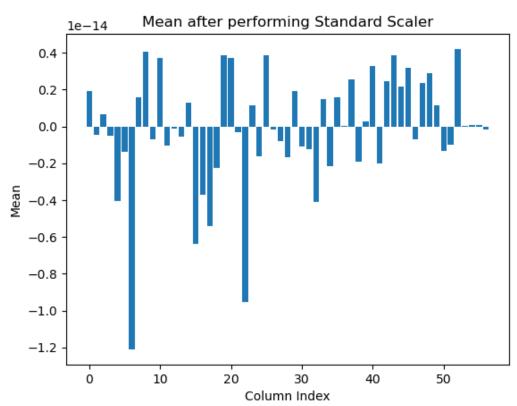








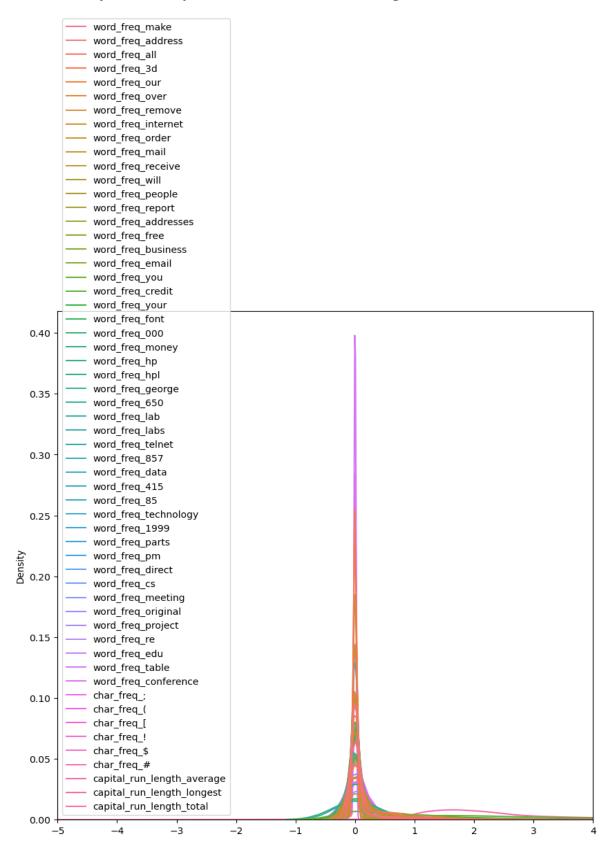








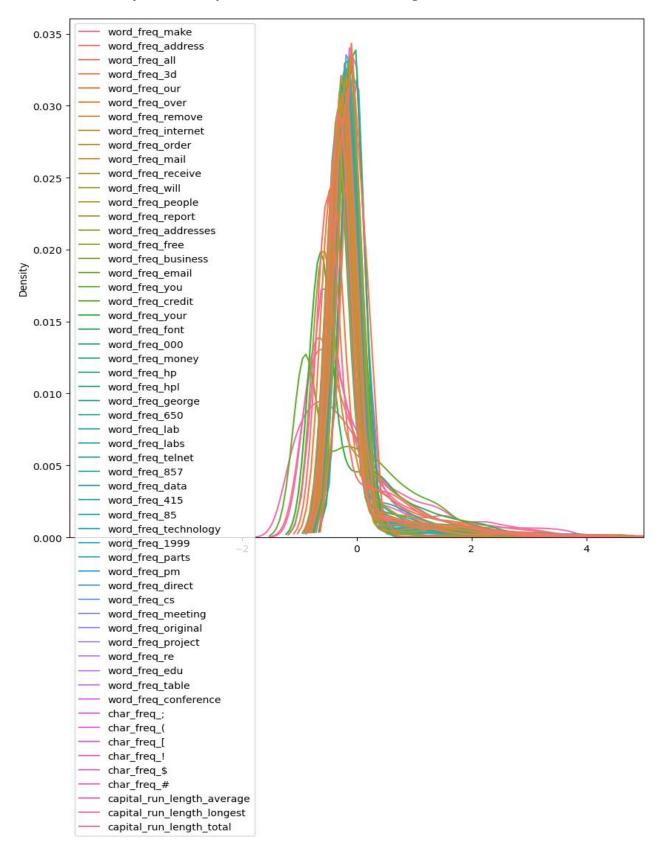
Kernel Density Estimate plot before Standard Scaling







Kernel Density Estimate plot after Standard Scaling



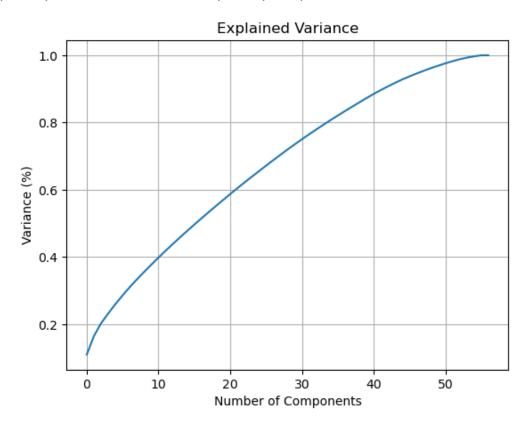




Dimensionality reduction techniques

Dimensionality reduction is a feature selection approach that allows us to use fewer features than the original dataset while maintaining a high level of information in the final model.

Principal component analysis is used to pick features using dimensionality reduction methods. As implied by the name, it extracts the primary components from the data.



From the diagram above, 44 principal components explain almost 90% of the variance in data.

So, instead of giving all the columns as inputs, 44 principal components of the data are entered to the machine learning algorithm.

Splitting the dataset into the Training set and Test set

The training set is given 80% of the dataset and testing set is given 20% of the dataset.

from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size = 0.20, random_state = 0)





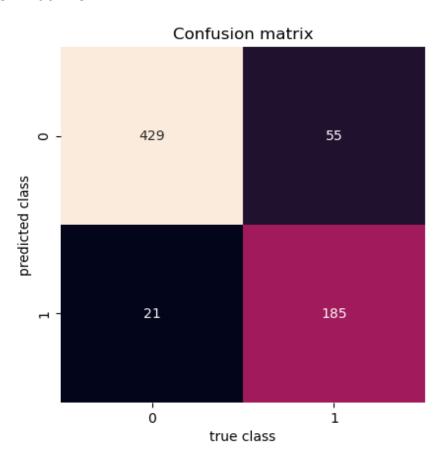
K Nearest Neighbors (KNN) Classification

Classification Report of KNN

Classification Report :

support	f1-score	recall	precision	
450	0.92	0.95	0.89	0
240	0.83	0.77	0.90	1
690 690 690	0.89 0.87 0.89	0.86 0.89	0.89	accuracy macro avg weighted avg

Confusion matrix of KNN







True Positive: 429 mails are predicted as Not Spam and it is correct.

False Positive: 55 mails are predicted as Spam, but it is Not Spam.

True Negative: 21 mails are predicted as Spam, and it is correct.

False Negative: 185 mails are predicted as Spam, but it is Not spam.

Accuracy of testing dataset

Accuracy score of email prediction using KNN: 88.98550724637681

Accuracy of training dataset

Accuracy score of email prediction using KNN: 92.8156748911466





Decision Tree Classification

Accuracy of testing dataset of Decision Tree Classification

Accuracy score of email prediction using Decision Trees = 0.8695652173913043

Summary of the test dataset

Classification Report :

	precision	recall	f1-score	support
0 1	0.88 0.84	0.92 0.78	0.90 0.81	450 240
accuracy macro avg weighted avg	0.86 0.87	0.85 0.87	0.87 0.85 0.87	690 690 690

Accuracy of training dataset of Decision Tree Classification

Accuracy score of email prediction using Decision Trees = 0.9992743105950653

Summary of the training dataset

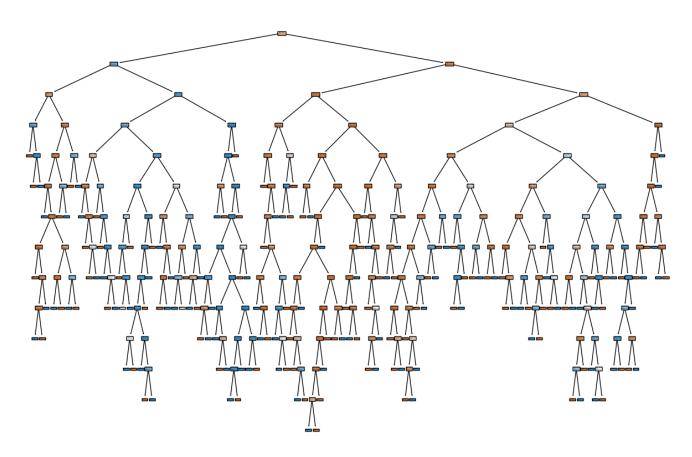
Classification Report :

precision	recall	f1-score	e support	-	
0		.00	1.00	1.00	1879 877
accuracy macro avg weighted avg		.00	1.00	1.00 1.00 1.00	2756 2756 2756





Visualizing final decision tree



Confusion Matrix before pruning the decision tree.

The Training Dataset accuracy is very high when compared to the Testing Dataset accuracy. This depicts that the model is overfitted. To avoid overfitting the Decision tree should be Pruned.

A decision tree can be pruned to minimize its size by removing branches that don't significantly improve the tree's ability to classify data. Pruning results, a decision tree that is simpler and performs better when generalizing to new data.

In the context of a Decision tree, the (TPR) true positive rate measures the proportion of positive instances correctly identified as positive.

The equation is below: TPR = TP / (TP + FN)

FN (False negative) is the number of positive instances that are incorrectly classified as positive.

In the context of a Decision tree, the (FPR) false positive rate measures the proportion of negative instances incorrectly identified as positive.

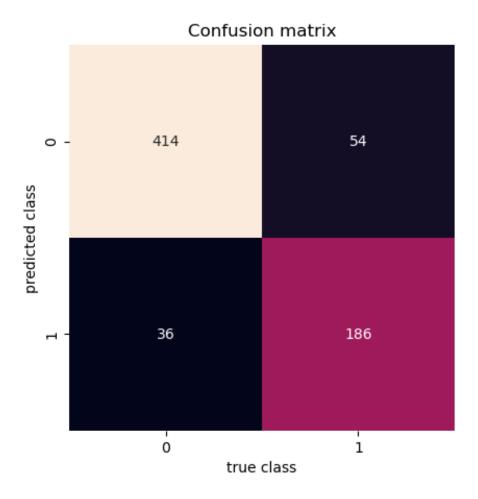




The equation is below: FPR = FP / (FP + TN)

TN (True negative) is the number of negative instances that are correctly classified as negative.

A good decision tree classifier model must have a high TPR while maintaining a lower FPR.



True Positive: 414 mails are predicted as Not Spam and it is correct.

False Positive: 54 mails are predicted as Spam, but it is Not Spam.

True Negative: 186 mails are predicted as Spam, and it is correct.

False Negative: 36 mails are predicted as Spam, but it is Not spam.



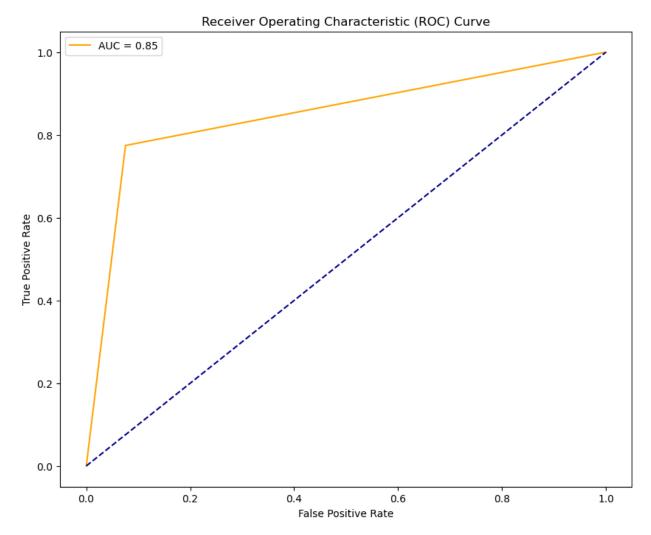


ROC Curve

An illustration of the performance of a binary classifier, such as a decision tree classifier, when the discrimination threshold is changed can be shown using a ROC (Receiver Operating Characteristic) curve.

A decision tree classifier's ROC curve will be in the top left corner of the plot if it performs well at classifying data.

The decision tree classifier's overall performance is assessed by the area under the ROC curve, with an AUC of 1.0 denoting excellent classification performance and an AUC of 0.5 denoting random guessing.



The AUC score for the above graph is 0.8475



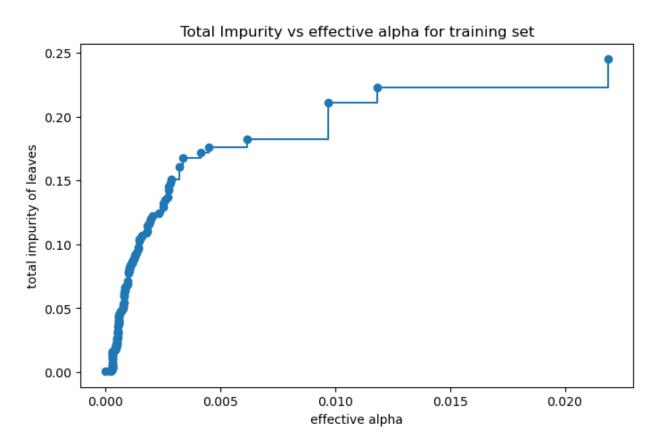


Pruning of Decision Tree

Since the training dataset was overfitted the tree needs to be pruned. The pruning process is below.

A graph is plotted between 'Total impurity of leaves' as Y axis and 'Effective alpha' as the X axis. Using the graph, we can find the optimal 'ccp alpha' value required for pruning.

The regularization parameter ccp alpha balances the accuracy and model complexity in decision trees. It helps the model perform better when it comes to generalization and prevents overfitting.



Accuracy of testing dataset after pruning

Accuracy score of email prediction after pruning the decision tree: 0.8782608695652174

Accuracy of training dataset after pruning

Accuracy score of email prediction after pruning the decision tree: 0.918722786647315

The above values, confirms that overfitting of the training dataset is treated properly.



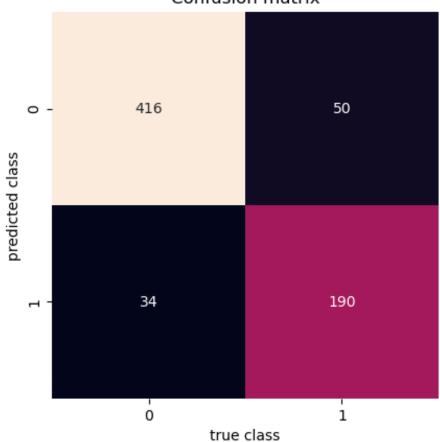


Classification Report and Confusion Matrix of Pruned Decision Tree.

Classification Report :

support	f1-score	recall	precision	
450	0.91	0.92	0.89	0
240	0.82	0.80	0.85	1
690	0.88			accuracy
690	0.87	0.86	0.87	macro avg
690	0.88	0.88	0.88	weighted avg









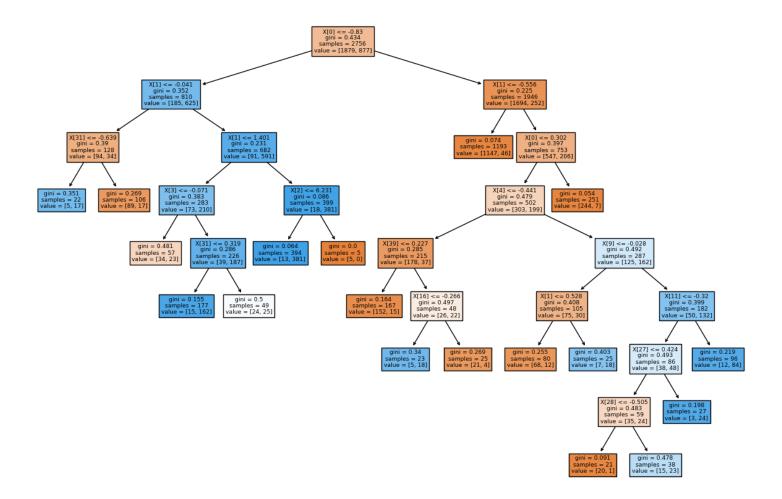
True Positive: 416 mails are predicted as Not Spam and it is correct.

False Positive: 50 mails are predicted as Spam, but it is Not Spam.

True Negative: 190 mails are predicted as Spam, and it is correct.

False Negative: 34 mails are predicted as Spam, but it is Not spam.

Decision tree obtained after pruning.







Comparisons

testing dataset after pruning
95652174
95

Accuracy of training dataset before pruning	Accuracy of training dataset after pruning
0.9992743105950653	0.918722786647315

Accuracy of testing dataset of KNN Algorithm	Accuracy of training dataset of KNN Algorithm
88.98550724637681	92.8156748911466
00.30330724037001	32.0130740311400

Limitations:

- Although PCA can make the dataset less dimensional, there is a possibility that some of the crucial features might be lost in the process. To determine the ideal number that maintains the most data, you can experiment with different principal component counts.
- Data set bias The dataset has a total of 4,601 email messages, out of which 1,813 are spam and 2,788 are non-spam. This can affect the result because more data is towards non-spam.
- The dataset is outdated because the dataset is created on 1999-07-01. The results obtained may not be helpful to come up with decisions to present days emails.
- Less amount of data When compared to the number of emails circulated in the world, 4601 email messages wouldn't be the best number to classify mails as spam or not.
- Decision trees are prone to overfitting therefore the tree should be pruned using the relevant parameters.





Future Enhancements:

- Selecting other types of models and classification algorithms, such as Random Forest, SVM, or Naive Bayes, to achieve better accuracy.
- Feature Engineering techniques can be utilized such as adding or removing specific features to see, if possible, to improve the performance of the model. Some examples are scaling techniques, such as Min-Max scaling or Normalization, chi-squared test.
- Can use Hyperparameter Tuning techniques for both KNN and Decision tree algorithms:
 - The above can be achieved by assigning maximum depth or minimum number of samples per leaf, using techniques like Grid Search or Random Search.
 - By changing the number of neighbors for KNN, using techniques like Grid Search or Random Search.
- Data Augmentation techniques can be used to generate more data to train the model. Some augmentation techniques are SMOTE or Data Synthesis. This approach can create data similar to the original data.
- Can do the classifications with recent data and data volume can be increased further to make timely decisions.





The link to git repository

https://github.com/DinethHasaranga/Machine-Learning-CW

Code:

#importing necessary libraries

import pandas as pd import numpy as np import matplotlib.pyplot as plt import seaborn as sn import re from sklearn.preprocessing import StandardScaler from sklearn.metrics import classification report, confusion matrix from sklearn.metrics import confusion_matrix import seaborn as sns from sklearn.tree import DecisionTreeClassifier from sklearn.metrics import roc_curve,auc from sklearn.metrics import roc_auc_score from sklearn.model_selection import train_test_split from sklearn.metrics import accuracy_score from sklearn.decomposition import PCA from sklearn import tree from sklearn.neighbors import KNeighborsClassifier

print("No of rows in dataset before preprocessing : ", len(Data_set))

#Loading the dataset

```
with open("C:\\Users\\Admin\\Desktop\\ML CW\\spambase.names") as spam:
text = spam.read()
labels = re.findall(r'\n(\w*_?\W?):', text)

Data_set = pd.read_csv("C:\\Users\\Admin\\Desktop\\ML CW\\spambase.data", header=None,
names=labels +['spam'])

Data_array=Data_set.values
# print(Data_array)

# printing first 5 rows
Data_set.head()
```





Finding the duplicates in the dataset

#Checking the availability of duplicates Data_set.duplicated()

Dropping the duplicate values

Data_set.drop_duplicates(inplace=True)
print("No of rows in dataset after removing duplicates : ", len(Data_set))

Finding the outliers in the dataset

fig = plt.figure(figsize =(100, 50))
Data_set.plot.box(title='Boxplot of Spam No spam email',rot=90)
plt.show()

Boxplot of capital_run_length_total

sn.boxplot(x = Data_set['capital_run_length_total'])

Boxplot of capital_run_length_average

sn.boxplot(x = Data_set['capital_run_length_average'])

Boxplot of capital_run_length_longest

sn.boxplot(x = Data_set['capital_run_length_longest'])

Making all the outliers as Null values from IQR technique

```
for x in ['capital_run_length_total','capital_run_length_longest','capital_run_length_average']:
    q75,q25 = np.percentile(Data_set.loc[:,x],[75,25])
    intr_qr = q75-q25

max = q75+(1.5*intr_qr)
min = q25-(1.5*intr_qr)

Data_set.loc[Data_set[x] < min,x] = np.nan
Data_set.loc[Data_set[x] > max,x] = np.nan
```

Boxplot of capital_run_length_total without outliers

sn.boxplot(x = Data set['capital run length total'])

Boxplot of capital_run_length_longest without outliers

sn.boxplot(x = Data_set['capital_run_length_longest'])

Boxplot of capital_run_length_average without outliers

sn.boxplot(x = Data_set['capital_run_length_average'])





Finding the null values in the dataset

Data_set.isna().sum().any()
Data_set.isna().sum()

Removing all the Null values

Drop all rows with NaN values df2=Data_set.dropna() df2=Data_set.dropna(axis=0)

Removing the target column

Reset index after drop df2=Data_set.dropna().reset_index(drop=True)

Summary of dataset before performing Standard Scaler

data.describe()

Kernel Density Plot

fig, ax = plt.subplots(figsize=(10,10)) sns.kdeplot(data=data, ax=ax) ax.set_xlim(-5, 4) plt.show()

Calculate the standard deviation of all columns std dev = data.std()

Plot the standard deviation of all columns
plt.bar(range(len(std_dev)), std_dev)
plt.title("Standard Deviation before performing Standard Scaler")
plt.xlabel("Column Index")
plt.ylabel("Standard Deviation")
plt.show()

Calculate the mean of all columns mean = data.mean()

Plot the mean of all columns
plt.bar(range(len(mean)), mean)
plt.title("Mean before performing Standard Scaler")
plt.xlabel("Column Index")
plt.ylabel("Mean")
plt.show()





Performing Standard Scaling for the dataset

```
scaler=StandardScaler()
scaled_data=scaler.fit_transform(data)
df=pd.DataFrame(data=scaled_data, columns= data.columns)
df
```

Summmary of dataset after performing Standard Scaling

```
print("No of rows in dataset after preprocessing : ", len(Data_set))
data.describe()
```

Performing PCA to the Dataset

```
pca = PCA()
principalComponents = pca.fit_transform(df)

plt.figure()
plt.plot(np.cumsum(pca.explained_variance_ratio_))

plt.xlabel('Number of Components')
plt.ylabel('Variance (%)') #for each component

plt.title('Explained Variance')
plt.grid(True)
plt.show()
```

KDE plot after performing Standard Scaler

```
fig, ax = plt.subplots(figsize=(10,10))
sns.kdeplot(data=df, ax=ax)
ax.set_xlim(-5, 5)
plt.show()
```

Calculate the standard deviation of all columns std_dev = df.std()

Plot the standard deviation of all columns
plt.bar(range(len(std_dev)), std_dev)
plt.title("Standard Deviation after performing Standard Scaler")
plt.xlabel("Column Index")
plt.ylabel("Standard Deviation")
plt.show()

Calculate the mean of all columns mean = df.mean()





```
# Plot the mean of all columns
plt.bar(range(len(mean)), mean)
plt.title("Mean after performing Standard Scaler")
plt.xlabel("Column Index")
plt.ylabel("Mean")
plt.show()
pca = PCA()
principalComponents = pca.fit transform(df)
plt.figure()
plt.plot(np.cumsum(pca.explained_variance_ratio_))
plt.xlabel('Number of Components')
plt.ylabel('Variance (%)') #for each component
plt.title('Explained Variance')
plt.grid(True)
plt.show()
Introducing the PCA components
pca = PCA(n_components=44)
new_data = pca.fit_transform(df)
# This will be the new data fed to the algorithm.
principal_Df = pd.DataFrame(data = new_data
       , columns = ['PC1',
'PC2','PC3','PC4','PC5','PC6','PC7','PC8','PC9','PC10','PC11','PC12','PC13','PC14','PC15','PC16','PC17','PC18'
,'PC19','PC20', 'PC21',
PC22','PC23','PC24','PC25','PC26','PC27','PC28','PC29','PC30','PC31','PC32','PC33','PC34','PC35','PC36','PC
37','PC38','PC39','PC40','PC41','PC42','PC43','PC44'])
Dataset after performing PCA
principal_Df.head()
# principal_Df
print(pca.explained_variance_)
print(pca.components )
```

Build the predictive model by appling Decision Tree algorithm

X = principal_Df.iloc[:,0:44].values
y = Data_set.iloc[:, 57].values

Splitting the dataset into the Training set and Test set

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size = 0.20, random_state = 0)





Applying KNN Algorithm

```
X = principal_Df.iloc[:,0:44].values
y = Data_set.iloc[:, 57].values
```

Splitting the dataset into the Training set and Test set

```
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size = 0.20, random_state = 0)
```

```
# Fitting classifier to the Training set classifier = KNeighborsClassifier(n_neighbors=5, metric='minkowski', p=2) classifier.fit(X_train,y_train)
```

Predicting the Test set results
y_pred = classifier.predict(X_test)

Classification Report

```
print('Classification Report : \n\n')
print(classification_report(y_test, y_pred))
```

Accuracy of testing dataset

print("Accuracy score of email prediction using KNN: ",accuracy_score(y_pred,y_test)*100)

Accuracy of training dataset

```
y_pred2 = classifier.predict(X_train)
print("Accuracy score of email prediction using KNN : ",accuracy_score(y_pred2,y_train)*100)
```

Visualization

Summary of the predictions made by the classifier mat = confusion_matrix(y_test, y_pred) sns.heatmap(mat.T, square=True, annot=True, fmt='d', cbar=False)

plt.title('Confusion matrix') plt.xlabel('true class') plt.ylabel('predicted class')





Applying Decision Tree Algorithm

Fitting classifier to the Training set

clf = DecisionTreeClassifier(random_state=0,criterion='gini')
clf.fit(X_train,y_train)

Accuracy of testing dataset

predictions_test=clf.predict(X_test)
accuracy_score(y_test, predictions_test)

Accuracy of training dataset

predictions_train = clf.predict(X_train)
accuracy_score(y_train,predictions_train)

Visualizing final decision tree

from sklearn import tree plt.figure(figsize=(15,10)) tree.plot_tree(clf,filled=True) plt.show()

Summary of the test dataset

print('Classification Report : \n\n')
print(classification_report(y_test,predictions_test))

Summary of the predictions made by the classifier mat = confusion_matrix(y_test, predictions_test) sns.heatmap(mat.T, square=True, annot=True, fmt='d', cbar=False)

plt.title('Confusion matrix') plt.xlabel('true class') plt.ylabel('predicted class')

Summary of the training dataset

print(classification_report(y_train,predictions_train))
print(confusion_matrix(y_train,predictions_train))

Evaluating the false positive rate and true positive rate

dt_probs = clf.predict_proba(X_test)[:,1]
fpr_dt, tpr_dt, thresholds_dt = roc_curve(y_test,dt_probs)

Plotting ROC curve for our Decision Tree

auc_score_dt = auc(fpr_dt,tpr_dt)
auc_score_dt

def plot_roc_curve(fpr, tpr):





```
plt.figure(figsize=(10,8))
  plt.plot(fpr dt, tpr dt, color='orange', label='AUC = %0.2f' % auc score dt)
  plt.plot([0, 1], [0, 1], color='darkblue', linestyle='--')
  plt.xlabel('False Positive Rate')
  plt.ylabel('True Positive Rate')
  plt.title('Receiver Operating Characteristic (ROC) Curve')
  plt.legend()
  plt.show()
plot_roc_curve(fpr_dt,tpr_dt)
Pruning the decision tree
path = clf.cost_complexity_pruning_path(X_train, y_train)
ccp_alphas, impurities = path.ccp_alphas, path.impurities
fig, ax = plt.subplots(figsize=(8,5))
ax.plot(ccp_alphas[:-1], impurities[:-1], marker='o', drawstyle="steps-post")
ax.set xlabel("effective alpha")
ax.set_ylabel("total impurity of leaves")
ax.set_title("Total Impurity vs effective alpha for training set")
clf = DecisionTreeClassifier(random_state=0, ccp_alpha=0.003)
clf.fit(X_train,y_train)
Accuracy of testing dataset after pruning
pred=clf.predict(X_test)
accuracy_score(y_test, pred)
Accuracy of training dataset after pruning
pred_1 = clf.predict(X_train)
accuracy_score(y_train,pred_1)
print('Classification Report : \n\n')
print(classification_report(y_test,pred))
# Summary of the predictions made by the classifier
mat = confusion matrix(y test, pred)
sns.heatmap(mat.T, square=True, annot=True, fmt='d', cbar=False)
plt.title('Confusion matrix')
plt.xlabel('true class')
plt.ylabel('predicted class')
```





Decision tree obtained after pruning

plt.figure(figsize=(15,10))
tree.plot_tree(clf,filled=True)
plt.show()