

Project File Machine Learning and Artificial Intelligence

Topic-Classification of mushrooms as edible or poisonous based their different features

EAEPCO9 ECAM – 1

Submitted To -

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<u>Acknowledgement</u>

In successfully completing this project, many people have I would like to express my special thanks of gratitude to my teacher **Ms Kavya Gupta** and our University **Netaji Subhas University of technology** (NSUT) who gave me the golden opportunity to do this project on the topic MUSHROOM CLASSIFICATION. It helped me in doing a lot of Research and i came to know about a lot of things related to this topic. Finally, I would also like to thank my parents and friends who helped me a lot in finalising this project within the limited time frame.

Abstract

Predicting if a set of mushrooms is edible or not corresponds to the task of classifying them into two groups—edible or poisonous—on the basis of a classification rule. To support this binary task, I have collected the largest and most comprehensive attribute based data available. In the work, we detail the creation, curation and simulation of a data set for binary classification. We evaluated different machine learning algorithms, namely, naive Bayes, logistic regression,KNN, Random forest, decision tree and stochastic gradient method. The data set contains 23 families and is the largest available. I further provide a fully reproducible workflow and provide the data under the FAIR principles.

Methodology

Libraries Used:

- Sklearn
- Pandas
- Matplotlib
- Seaborn
- Numpy

Classification Models Used:

- Naïve-Bayes
- Logistic regression
- KNN
- Random forest
- Decision tree
- Stochastic gradient method

<u>PROJECT ON THE</u> <u>CLASSIFICATION OF MUSHROOMS</u> INTO EDIBLE OR POISONOUS

IMPORTING NECESSARY LIBRARIES

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sn
from sklearn import preprocessing, tree
from sklearn.metrics import
roc_curve,accuracy_score,confusion_matrix,classification_report,ro
c_auc_score
from sklearn.model selection import train test split
from sklearn.linear_model import LogisticRegression
from sklearn.naive_bayes import GaussianNB
from sklearn.neighbors import KNeighborsClassifier
from sklearn.ensemble import RandomForestClassifier
from sklearn.tree import DecisionTreeClassifier
from sklearn import feature selection
%matplotlib inline
df=pd.read csv("mushroom data.csv")
df.shape
 (61069, 21)
```

df.head()

class	cap- diameter	cap- shape	cap- surface	cap- color	does- bruise- or- bleed	gill- attachment	gill- spacing	gill- color	stem- height	 stem- root	stem- surface	stem- color	veil- type	veil- color	has- ring	ring- type	spore- print- color
р	15.26		g			е	NaN	w	16.95		у	w		w		g	NaN
р	16.60		g				NaN	w	17.99		у	w		w		g	NaN
р	14.07		g			е	NaN	w	17.80		у	w		w		g	NaN
р	14.17		h	е			NaN	w	15.77		у	w		w		р	NaN
р	14.64		h			е	NaN	w	16.53		У	w		w		р	NaN

Converting String/Char Values to Int

```
le_cap_shape=preprocessing.LabelEncoder()
le cap shape.fit(df['cap-shape'])
df['cap-shape']=le cap shape.transform(df['cap-shape'])
le cap surface=preprocessing.LabelEncoder()
le cap surface.fit(df['cap-surface'])
df['cap-surface']=le_cap_surface.transform(df['cap-surface'])
le cap color=preprocessing.LabelEncoder()
le cap color.fit(df['cap-color'])
df['cap-color']=le cap color.transform(df['cap-color'])
le gill color=preprocessing.LabelEncoder()
le gill color.fit(df['gill-color'])
df['gill-color']=le gill color.transform(df['gill-color'])
le stem color=preprocessing.LabelEncoder()
le stem color.fit(df['stem-color'])
df['stem-color']=le stem color.transform(df['stem-color'])
le has ring=preprocessing.LabelEncoder()
le has ring.fit(df['has-ring'])
df['has-ring']=le has ring.transform(df['has-ring'])
le habitat=preprocessing.LabelEncoder()
le habitat.fit(df['habitat'])
df['habitat']=le habitat.transform(df['habitat'])
le class=preprocessing.LabelEncoder()
le class.fit(df['class'])
df['class']=le class.transform(df['class'])
le does bruise=preprocessing.LabelEncoder()
le does bruise.fit(df['does-bruise-or-bleed'])
df['does-bruise-or-bleed']=le does bruise.transform(df['does-
bruise-or-bleed'])
le gill attachment=preprocessing.LabelEncoder()
le gill attachment.fit(df['gill-attachment'])
df['gill-attachment']=le gill attachment.transform(df['gill-
attachment'l)
le_stem_root=preprocessing.LabelEncoder()
le_stem_root.fit(df['stem-root'])
df['stem-root']=le stem_root.transform(df['stem-root'])
```

```
le stem surface=preprocessing.LabelEncoder()
le_stem_surface.fit(df['stem-surface'])
df['stem-surface']=le stem surface.transform(df['stem-surface'])
le_veil_type=preprocessing.LabelEncoder()
le_veil_type.fit(df['veil-type'])
df['veil-type']=le veil type.transform(df['veil-type'])
le veil color=preprocessing.LabelEncoder()
le_veil_color.fit(df['veil-color'])
df['veil-color']=le veil color.transform(df['veil-color'])
le ring type=preprocessing.LabelEncoder()
le ring type.fit(df['ring-type'])
df['ring-type']=le ring type.transform(df['ring-type'])
le season=preprocessing.LabelEncoder()
le season.fit(df['season'])
df['season']=le season.transform(df['season'])
df=df.drop(['gill-spacing','spore-print-color'],axis=1)
df.head()
```

class	cap- diameter	cap- shape	cap- surface	cap- color	does- bruise- or- bleed	gill- attachment				stem- root	stem- surface	stem- color		veil- color		habitat	season
1	15.26	6	2	6	0	2	10	16.95	17.09	4	7	11	0	4	2	0	3
1	16.60	6	2	6	0	2	10	17.99	18.19	4	7	11		4	2		2
1	14.07	6	2	6	0	2	10	17.80	17.74	4	7	11	0	4	2		3
1	14.17	2	3		0	2	10	15.77	15.98	4		11		4	5		3
1	14.64	6	3	6	0	2	10	16.53	17.20	4	7	11	0	4	5	0	3

Visualizing the correlation between all the features

```
f, ax = plt.subplots(figsize=(20, 16))
corr = df.corr()
sn.heatmap(corr, annot=True)
plt.show()
```

					0.005	0.00	0.0000	0.053	0.10				0.005	0.050		2.252		2.242	
dass -	- 1	-0.18	-0.11	-0.033	0.026	-0.02	-0.0038	-0.061	-0.12	-0.2	-0.011	-0.17	-0.095	-0.053	-0.076	0.058	0.033	-0.042	-0.054
cap-diameter -	-0.18	1	0.1	0.035	0.042	0.18	0.011	0.22	0.42	0.7	-0.074	0.043	0.0087	-0.082	-0.041	0.077	0.061	-0.13	0.037
cap-shape -	-0.11	0.1	1	-0.046	0.016	0.19	-0.15	0.12	0.034	0.16	-0.026	0.12	0.0091	0.034	-0.1	0.036	0.14	-0.12	-0.027
cap-surface -	-0.033	0.035	-0.046	1	0.096	-0.09	0.076	-0.027	0.046	0.063	-0.11	-0.033	0.12	-0.0017	0.027	0.038	-0.071	0.039	-0.0012
cap-color -	0.026	0.042	0.016	0.096	1	0.011	0.01	0.14	-0.011	0.00057	0.11	0.054	0.27	-0.051	0.0044	0.071	0.018	0.032	0.0084
does-bruise-or-bleed -	-0.02	0.18	0.19	-0.09	0.011	1	-0.078	0.019	0.077	0.18	-0.0037	0.24	-0.025	0.039	0.028	0.019	-0.011	-0.1	-0.051
gill-attachment -	-0.0038	0.011	-0.15	0.076	0.01	-0.078	1	-0.033	0.1	0.06	-0.19	0.065	0.098	-0.038	0.0042	-0.012	-0.037	-0.13	-0.033
gill-color -	-0.061	0.22	0.12	-0.027	0.14	0.019	-0.033	1	0.16	0.11	-0.11	0.038	0.15	-0.18	-0.011	-0.056	0.043	-0.076	-0.011
stem-height -	-0.12	0.42	0.034	0.046	-0.011	0.077	0.1	0.16	1	0.44	-0.073	0.14	0.14	-0.3	-0.24	0.28	0.23	-0.021	-0.021
stem-width -	-0.2	0.7	0.16	0.063	0.00057	0.18	0.06	0.11	0.44	1	-0.084	0.16	0.058	-0.04	-0.021	-0.00098	-0.018	-0.11	0.014
stem-root -	-0.011	-0.074	-0.026	-0.11	0.11	-0.0037	-0.19	-0.11	-0.073	-0.084	1	0.1	0.041	0.14	0.13	-0.1	-0.0072	0.027	-0.027
stem-surface -	-0.17	0.043	0.12	-0.033	0.054	0.24	0.065	0.038	0.14	0.16	0.1	1	0.25	-0.073	-0.072	-0.0081	0.024	-0.017	-0.042
stem-color -	-0.095	0.0087	0.0091	0.12	0.27	-0.025	0.098	0.15	0.14	0.058	0.041	0.25	1	-0.18	-0.032	0.15	0.091	0.04	-0.025
veil-type -	-0.053	-0.082	0.034	-0.0017	-0.051	0.039	-0.038	-0.18	-0.3	-0.04	0.14	-0.073	-0.18	1	0.46	-0.35	-0.15	0.12	-0.02
veil-color -	-0.076	-0.041	-0.1	0.027	0.0044	0.028	0.0042	-0.011	-0.24	-0.021	0.13	-0.072	-0.032	0.46	1	-0.36	-0.31	0.083	0.054
has-ring -	0.058	0.077	0.036	0.038	0.071	0.019	-0.012	-0.056	0.28	-0.00098	-0.1	-0.0081	0.15	-0.35	-0.36	1	0.65	0.1	0.039
ring-type -	0.033	0.061	0.14	-0.071	0.018	-0.011	-0.037	0.043	0.23	-0.018	-0.0072	0.024	0.091	-0.15	-0.31	0.65	1	-0.039	0.03
habitat -	-0.042	-0.13	-0.12	0.039	0.032	-0.1	-0.13	-0.076	-0.021	-0.11	0.027	-0.017	0.04	0.12	0.083	0.1	-0.039	1	0.0044
season -	-0.054	0.037	-0.027	-0.0012	0.0084	-0.051	-0.033	-0.011	-0.021	0.014	-0.027	-0.042	-0.025	-0.02	0.054	0.039	0.03	0.0044	1
	dass –	cap-diameter –	cap-shape -	cap-surface -	cap-color -	does-bruise-or-bleed –	gill-attachment –	gill-color –	stem-height –	stem-width -	stem-root -	stem-surface -	stem-color -	veil-type –	veil-color –	has-ring –	ring-type –	habitat –	season –

Splitting Dataset

```
xdf=df.copy()
xdf.drop(['class'],axis=1,inplace=True)

ydf=df['class'].copy()
ydf

X_train, X_test,Y_train,Y_test =
train_test_split(xdf,ydf,test_size=0.2,random_state=0)
```

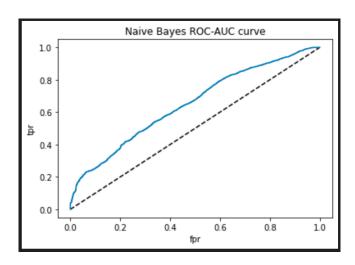
Naive Bayes Model

```
NB_model = GaussianNB()
NB_model.fit(X_train,Y_train)
Y_hat_NB=NB_model.predict(X_test)
print("Accuracy : {}
%".format(accuracy_score(Y_test,Y_hat_NB)*100))
print("Train Accuracy : {}
%".format(accuracy_score(Y_train,NB_model.predict(X_train))*100))
print(classification_report(Y_test,Y_hat_NB))
```

Accuracy : 59.56279679056819% Train Accuracy : 60.04707808822024%						
,	recision	recall	f1-score	support		
0	0.53	0.59	0.56	5302		
1	0.66	0.60	0.63	6912		
accuracy			0.60	12214		
macro avg	0.59	0.60	0.59	12214		
weighted avg	0.60	0.60	0.60	12214		

ROC-AUC curve

```
y_pred_proba = NB_model.predict_proba(X_test)[:,1]
fpr, tpr, thresholds = roc_curve(Y_test, y_pred_proba)
plt.plot([0,1],[0,1],'k--')
plt.plot(fpr,tpr)
plt.xlabel('fpr')
plt.ylabel('tpr')
plt.title('Naive Bayes ROC-AUC curve')
plt.show()
```



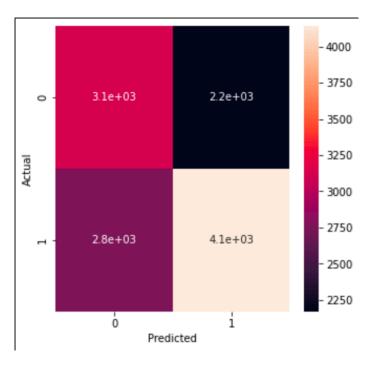
ROC-AUC Score

roc_auc_score(Y_test,y_pred_proba)

0.6492078952124984

Confusion matrix

```
cm=confusion_matrix(Y_test,Y_hat_NB)
plt.figure(figsize=(5,5))
sn.heatmap(cm,annot=True)
plt.xlabel('Predicted')
plt.ylabel('Actual')
```



Logistic Regression Model

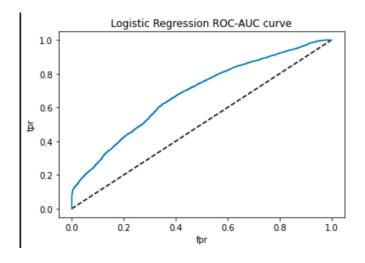
```
LR_model = LogisticRegression(C=0.01,solver='sag')
LR_model.fit(X_train,Y_train)
Y_hat_LR=LR_model.predict(X_test)
print("Accuracy : {}
%".format(accuracy_score(Y_test,Y_hat_LR)*100))
print("Train Accuracy : {}
%".format(accuracy_score(Y_train,LR_model.predict(X_train))*100))
```

print(classification_report(Y_test,Y_hat_LR))

Accuracy : 64	Accuracy : 64.09857540527264%							
Train Accurac	cy : 64.1367	3114317879	8					
	precision	recall	f1-score	support				
0	0.61	0.49	0.54	5302				
1	0.66	0.76	0.70	6912				
accuracy			0.64	12214				
macro avg	0.63	0.62	0.62	12214				
weighted avg	0.64	0.64	0.63	12214				

ROC-AUC curve

```
y_pred_proba = LR_model.predict_proba(X_test)[:,1]
fpr, tpr, thresholds = roc_curve(Y_test, y_pred_proba)
plt.plot([0,1],[0,1],'k--')
plt.plot(fpr,tpr)
plt.xlabel('fpr')
plt.ylabel('tpr')
plt.title('Logistic Regression ROC-AUC curve')
plt.show()
```



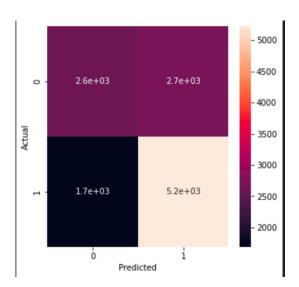
ROC-AUC Score

roc_auc_score(Y_test,y_pred_proba)

0.6815160596280929

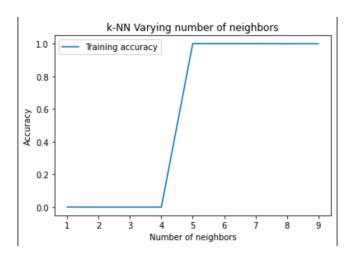
Confusion matrix

```
cm=confusion_matrix(Y_test,Y_hat_LR)
plt.figure(figsize=(5,5))
sn.heatmap(cm,annot=True)
plt.xlabel('Predicted')
plt.ylabel('Actual')
plt.show()
```



KNN Model

```
Ks=100
train_accuracy =np.zeros((Ks-1))
test_accuracy = np.zeros((Ks-1))
mean_acc=np.zeros((Ks-1))
for n in range(5,Ks):
    knn=KNeighborsClassifier(n neighbors=n)
    knn.fit(X_train,Y_train)
    Y_hat_knn=knn.predict(X_test)
    mean_acc[n-1]=accuracy_score(Y_test,Y_hat_knn)
    train_accuracy[n-1] = knn.score(X_train, Y_train)
test_accuracy[n-1] = knn.score(X_test, Y_test)
print("Max accuracy is: ",mean_acc.max(),"with
k=",mean acc.argmax()+1)
Max accuracy is: 0.9988537743572949 with k= 5
empty= np.arange(1,10)
plt.title('k-NN Varying number of neighbors')
plt.plot(empty, train_accuracy, label='Training accuracy')
plt.legend()
plt.xlabel('Number of neighbors')
plt.ylabel('Accuracy')
plt.show()
```

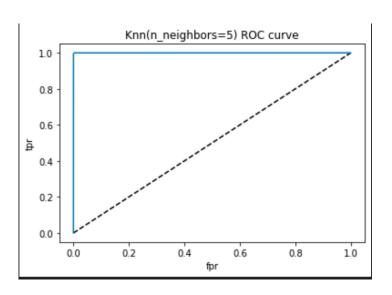


```
KNN_model=KNeighborsClassifier(n_neighbors=mean_acc.argmax()+1)
KNN_model.fit(X_train,Y_train)
Y_hat_KNN=KNN_model.predict(X_test)
print("Accuracy : {}
%".format(accuracy_score(Y_test,Y_hat_KNN)*100))
print("Train Accuracy : {}
%".format(accuracy_score(Y_train,KNN_model.predict(X_train))*100))
print(classification report(Y test,Y hat KNN))
```

Accuracy : 99.	Accuracy : 99.8853774357295%							
Train Accuracy	: 99.94678	129157711	%					
	precision	recall	f1-score	support				
0	1.00	1.00	1.00	5302				
1	1.00	1.00	1.00	6912				
accuracy			1.00	12214				
macro avg	1.00	1.00	1.00	12214				
weighted avg	1.00	1.00	1.00	12214				

ROC-AUC curve

```
y_pred_proba = knn.predict_proba(X_test)[:,1]
fpr, tpr, thresholds = roc_curve(Y_test, y_pred_proba)
plt.plot([0,1],[0,1],'k--')
plt.plot(fpr,tpr, label='Knn')
plt.xlabel('fpr')
plt.ylabel('tpr')
plt.title('Knn(n_neighbors=5) ROC-AUC curve')
plt.show()
```



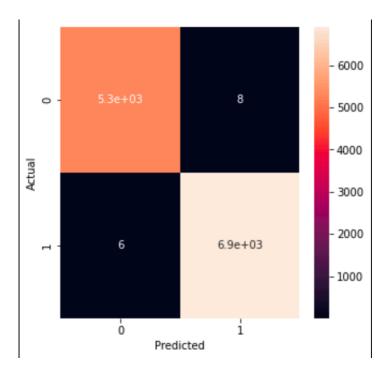
ROC-AUC Score

roc_auc_score(Y_test,y_pred_proba)

0.9998268909705631

Confusion matrix

cm=confusion_matrix(Y_test,Y_hat_KNN)
plt.figure(figsize=(5,5))
sn.heatmap(cm,annot=True)
plt.xlabel('Predicted')
plt.ylabel('Actual')



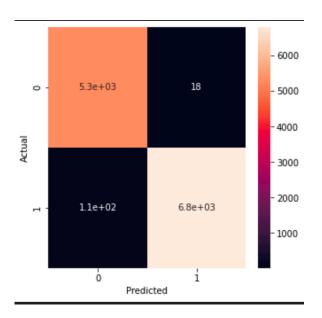
Random Forest Model

```
RF_model =
RandomForestClassifier(n_estimators=100,max_depth=10,random_state=
0)
RF_model.fit(X_train,Y_train)
Y_hat_RF=RF_model.predict(X_test)
print("Accuracy : {}
%".format(accuracy_score(Y_test,Y_hat_RF)*100))
print("Train Accuracy : {}
%".format(accuracy_score(Y_train,RF_model.predict(X_train))*100))
print(classification_report(Y_test,Y_hat_RF))
```

Accuracy : 98.	Accuracy : 98.97658424758474%						
Train Accuracy	: 99.09528	195681098	%				
1	precision	recall	f1-score	support			
0	0.98	1.00	0.99	5302			
1	1.00	0.98	0.99	6912			
accuracy			0.99	12214			
macro avg	0.99	0.99	0.99	12214			
weighted avg	0.99	0.99	0.99	12214			

Confusion matrix

```
cm=confusion_matrix(Y_test,Y_hat_RF)
plt.figure(figsize=(5,5))
sn.heatmap(cm,annot=True)
plt.xlabel('Predicted')
plt.ylabel('Actual')
plt.show()
```



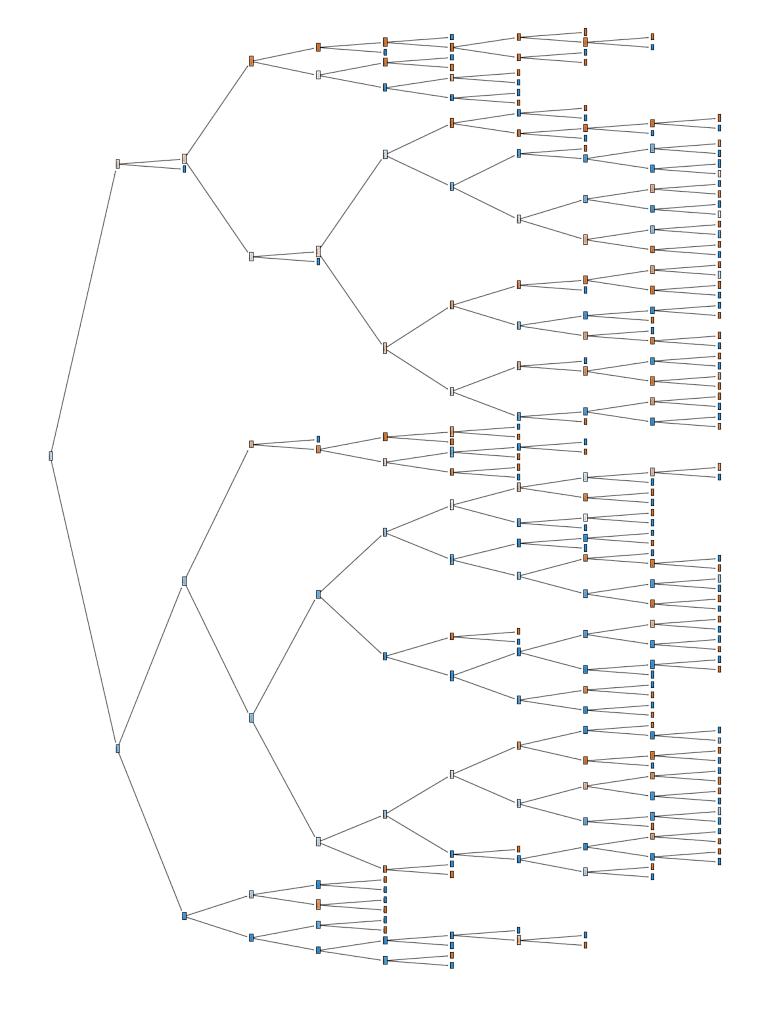
Decision Tree

```
DT_model = DecisionTreeClassifier(max_depth=10,random_state=0)
DT_model.fit(X_train,Y_train)
Y_hat_DT=DT_model.predict(X_test)
print("Accuracy : {}
%".format(accuracy_score(Y_test,Y_hat_DT)*100))
print("Train Accuracy : {}
%".format(accuracy_score(Y_train,DT_model.predict(X_train))*100))
print(classification_report(Y_test,Y_hat_DT))
```

Accuracy : 95	Accuracy : 95.82446373014574%							
Train Accurac	y: 96.08023	374373145%						
	precision	recall	f1-score	support				
0	0.95	0.95	0.95	5302				
1	0.96	0.96	0.96	6912				
accuracy			0.96	12214				
macro avg	0.96	0.96	0.96	12214				
weighted avg	0.96	0.96	0.96	12214				

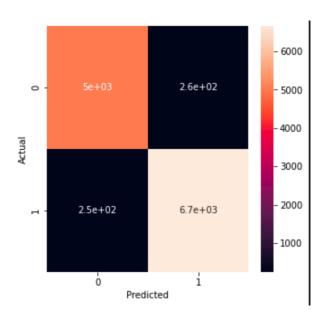
Visualizing the Decision Tree

```
feature_names=df.columns[:-1]
fig=plt.figure(figsize=(25,20))
_=tree.plot_tree(DT_model,feature_names=feature_names,filled=True)
fig.savefig('DT.png')
plt.show()
```



Confusion matrix

```
cm=confusion_matrix(Y_test,Y_hat_DT)
plt.figure(figsize=(5,5))
sn.heatmap(cm,annot=True)
plt.xlabel('Predicted')
plt.ylabel('Actual')
plt.show()
```



Stochastic Gradient Descent Model

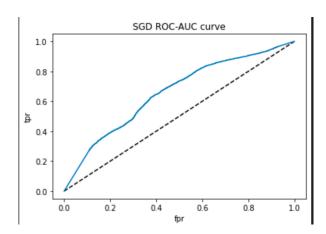
```
from sklearn.linear_model import SGDClassifier
sgd_model= SGDClassifier(loss= 'modified_huber', shuffle=True,
random_state=1)
sgd_model.fit(X_train, Y_train)
Y_hat_SGD=sgd_model.predict(X_test)

print("Accuracy : {}
%".format(accuracy_score(Y_test,Y_hat_SGD)*100))
print("Train Accuracy : {}
%".format(accuracy_score(Y_train,sgd_model.predict(X_train))*100))
print(classification_report(Y_test,Y_hat_SGD))
```

Accuracy : 63.	Accuracy : 63.97576551498281%							
Train Accuracy	Train Accuracy : 63.10306007573432%							
	precision	recall	f1-score	support				
0	0.64	0.38	0.48	5302				
1	0.64	0.84	0.72	6912				
accuracy			0.64	12214				
macro avg	0.64	0.61	0.60	12214				
weighted avg	0.64	0.64	0.62	12214				

ROC-AUC curve

```
y_pred_proba = sgd_model.predict_proba(X_test)[:,1]
fpr, tpr, thresholds = roc_curve(Y_test, y_pred_proba)
plt.plot([0,1],[0,1],'k--')
plt.plot(fpr,tpr)
plt.xlabel('fpr')
plt.ylabel('tpr')
plt.title('SGD_ROC-AUC_curve')
plt.show()
```



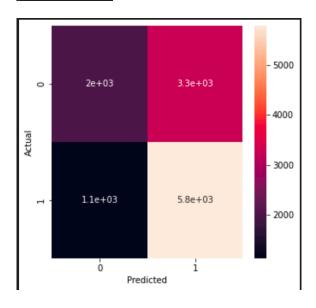
ROC-AUC Score

roc_auc_score(Y_test,y_pred_proba)

0.6576904013771883

Confusion matrix

```
cm=confusion_matrix(Y_test,Y_hat_SGD)
plt.figure(figsize=(5,5))
sn.heatmap(cm,annot=True)
plt.xlabel('Predicted')
plt.ylabel('Actual')
plt.show()
```



Final Accuracies of All the Models

```
data = [['Naive Bayes',
    accuracy_score(Y_train,NB_model.predict(X_train))*100,accuracy_sco
    re(Y_test,Y_hat_NB)*100], ['Logistic
    Regression',accuracy_score(Y_train,LR_model.predict(X_train))*100,
    accuracy_score(Y_test,Y_hat_LR)*100],
    ['KNN',accuracy_score(Y_train,KNN_model.predict(X_train))*100,accu
    racy_score(Y_test,Y_hat_KNN)*100], ['Random
    Forest',accuracy_score(Y_train,RF_model.predict(X_train))*100,accu
    racy_score(Y_test,Y_hat_RF)*100], ['Decision
    Tree',accuracy_score(Y_train,DT_model.predict(X_train))*100,accura
    cy_score(Y_test,Y_hat_DT)*100], ['SGD Method',
    accuracy_score(Y_train,sgd_model.predict(X_train))*100,accuracy_score(Y_test,Y_hat_SGD)*100]]
    accuracy_matrix = pd.DataFrame(data, columns = ['Algo Used',
    'Train Accuracy','Test Accuracy'],index=['1','2','3','4','5','6'])
```

accuracy_matrix

	Algo Used	Train Accuracy	Test Accuracy
1	Naive Bayes	60.047078	59.562797
2	Logistic Regression	64.136731	64.098575
3	KNN	99.946781	99.885377
4	Random Forest	99.095282	98.976584
5	Decision Tree	96.080237	95.824464
6	SGD Method	63.103060	63.975766

By Looking at the above table it is clear that KNN has the best accuracy for the given data set

Conclusion

To classify the same data, we use different machine learning models and they give different accuracies. Using **Naive Bayes**, we received an accuracy of 60.05% on training data and 59.56% on testing data. Using **Logistic Regression**, we received an accuracy of 64.14% on training data and 64.1% on testing data.

For **Random forest** we received accuracy of 99.1% for training data and 98.98% for testing data. For **KNN** classifier it was 99.95% and 99.88% respectively.

Decision Tree was able to classify the training data with a 96.08% accuracy and the testing data with 95.82% accuracy while for **SGD** it was 63.1% and 63.98% respectively.

Hence, we conclude that the best classifiers for the given dataset and parameters are **Random Forest** and **KNN**.

References

- https://archive.ics.uci.edu/ml/datasets/
 Secondary+Mushroom+Dataset Dataset from UCI
- 2) https://www.ncbi.nlm.nih.gov/pmc/articles/
 PMC8046754/ Mushroom data creation, curation, and simulation to support classification tasks-Reference Paper
- 3) Product of bachelor thesis at Philipps-Universität Marburg, Bioinformatics Division, supervised by Dr. G. Hattab -Research Work for collecting and creation of the dataset