```
# Balanced dataset using Random Undersampling method
# Then applied Logistic Regression, Naive Bayes Classifier, and Decision Tree algorithms
# Compared changes in accuracy rates when 10-fold cross-validation applied
import sys
!{sys.executable} -m pip install -U pandas-profiling[notebook]
!jupyter nbextension enable --py widgetsnbextension
!pip install matplotlib
!pip install graphviz
     Requirement already satisfied: pandas-profiling[notebook] in /usr/local/lib/python3.7/di
    WARNING: pandas-profiling 1.4.1 does not provide the extra 'notebook'
    Requirement already satisfied: jinja2>=2.8 in /usr/local/lib/python3.7/dist-packages (fr
     Requirement already satisfied: pandas>=0.19 in /usr/local/lib/python3.7/dist-packages (1
     Requirement already satisfied: matplotlib>=1.4 in /usr/local/lib/python3.7/dist-packages
     Requirement already satisfied: six>=1.9 in /usr/local/lib/python3.7/dist-packages (from
     Requirement already satisfied: MarkupSafe>=0.23 in /usr/local/lib/python3.7/dist-package
     Requirement already satisfied: pyparsing!=2.0.4,!=2.1.2,!=2.1.6,>=2.0.1 in /usr/local/li
     Requirement already satisfied: numpy>=1.11 in /usr/local/lib/python3.7/dist-packages (fr
    Requirement already satisfied: kiwisolver>=1.0.1 in /usr/local/lib/python3.7/dist-packag
     Requirement already satisfied: python-dateutil>=2.1 in /usr/local/lib/python3.7/dist-pac
     Requirement already satisfied: cycler>=0.10 in /usr/local/lib/python3.7/dist-packages (1
     Requirement already satisfied: typing-extensions in /usr/local/lib/python3.7/dist-packag
    Requirement already satisfied: pytz>=2017.3 in /usr/local/lib/python3.7/dist-packages (1
     Enabling notebook extension jupyter-js-widgets/extension...
    Paths used for configuration of notebook:
             /root/.jupyter/nbconfig/notebook.json
           - Validating: OK
    Paths used for configuration of notebook:
             /root/.jupyter/nbconfig/notebook.json
     Requirement already satisfied: matplotlib in /usr/local/lib/python3.7/dist-packages (3.2
     Requirement already satisfied: pyparsing!=2.0.4,!=2.1.2,!=2.1.6,>=2.0.1 in /usr/local/li
    Requirement already satisfied: numpy>=1.11 in /usr/local/lib/python3.7/dist-packages (fr
     Requirement already satisfied: cycler>=0.10 in /usr/local/lib/python3.7/dist-packages (1
     Requirement already satisfied: python-dateutil>=2.1 in /usr/local/lib/python3.7/dist-pac
     Requirement already satisfied: kiwisolver>=1.0.1 in /usr/local/lib/python3.7/dist-packas
     Requirement already satisfied: typing-extensions in /usr/local/lib/python3.7/dist-packas
     Requirement already satisfied: six>=1.5 in /usr/local/lib/python3.7/dist-packages (from
     Requirement already satisfied: graphviz in /usr/local/lib/python3.7/dist-packages (0.10
from google.colab import files
uploaded = files.upload()
```

. , ,,

Choose files | stroke preprocessed.arff

• **stroke preprocessed.arff**(n/a) - 368681 bytes, last modified: 06/03/2022 - 100% done Saving stroke preprocessed.arff to stroke preprocessed.arff

import pandas as pd
from scipy.io import arff

```
import numpy as np
# Timing how long predictors take to run for efficiency calculations
# Import libraries
import time
data file = "stroke preprocessed.arff"
data = arff.loadarff(data_file)
df = pd.DataFrame(data[0])
for col in df.columns:
  if df[col].dtype == 'object':
    # Ensure data isn't read as bytes but rather as strings from file
    df[col] = df[col].str.decode('utf-8')
# Examine data types
print(df.dtypes)
     "id"
                            float64
     "gender"
                             object
     "age"
                            float64
     "hypertension"
                             object
     "heart_disease"
                             object
     "ever_married"
                             object
     "work type"
                             object
     "residence_type"
                             object
     "avg_glucose_level"
                            float64
     "bmi"
                            float64
     "smoking_status"
                             object
     "stroke"
                             object
     dtype: object
# Display first 10 rows
df.head(10)
```

	"id"	"gender"	"age"	"hypertension"	"heart_disease"	"ever_married"	"work_type
0	9046.0	Male	67.0	0	1	Yes	Privat
1	51676.0	Female	61.0	0	0	Yes	Sel employe
2	31112.0	Male	80.0	0	1	Yes	Privat

Examine meta info about data
df.info()

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 5110 entries, 0 to 5109
Data columns (total 12 columns):

#	Column	Non-Null Count	Dtype
0	"id"	5110 non-null	float64
1	"gender"	5110 non-null	object
2	"age"	5110 non-null	float64
3	"hypertension"	5110 non-null	object
4	"heart_disease"	5110 non-null	object
5	"ever_married"	5110 non-null	object
6	"work_type"	5110 non-null	object
7	"residence_type"	5110 non-null	object
8	"avg_glucose_level"	5110 non-null	float64
9	"bmi"	5110 non-null	float64
10	"smoking_status"	5110 non-null	object
11	"stroke"	5110 non-null	object
المارات المسلم	£1+C4/4\ -b	+ (0)	

dtypes: float64(4), object(8)

memory usage: 479.2+ KB

The original 201 null values were all from bmi column, and they have been replaced by place
Convert the 5000 values back into null values
df = df.replace(5000.0, np.nan)

Check head of dataset again
df.head(10)

	"id"	"gender"	"age"	"hypertension"	"heart_disease"	"ever_married"	"work_type
0	9046.0	Male	67.0	0	1	Yes	Privat
1	51676.0	Female	61.0	0	0	Yes	Sel employe
2	31112.0	Male	80.0	0	1	Yes	Privat
3	60182.0	Female	49.0	0	0	Yes	Privat
А	166E N	Eomolo	70 O	1	0	Voc	Sel

Check structure of data types to ensure bmi remains float
df.info()

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 5110 entries, 0 to 5109
Data columns (total 12 columns):

#	Column	Non-Null Count	Dtype
0	"id"	5110 non-null	float64
1	"gender"	5110 non-null	object
2	"age"	5110 non-null	float64
3	"hypertension"	5110 non-null	object
4	"heart_disease"	5110 non-null	object
5	"ever_married"	5110 non-null	object
6	"work_type"	5110 non-null	object
7	"residence_type"	5110 non-null	object
8	"avg_glucose_level"	5110 non-null	float64
9	"bmi"	4909 non-null	float64
10	"smoking_status"	5110 non-null	object
11	"stroke"	5110 non-null	object
		4 - 4	

dtypes: float64(4), object(8)

memory usage: 479.2+ KB

```
# Remove records with NAs from dataset
df_noNA = df
df_noNA = df_noNA.dropna()
df_noNA.head(10)
```

	"id"	"gender"	"age"	"hypertension"	"heart_disease"	"ever_married"	"work_typ
0	9046.0	Male	67.0	0	1	Yes	Priva
2	31112.0	Male	80.0	0	1	Yes	Priva
3	60182.0	Female	49.0	0	0	Yes	Priva
4	1665.0	Female	79.0	1	0	Yes	S ₍ employ
E	E6660 0	Mala	01 N	^	^	Voc	Driv.

See if data is imbalanced on the variable of interest, stroke

Count how many '1's (stroke) and '0's (no stroke) appear
print(df_noNA['"stroke"'].value_counts())

Dataset is quite unbalanced on the stroke variable

Around 209/(4700+209) = 4.3% of dataset is positive for stroke

0 47001 209

Name: "stroke", dtype: int64

+_+

Change 'stroke' attribute into data type float
df_noNA['"stroke"'] = df_noNA['"stroke"'].astype(float)
df_noNA.head(10)

/usr/local/lib/python3.7/dist-packages/ipykernel_launcher.py:2: SettingWithCopyWarning: A value is trying to be set on a copy of a slice from a DataFrame.

Thy using loc[now indexen col indexen] - value instead

print(df_noNA.dtypes)

```
"id"
                      float64
"gender"
                      object
"age"
                      float64
"hypertension"
                      object
"heart_disease"
                      object
"ever_married"
                       object
"work type"
                       object
"residence_type"
                       object
"avg_glucose_level"
                      float64
"bmi"
                      float64
"smoking_status"
                     object
"stroke"
                      float64
dtype: object
 • 0000E.0 MIGIO 1 1.0
```

See if there are any extreme values in numeric data
df_noNA.describe()

	"id"	"age"	"avg_glucose_level"	"bmi"	"stroke"	1
count	4909.000000	4909.000000	4909.000000	4909.000000	4909.000000	
mean	37064.313506	42.865374	105.305150	28.893237	0.042575	
std	20995.098457	22.555115	44.424341	7.854067	0.201917	
min	77.000000	0.080000	55.120000	10.300000	0.000000	
25%	18605.000000	25.000000	77.070000	23.500000	0.000000	
50%	37608.000000	44.000000	91.680000	28.100000	0.000000	
75%	55220.000000	60.000000	113.570000	33.100000	0.000000	
max	72940.000000	82.000000	271.740000	97.600000	1.000000	

```
# Normalize continuous numeric variables
# Such as age, avg_glucose_level, and bmi
# Using z-score methods

# Import libraries for normalization
from sklearn import preprocessing
scaler = preprocessing.MinMaxScaler()

# Only need to normalize continuous numeric variables
var_to_norm = ['"age"', '"avg_glucose_level"', '"bmi"']
df_noNA[var_to_norm] = scaler.fit_transform(df_noNA[var_to_norm])
```

Examine first 10 rows of normalized dataset

```
df_noNA.head()
```

The 3 columns are now standarized to values between 0-1

/usr/local/lib/python3.7/dist-packages/pandas/core/frame.py:3678: SettingWithCopyWarning
A value is trying to be set on a copy of a slice from a DataFrame.

Try using .loc[row indexer,col indexer] = value instead

See the caveats in the documentation: https://pandas.pydata.org/pandas-docs/stable/user_self[col] = igetitem(value, i)

"work_t	"ever_married"	"heart_disease"	"hypertension"	"age"	"gender"	"id"	
Pr	Yes	1	0	0.816895	Male	9046.0	0
Pr	Yes	1	0	0.975586	Male	31112.0	2
Pr	Yes	0	0	0.597168	Female	60182.0	3
empl	Yes	0	1	0.963379	Female	1665.0	4
Pr	Yes	0	0	0.987793	Male	56669.0	5



```
# Create list of categorical columns
cat_cols = ['"gender"', '"hypertension"', '"heart_disease"', '"ever_married"', '"work_type"',
```

```
# Create copy of a data frame in memory w/ a different name
df_dummy = df_noNA.copy()
# Convert only categorical feature into dummy/one-hot features
df_dummy = pd.get_dummies(df_noNA, columns = cat_cols, prefix = cat_cols)
# Print dataset
df_dummy
```

		"id"	"age"	"avg_glucose_level	"bmi"	"stroke"	"gender"_Female	"gend
	0	9046.0	0.816895	0.80126	5 0.301260	1.0	0	
	2	31112.0	0.975586	0.23451	2 0.254296	1.0	0	
	3	60182.0	0.597168	0.53600	3 0.276060	1.0	1	
	4	1665.0	0.963379	0.54934	9 0.156930	1.0	1	
	5	56669.0	0.987793	0.60516	1 0.214204	1.0	0	
			set split selection	import train_test_s	olit			
	5106	44873 N	0 987793	0.323510	340206	0.0	1	
from from impor warni	<pre># Balancing Dataset: Import libraries needed for Random Undersampling method from collections import Counter from imblearn.under_sampling import RandomUnderSampler import warnings warnings.simplefilter(action='ignore', category=FutureWarning)</pre>							
	"id"			float64				
	"age"	.1		float64				
	"bmi"	glucose_le	eveī	float64 float64				
	"strok	'e"		float64				
		r"_Female	<u>م</u>	uint8				
	_	r"_Male	-	uint8				
	_	r"_Other		uint8				
	_	tension"	_0	uint8				
	"hyper	tension"	_1	uint8				
		_disease		uint8				
		_disease'		uint8				
		married"		uint8				
	_	married" type"_Gov	_	uint8 uint8				
	_	—	vc_job ver_worked					
		type"_Pr:		uint8				
			lf-employe					
		type"_ch:		uint8				
		lence_type		uint8				
	"resid	lence_type	e"_Urban	uint8				
			s"_Unknown					
			s"_formerl					
			s"_never s					
		.ng_status object	s"_smokes	uint8				
)						

[#] Partition the class of interest, stroke (Dependent Variable), from the Independent Variable

```
iv classes = df dummy.iloc[:,:-1]
dv_class = df_dummy.iloc[:,-1]
# Set class name as "stroke", all other attributes will be used as features
class_col_name = '"stroke"'
# Obtain necessary dummy feature names
dummy feature name = df dummy.columns.values.tolist()
dummy_feature_names = dummy_feature_name[5:]
# 70% training, 30% test set split
x_train, x_test, y_train, y_test = train_test_split(df_dummy.loc[:, dummy_feature_names], df_
# Display the class distribution in its original split
print("Original class split prior to undersampling: ", Counter(y_train))
     Original class split prior to undersampling: Counter({0.0: 3291, 1.0: 145})
# Implement Random Undersampler Model
undersampler = RandomUnderSampler(sampling strategy = 'majority')
# Apply data into Random Undersampler Model
x_train_under, y_train_under = undersampler.fit_resample(x_train, y_train)
# Display the class distribution after the Random Undersampler is applied
print("Class split after Random Undersampling: ", Counter(y_train_under))
     Class split after Random Undersampling: Counter({0.0: 145, 1.0: 145})
# Import required libraries
from sklearn.svm import SVC
from sklearn.metrics import classification report, roc auc score
# Evaluate the Random Undersampler
model = SVC()
clf_undersampler = model.fit(x_train_under, y_train_under)
pred_undersampler = clf_undersampler.predict(x_test)
# Display ROC AUC score
print("Random Undersampled data - ROC AUC: ", roc_auc_score(y_test, pred_undersampler))
     Random Undersampled data - ROC AUC: 0.6595768275372603
start = time.time()
# Import needed libraries for Logistic Regression Model
```

```
from sklearn.linear model import LogisticRegression
# Begin to Implement Logistic Regression Model
log regr = LogisticRegression()
# Apply data into Logistic Regression Model
log regr.fit(x train, y train)
y_pred = log_regr.predict(x_test)
# Obtain Confusion Matrix and Evaluation Metrics for the Logistic Regression Model
from sklearn import metrics
cnf_matrix = metrics.confusion_matrix(y_test, y_pred)
cnf matrix
    array([[1409,
                      0],
           [ 64,
                      0]])
# Display Evaluation Metrics for Logistic Regression Model
print("Logistic Regression Accuracy:\t", metrics.accuracy_score(y_test, y_pred))
print("Logistic Regression Precision:\t",metrics.precision score(y test, y pred))
print("Logistic Regression Recall:\t",metrics.recall_score(y_test, y_pred))
    Logistic Regression Accuracy:
                                      0.956551255940258
     Logistic Regression Precision:
                                      0.0
    Logistic Regression Recall:
                                      0.0
     /usr/local/lib/python3.7/dist-packages/sklearn/metrics/_classification.py:1318: Undefine
       _warn_prf(average, modifier, msg_start, len(result))
# Import libraries for cross-validation
from sklearn.model selection import cross val score, cross val predict
# 10-Fold Cross Validation for Logistic Regression
cv lr = cross val score(log regr, df dummy, df dummy[class col name], cv=10)
print("Cross-validated scores:\t", cv_lr)
# Increased Accuracy score from 0.957 to 1
     /usr/local/lib/python3.7/dist-packages/sklearn/linear model/ logistic.py:818: Convergence
    STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.
    Increase the number of iterations (max iter) or scale the data as shown in:
         https://scikit-learn.org/stable/modules/preprocessing.html
    Please also refer to the documentation for alternative solver options:
         https://scikit-learn.org/stable/modules/linear model.html#logistic-regression
       extra_warning_msg=_LOGISTIC_SOLVER_CONVERGENCE_MSG,
     /usr/local/lib/python3.7/dist-packages/sklearn/linear_model/_logistic.py:818: Convergence
    STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.
    Increase the number of iterations (max_iter) or scale the data as shown in:
         https://scikit-learn.org/stable/modules/preprocessing.html
```

```
Please also refer to the documentation for alternative solver options:
         https://scikit-learn.org/stable/modules/linear_model.html#logistic-regression
       extra warning msg= LOGISTIC SOLVER CONVERGENCE MSG,
     Cross-validated scores: [1.
                                          1.
                                                                1.
                                                                            1.
                                                                                       1.
                            0.95723014 0.95918367]
      1.
                 1.
# Cross validation accuracy for Logistic Regression (R2 score)
predictions = cross val predict(log regr, df dummy, df dummy[class col name], cv=10)
accuracy = metrics.r2 score(df dummy[class col name], predictions)
print("Cross-Predicted Accuracy for Logistic Regression: ", accuracy)
     /usr/local/lib/python3.7/dist-packages/sklearn/linear model/ logistic.py:818: Convergence
     STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.
     Increase the number of iterations (max iter) or scale the data as shown in:
         https://scikit-learn.org/stable/modules/preprocessing.html
     Please also refer to the documentation for alternative solver options:
         https://scikit-learn.org/stable/modules/linear model.html#logistic-regression
       extra warning msg= LOGISTIC SOLVER CONVERGENCE MSG,
     /usr/local/lib/python3.7/dist-packages/sklearn/linear model/ logistic.py:818: Convergence
     STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.
     Increase the number of iterations (max iter) or scale the data as shown in:
         https://scikit-learn.org/stable/modules/preprocessing.html
     Please also refer to the documentation for alternative solver options:
         https://scikit-learn.org/stable/modules/linear model.html#logistic-regression
       extra warning msg= LOGISTIC SOLVER CONVERGENCE MSG,
     Cross-Predicted Accuracy for Logistic Regression: 0.7951043469408531
end = time.time()
print("Time to run Logistic Regression: ", end - start)
     Time to run Logistic Regression: 2.5281853675842285
start = time.time()
# Naive Bayes modeling
from sklearn.naive bayes import MultinomialNB
# Create Multinomial NB Classifier
nb = MultinomialNB()
# Train model using training sets
nb.fit(x train under, y train under)
     MultinomialNB()
```

```
# Predict response for test dataset
y pred = nb.predict(x test)
# Print Naive Bayes output
print("Number of features used: ", nb.n_features_)
print("Classes: ", nb.classes )
print("Number of records for classes: ", nb.class_count_)
print("Log prior probability for classes: ", nb.class_log_prior_)
print("Log conditional probability for each feature given a class: ", nb.feature_log_prob_)
     Number of features used: 20
     Classes: [0. 1.]
     Number of records for classes: [145. 145.]
     Log prior probability for classes: [-0.69314718 -0.69314718]
     Log conditional probability for each feature given a class: [[-2.32703619 -3.11351531 -
       -4.86271516 -3.13549422 -2.31718389 -4.05178495 -6.94215671 -2.38827981
       -3.80666249 -4.37720735 -2.57270885 -2.722649 -3.27859506 -3.64631984
       -2.84781214 -3.80666249]
      [-2.49950545 -2.81502232 -6.94215671 -2.28819636 -3.20448709 -2.16303321
       -3.6099522 -4.2341065 -2.05935478 -3.85111425 -6.94215671 -2.4312972
       -3.38680864 -6.94215671 -2.75250196 -2.54770755 -3.94642443 -3.18095659
       -2.93482352 -3.5081695 ]]
# Get Naive Bayes Classifier Confusion matrix
from sklearn.metrics import confusion matrix
cf = confusion matrix(y test, y pred)
print("Confusion Matrix")
print(cf)
tn, fp, fn, tp = cf.ravel()
print("TP: ", tp, ", FP: ", fp, ", TN: ", tn, ", FN: ", fn)
     Confusion Matrix
     [[927 482]
      [ 24 40]]
     TP: 40 , FP: 482 , TN: 927 , FN: 24
# Get Naive Bayes Classifier Classifier report
from sklearn.metrics import classification report
from sklearn import metrics
print(classification report(y test, y pred))
                   precision
                                recall f1-score
                                                   support
              0.0
                        0.97
                                  0.66
                                            0.79
                                                      1409
              1.0
                        0.08
                                  0.62
                                            0.14
                                                        64
                                            0.66
                                                      1473
         accuracy
                        0.53
                                  0.64
                                            0.46
                                                      1473
        macro avg
```

Import libraries for plotting the decision tree

```
# Display Evaluation Metrics for Naive Bayes Classifier
print("Naive Bayes Classifier Accuracy:\t", metrics.accuracy_score(y_test, y_pred))
print("Naive Bayes Classifier Precision:\t",metrics.precision_score(y_test, y_pred))
print("Naive Bayes Classifier Recall:\t\t",metrics.recall_score(y_test, y_pred))
    Naive Bayes Classifier Accuracy:
                                              0.6564833672776647
    Naive Bayes Classifier Precision:
                                              0.07662835249042145
    Naive Bayes Classifier Recall:
                                              0.625
# 10-Fold Cross Validation for Naive Bayes Classifier
cv_nb = cross_val_score(nb, df_dummy, df_dummy[class_col_name], cv=10)
print("Cross-validated scores:\t", cv nb)
# Increased Accuracy score from 0.545 to 0.998
    Cross-validated scores: [0.99796334 0.99592668 0.99796334 0.99796334 0.99592668 0.99389
      0.99592668 0.99796334 0.99796334 0.99795918]
# Cross validation accuracy for Naive Bayes Classifier (R2 score)
predictions = cross_val_predict(nb, df_dummy, df_dummy[class_col_name], cv=10)
accuracy = metrics.r2_score(df_dummy[class_col_name], predictions)
print("Cross-Predicted Accuracy for Naive Bayes Classifier: ", accuracy)
    Cross-Predicted Accuracy for Naive Bayes Classifier: 0.9250381757100682
end = time.time()
print("Time to run Naive Bayes Classifier: ", end - start)
    Time to run Naive Bayes Classifier: 0.42337512969970703
start = time.time()
# Decision tree on dummy encoded data
from sklearn import tree
clf = tree.DecisionTreeClassifier(max depth = 5) # 5 levels set
clf = clf.fit(x_train_under, y_train_under)
import graphviz
# Obtain unique class values to show on tree
class values = df dummy[class col name]. unique()
print("class names: ", class_values)
    class names: [1. 0.]
```

```
import matplotlib
from matplotlib import pyplot as plt

# Plot decision tree
fig = plt.figure(figsize=(25,20))
_ = tree.plot_tree(clf, feature_names = dummy_feature_names, class_names = str(class_values),
```

Save decision tree figure fig.savefig("decision_tree7.png")

"smoking_status" Unknown <= 0.5 | "residence_type" Urban <= 0.5 | gini = 0.257 | gini = 0.455 |

"heart_disease"_0 <= 0.5 gini = 0.499

Perform prediction on test set y_pred = clf.predict(x_test)

Get decision tree confusion matrix cf = confusion_matrix(y_test, y_pred) print("Confusion Matrix") print(cf) tn, fp, fn, tp = cf.ravel()

> Confusion Matrix [[740 669] [22 42]]

TP: 42 , FP: 669 , TN: 740 , FN: 22

print("TP: ", tp, ", FP: ", fp, ", TN: ", tn, ", FN: ", fn)

Get decision tree report from sklearn.metrics import classification_report from sklearn import metrics print(classification_report(y_test, y_pred))

Undersampling can lead to overfitting

	precision	recall	f1-score	support
0.0	0.97	0.53	0.68	1409
1.0	0.06	0.66	0.11	64
accuracy			0.53	1473
macro avg	0.52	0.59	0.40	1473
weighted avg	0.93	0.53	0.66	1473

Display Evaluation Metrics for Decision Tree print("Decision Tree Accuracy:\t\t", metrics.accuracy_score(y_test, y_pred)) print("Decision Tree Precision:\t",metrics.precision_score(y_test, y_pred)) print("Decision Tree Recall:\t\t",metrics.recall_score(y_test, y_pred))

Decision Tree Accuracy:

0.5308893414799728

```
Decision Tree Precision:
                                      0.05907172995780591
     Decision Tree Recall:
                                      0.65625
# 10-Fold Cross Validation for Decision Tree
cv dt = cross val score(clf, df dummy, df dummy[class col name], cv=10)
print("Cross-validated scores:\t", cv_dt)
# Increased Accuracy score from 0.759 to 0.998
     Cross-validated scores: [1. 1. 1. 1. 1. 1. 1. 1. 1. ]
# Cross validation accuracy for Decision Tree (R2 score)
predictions = cross_val_predict(clf, df_dummy, df_dummy[class_col_name], cv=10)
accuracy = metrics.r2_score(df_dummy[class_col_name], predictions)
print("Cross-Predicted Accuracy for Decision Tree: ", accuracy)
     Cross-Predicted Accuracy for Decision Tree: 1.0
end = time.time()
print("Time to run Decision Tree: ", end - start)
     Time to run Decision Tree: 3.2847201824188232
```