```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
import warnings
from sklearn.model_selection import KFold, StratifiedKFold, cross_val_score
from sklearn import linear_model, tree, ensemble
```

dataframe=pd.read\_csv("/content/Heart\_Disease\_Prediction.csv")
dataframe=dataframe.dropna()
dataframe.head(10)

	Age	Sex	Chest pain type	ВР	Cholesterol	FBS over 120	EKG results	Max HR	Exercise angina	ST depression	Slope of ST
0	70	1	4	130	322	0	2	109	0	2.4	2
1	67	0	3	115	564	0	2	160	0	1.6	2
2	57	1	2	124	261	0	0	141	0	0.3	1
3	64	1	4	128	263	0	0	105	1	0.2	2
4	74	0	2	120	269	0	2	121	1	0.2	1
5	65	1	4	120	177	0	0	140	0	0.4	1
6	56	1	3	130	256	1	2	142	1	0.6	2
7	59	1	4	110	239	0	2	142	1	1.2	2
4											•

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- ◀											•

dataframe.isna().sum()

```
Age
Sex
                          0
Chest pain type
BP
Cholesterol
                          0
FBS over 120
EKG results
                          0
Max HR
Exercise angina
                          0
ST depression
Slope of ST
Number of vessels fluro
Thallium
Heart Disease
dtype: int64
```

```
plt.figure(figsize=(15,10))
sns.heatmap(dataframe.corr(),linewidth=.01,annot=True,cmap="winter")
plt.show()
plt.savefig('correlationfigure')
```

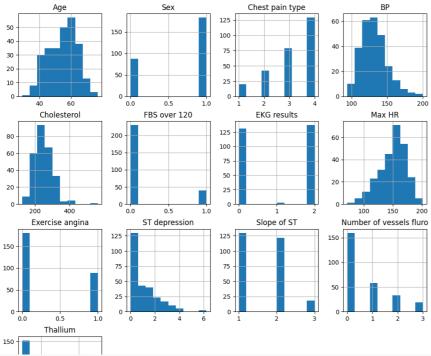
 $\supseteq$ 

<ipython-input-25-762dfe51a80e>:2: FutureWarning: The default value of numeric\_only in DataFrame.corr is deprecated. In a future ver sns.heatmap(dataframe.corr(),linewidth=.01,annot=True,cmap="winter")

		(,,		•	,		,								
Age -	1	-0.094	0.097	0.27	0.22	0.12	0.13	-0.4	0.098	0.19	0.16	0.36	0.11		1.0
Sex -	-0.094	1	0.035	-0.063	-0.2	0.042	0.039	-0.076	0.18	0.097	0.051	0.087	0.39		- 0.8
Chest pain type -	0.097	0.035	1	-0.043	0.09	-0.099	0.074	-0.32	0.35	0.17	0.14	0.23	0.26		
BP -	0.27	-0.063	-0.043	1	0.17	0.16	0.12	-0.039	0.083	0.22	0.14	0.086	0.13	_	0.6
Cholesterol -	0.22	-0.2	0.09	0.17	1	0.025	0.17	-0.019	0.078	0.028	-0.0058	0.13	0.029		
FBS over 120 -	0.12	0.042	-0.099	0.16	0.025	1	0.053	0.022	-0.0041	-0.026	0.044	0.12	0.049	-	0.4
EKG results -	0.13	0.039	0.074	0.12	0.17	0.053	1	-0.075	0.095	0.12	0.16	0.11	0.0073		
Max HR -	-0.4	-0.076	-0.32	-0.039	-0.019	0.022	-0.075	1	-0.38	-0.35	-0.39	-0.27	-0.25	-	0.2
Exercise angina -	0.098	0.18	0.35	0.083	0.078	-0.0041	0.095	-0.38	1	0.27	0.26	0.15	0.32		
ST depression -	0.19	0.097	0.17	0.22	0.028	-0.026	0.12	-0.35	0.27	1	0.61	0.26	0.32	-	0.0
Slope of ST -	0.16	0.051	0.14	0.14	-0.0058	0.044	0.16	-0.39	0.26	0.61	1	0.11	0.28		
Number of vessels fluro -	0.36	0.087	0.23	0.086	0.13	0.12	0.11	-0.27	0.15	0.26	0.11	1	0.26	-	-0.2
Thallium -	0.11	0.39	0.26	0.13	0.029	0.049	0.0073	-0.25	0.32	0.32	0.28	0.26	1		
	Age -	- Sex -	Chest pain type -	- BP	Cholesterol -	FBS over 120 -	EKG results -	Max HR -	Exercise angina -	ST depression -	Slope of ST -	Number of vessels fluro -	Thallium -		0.4

<Figure size 640x480 with 0 Axes>

dataframe.hist(figsize=(12,12))
plt.savefig('featuresplot')



X=dataframe.iloc[:,:-1].values y=dataframe.iloc[:,-1].values JU | import pandas as pd from sklearn.model\_selection import train\_test\_split # Split the data into training and testing sets X\_train, X\_test, y\_train, y\_test = train\_test\_split(X, y, test\_size=0.5, random\_state=40) # For X\_train and y\_train: train\_data = pd.DataFrame(X\_train) train\_data['target'] = y\_train # Remove rows with NaN values train\_data\_cleaned = train\_data.dropna() # Separate X\_train and y\_train from the cleaned data X\_train = train\_data\_cleaned.drop('target', axis=1) y\_train = train\_data\_cleaned['target'] # For X\_test and y\_test: test\_data = pd.DataFrame(X\_test) test\_data['target'] = y\_test # Remove rows with NaN values test\_data\_cleaned = test\_data.dropna() # Separate X\_test and y\_test from the cleaned data X\_test = test\_data\_cleaned.drop('target', axis=1) y\_test = test\_data\_cleaned['target'] # Now X\_train\_cleaned, y\_train\_cleaned, X\_test\_cleaned, and y\_test\_cleaned do not contain NaN values and are 1-dimensional.

```
from sklearn.model_selection import cross_val_score, GridSearchCV
from sklearn.linear_model import LogisticRegression
lr=LogisticRegression()
model1=lr.fit(X_train,y_train)
prediction1=model1.predict(X_test)
from \ sklearn.metrics \ import \ confusion\_matrix
cm=confusion_matrix(y_test,prediction1)
cm
sns.heatmap(cm, annot=True,cmap='winter',linewidths=0.3, linecolor='black',annot_kws={"size": 20})
TP=cm[0][0]
TN=cm[1][1]
FN=cm[1][0]
FP=cm[0][1]
precision1=TP/(TP+FP)
recall1=TP/(TP+FN)
print('Testing Accuracy for Logistic Regression:',(TP+TN)/(TP+TN+FP+FP))
print('Testing Sensitivity for Logistic Regression:',recall1)
print('Testing Specificity for Logistic Regression:',(TN/(TN+FP)))
print('Testing Precision for Logistic Regression:',precision1)
print('Testing F1-measure for Logistic Regression:',(2*precision1*recall1)/(precision1+recall1))
     /usr/local/lib/python3.10/dist-packages/sklearn/linear model/ logistic.py:458: Conver
```

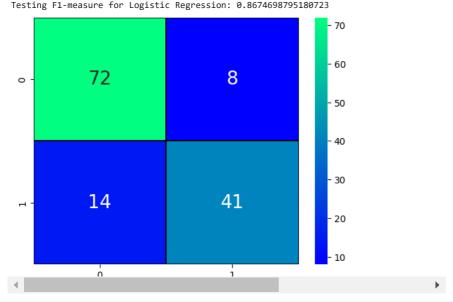
STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.

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

n\_iter\_i = \_check\_optimize\_result( Testing Accuracy for Logistic Regression: 0.837037037037 Testing Sensitivity for Logistic Regression: 0.8372093023255814 Testing Specificity for Logistic Regression: 0.8367346938775511 Testing Precision for Logistic Regression: 0.9



```
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import accuracy_score
rfc=RandomForestClassifier()
model3 = rfc.fit(X_train, y_train)
prediction3 = model3.predict(X_test)
cm3=confusion_matrix(y_test, prediction3)
sns.heatmap(cm3, annot=True,cmap='winter',linewidths=0.3, linecolor='black',annot_kws={"size": 20})
TP=cm3[0][0]
TN=cm3[1][1]
FN=cm3[1][0]
FP=cm3[0][1]
precision3=TP/(TP+FP)
recall3=TP/(TP+FN)
print(round(accuracy score(prediction3,y test)*100,2))
print('Testing Accuracy for Random Forest:',(TP+TN)/(TP+TN+FP+FP))
print('Testing Sensitivity for Random Forest:',recall3)
print('Testing \ Specificity \ for \ Random \ Forest:',(TN/(TN+FP)))
print('Testing Precision for Random Forest:',precision3)
print('Testing F1-measure for Random Forest:',(2*precision3*recall3)/(precision3+recall3))
```

81.48

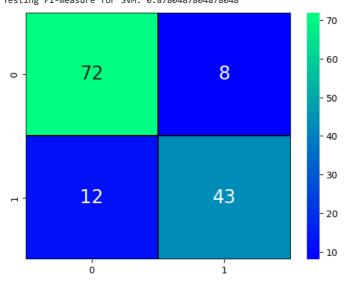
```
Testing Sensitivity for Random Forest: 0.8313253012048193
Testing Specificity for Random Forest: 0.7884615384615384
Testing Precision for Random Forest: 0.8625
Testing F1-measure for Random Forest: 0.8466257668711658

- 60
- 11
- 50
- 40
- 30
```

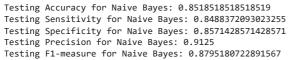
Testing Accuracy for Random Forest: 0.8148148148148

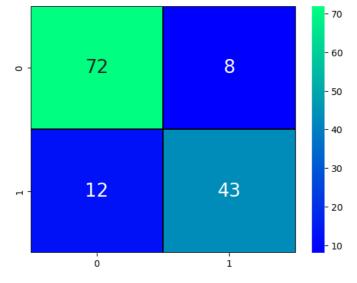
```
from sklearn.svm import SVC
from sklearn.metrics import confusion_matrix, classification_report
import seaborn as sns
svm = SVC(C=12, kernel='linear')
model4 = svm.fit(X_train, y_train)
prediction4 = model4.predict(X_test)
cm4 = confusion_matrix(y_test, prediction4)
sns.heatmap(cm4, annot=True, cmap='winter', linewidths=0.3, linecolor='black', annot_kws={"size": 20})
TP = cm4[0][0]
TN = cm4[1][1]
FN = cm4[1][0]
FP = cm4[0][1]
precision4=TP/(TP+FP)
recall4=TP/(TP+FN)
print('Testing Accuracy for SVM:', (TP + TN) / (TP + TN + FN + FP))
print('Testing Sensitivity for SVM:', recall4)
print('Testing Specificity for SVM:', (TN / (TN + FP)))
print('Testing Precision for SVM:', precision4)
print('Testing F1-measure for SVM:',(2*precision4*recall4)/(precision4+recall4))
```

Testing Accuracy for SVM: 0.8518518518518519
Testing Sensitivity for SVM: 0.8571428571428571
Testing Specificity for SVM: 0.8431372549019608
Testing Precision for SVM: 0.9
Testing F1-measure for SVM: 0.8780487804878048



```
from sklearn.naive_bayes import GaussianNB
nb = GaussianNB()
model5=nb.fit(X_train,y_train)
prediction5 = nb.predict(X_test)
cm5=confusion_matrix(y_test, prediction5)
#score_nb = round(accuracy_score(y_pred_nb,y_test)*100,2)
sns.heatmap(cm4, annot=True, cmap='winter', linewidths=0.3, linecolor='black', annot_kws={"size": 20})
TP = cm5[0][0]
TN = cm5[1][1]
FN = cm5[1][0]
FP = cm5[0][1]
precision5=TP/(TP+FP)
recall5=TP/(TP+FN)
print('Testing Accuracy for Naive Bayes:', (TP + TN) / (TP + TN + FN + FP))
print('Testing Sensitivity for Naive Bayes:', recall5)
\label{eq:print(Testing Specificity for Naive Bayes:', (TN / (TN + FP)))}
print('Testing Precision for Naive Bayes:', precision5)
print('Testing F1-measure for Naive Bayes:',(2*precision5*recall5)/(precision5+recall5))
#print("The accuracy score achieved using Naive Bayes is: "+str(score_nb)+" %")
```





Heart Disease ['Absence']

```
from \ sklearn.tree \ import \ Decision Tree Classifier
max accuracy = 0
for x in range(200):
    dt = DecisionTreeClassifier(random_state=x)
    dt.fit(X_train,y_train)
    y_pred_dt = dt.predict(X_test)
    current_accuracy = round(accuracy_score(y_pred_dt,y_test)*100,2)
    if(current_accuracy>max_accuracy):
        max_accuracy = current_accuracy
        best x = x
#print(max accuracy)
#print(best_x)
dt = DecisionTreeClassifier(random_state=best_x)
model6=dt.fit(X train,y train)
prediction6= dt.predict(X_test)
cm6=confusion_matrix(y_test, prediction6)
sns.heatmap(cm4, annot=True, cmap='winter', linewidths=0.3, linecolor='black', annot_kws={"size": 20})
TP = cm6[0][0]
TN = cm6[1][1]
FN = cm6[1][0]
FP = cm6[0][1]
precision6=TP/(TP+FP)
recall6=TP/(TP+FN)
\label{eq:print(Testing Accuracy for DecisionTreeClassifier:', (TP + TN) / (TP + TN + FN + FP))} \\
print('Testing Sensitivity for DecisionTreeClassifier:', recall6)
print('Testing Specificity for DecisionTreeClassifier:', (TN / (TN + FP)))
print('Testing Precision for DecisionTreeClassifier:', precision6)
print('Testing F1-measure for DecisionTreeClassifier:'.(2*precision6*pecall6)/(precision6+pecall6))
import numpy as np
# Assuming new_data is your new dataset with features for prediction
new_data = np.array([54, 0, 3, 110, 214, 0, 0, 158, 0, 1.6, 2, 0, 3])
# Reshape to 2D array
new_data_2d = new_data.reshape(1, -1) # or new_data[np.newaxis, :]
# Make predictions
new_prediction = model4.predict(new_data_2d)
# Display the predicted target values
print(" Heart Disease ",end='')
print(new_prediction)
```