

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
from sklearn.model_selection import train_test_split
from sklearn.metrics import confusion_matrix, accuracy_score, mean_squared_error, r2_score
from sklearn.neighbors import KNeighborsClassifier
from sklearn.svm import SVC
from sklearn.tree import DecisionTreeClassifier
from sklearn.ensemble import RandomForestClassifier
from sklearn.ensemble import GradientBoostingClassifier
import seaborn as sns
```

```
from google.colab import files
data=files.upload()
```

Choose Files No file chosen

Upload widget is only available when the cell has been executed in the current browser session. Please rerun this cell to enable.

Saving Dataset of Diabetes 1.csv to Dataset of Diabetes 1 (1).csv

```
df=pd.read_csv("/content/Dataset of Diabetes 1 (1).csv")
```

df

	ID	Gender	AGE	HbA1c	HDL	LDL	BMI	CLASS
0	502	F	50	4.9	2.4	1.4	24.0	N
1	735	M	26	4.9	1.1	2.1	23.0	N
2	420	F	50	4.9	2.4	1.4	24.0	N
3	680	F	50	4.9	2.4	1.4	24.0	N
4	504	M	33	4.9	0.8	2.0	21.0	N
...	...	...	...	...	...	...	...	...
494	397	M	55	11.7	1.3	2.3	30.0	Y
495	681	F	58	9.0	3.2	1.4	35.0	Y
496	749	M	55	10.0	1.2	3.4	33.0	Y
497	321	F	54	10.7	1.1	4.2	39.0	Y
498	381	M	58	8.0	0.9	2.0	29.0	Y

499 rows × 8 columns

```
arr=np.array(df)
arr
array([[502, 'F', 50, ..., 1.4, 24.0, 'N'],
       [735, 'M', 26, ..., 2.1, 23.0, 'N'],
       [420, 'F', 50, ..., 1.4, 24.0, 'N'],
       ...,
       [749, 'M', 55, ..., 3.4, 33.0, 'Y'],
       [321, 'F', 54, ..., 4.2, 39.0, 'Y'],
       [381, 'M', 58, ..., 2.0, 29.0, 'Y']], dtype=object)
```

```
df.isnull().sum()
ID      0
Gender  0
AGE      0
HbA1c   0
HDL      0
LDL      0
BMI      0
CLASS   0
dtype: int64
```

```
df.isnull().sum().sum()
0
```

```
df.shape
(499, 8)
```

```
df.replace({'Gender':{'M':0,'F':1}},inplace=True)
```

```
x=df.drop(columns='ID',axis=1)
df=x

x=df.drop(columns='CLASS',axis=1)
y=df['CLASS']
train_x,test_x,train_y,test_y=train_test_split(x,y,test_size=(0.3))
train_x.shape
```

(349, 6)

train\_x

	Gender	AGE	HbA1c	HDL	LDL	BMI
313	0	60	9.0	0.80	3.70	30.0
35	1	39	4.0	1.10	2.60	22.0
432	0	52	7.9	1.10	2.50	28.0
116	0	50	6.2	0.60	1.00	24.0
362	1	60	6.2	1.00	4.40	27.0
...	...	...	...	...	...	...
268	1	55	5.9	1.00	2.00	30.0
117	1	49	6.0	0.75	1.35	23.0
90	0	30	5.6	0.90	3.30	24.5
49	1	47	5.0	1.00	2.40	22.0
59	1	44	4.1	1.00	3.50	21.0

349 rows × 6 columns

test\_x.shape

(150, 6)

test\_x

	Gender	AGE	HbA1c	HDL	LDL	BMI
98	0	60	0.9	1.1	3.6	24.0
275	0	56	7.6	0.9	3.3	26.0
66	1	35	5.0	1.3	2.4	20.0
179	0	49	5.2	3.9	0.8	24.0
339	1	61	11.5	1.1	2.5	26.0
...	...	...	...	...	...	...
128	0	50	5.9	1.0	3.7	25.0
195	0	50	4.0	1.1	3.2	23.0
401	0	60	6.8	0.7	4.1	33.0
264	1	55	5.9	1.0	2.0	30.0
14	1	50	4.0	1.2	2.2	24.0

150 rows × 6 columns

```
pd.DataFrame(train_x).describe()
```

```

Gender      AGE      HbA1c      HDL      LDL      BMI

train_y.shape

(349,)

std      0.498739      9.138748      2.099589      0.587388      1.057292      5.331973

from sklearn.preprocessing import StandardScaler

sc=StandardScaler()

from sklearn.preprocessing import StandardScaler
sc=StandardScaler()
train_x_sc=sc.fit_transform(train_x)
pd.DataFrame(train_x_sc).describe()

```

	0	1	2	3	4	5
<b>count</b>	3.490000e+02	3.490000e+02	3.490000e+02	3.490000e+02	3.490000e+02	3.490000e+02
<b>mean</b>	1.030694e-16	2.048664e-16	1.864157e-16	-3.200242e-16	-1.072049e-16	4.549051e-16
<b>std</b>	1.001436e+00	1.001436e+00	1.001436e+00	1.001436e+00	1.001436e+00	1.001436e+00
<b>min</b>	-9.147907e-01	-2.517858e+00	-2.915789e+00	-1.163042e+00	-2.217844e+00	-1.744950e+00
<b>25%</b>	-9.147907e-01	-5.453945e-01	-7.694347e-01	-4.810838e-01	-7.023716e-01	-8.058646e-01
<b>50%</b>	-9.147907e-01	3.312557e-01	-1.016802e-01	-1.401044e-01	-1.340694e-01	-5.459612e-02
<b>75%</b>	1.093146e+00	5.504183e-01	6.137711e-01	2.008749e-01	7.183840e-01	8.657077e-01
<b>max</b>	1.093146e+00	2.851625e+00	3.189396e+00	9.236828e+00	6.874991e+00	2.809615e+00

```

from sklearn.linear_model import Perceptron
p=Perceptron()
p.fit(train_x_sc,train_y)
train_pred=p.predict(train_x_sc)
test_pred=p.predict(test_x)

```

/usr/local/lib/python3.7/dist-packages/sklearn/base.py:444: UserWarning: X has feature names, but Perceptron was fitted without feature names, but {self.\_\_class\_\_.\_\_name\_\_} was fitted without"

```

from sklearn.metrics import accuracy_score
print('Training Accuracy =',accuracy_score(train_pred,train_y))

```

Training Accuracy = 0.7449856733524355

```
print('Test Accuracy =',accuracy_score(test_y,test_pred))
```

Test Accuracy = 0.6866666666666666

```

from sklearn.svm import SVC
svc=SVC()

```

```
svc.fit(train_x_sc,train_y)
```

SVC()

```

train_pred_svc=svc.predict(train_x_sc)
test_pred_svc=svc.predict(test_x)

```

/usr/local/lib/python3.7/dist-packages/sklearn/base.py:444: UserWarning: X has feature names, but SVC was fitted without feature names, but {self.\_\_class\_\_.\_\_name\_\_} was fitted without"

```
print("Training",accuracy_score(train_pred_svc,train_y))
```

Training 0.9054441260744985

```
print("Test",accuracy_score(test_pred_svc,test_y))
```

Test 0.6866666666666666

```

svc1=SVC(kernel="rbf",C=0.5)
svc1.fit(train_x_sc,train_y)

```

```

train_pred_svc1=svc1.predict(train_x_sc)
test_pred_svc1=svc1.predict(test_x)
/usr/local/lib/python3.7/dist-packages/sklearn/base.py:444: UserWarning: X has feature names, but SVC was fitted without feature na
f"X has feature names, but {self.__class__.__name__} was fitted without"

print("Train",accuracy_score(train_pred_svc1,train_y))

Train 0.8681948424068768

print("test",accuracy_score(test_pred_svc1,test_y))

test 0.6866666666666666

from sklearn.neighbors import KNeighborsClassifier
knn=KNeighborsClassifier(n_neighbors=3)
knn.fit(train_x_sc,train_y)
train_pred_knn=knn.predict(train_x_sc)
test_pred_knn=knn.predict(test_x)

/usr/local/lib/python3.7/dist-packages/sklearn/base.py:444: UserWarning: X has feature names, but KNeighborsClassifier was fitted w
f"X has feature names, but {self.__class__.__name__} was fitted without"

print("Train accuracy",accuracy_score(train_pred_knn,train_y))
print("test accuracy",accuracy_score(test_pred_knn,test_y))

Train accuracy 0.9226361031518625
test accuracy 0.6866666666666666

from sklearn.tree import DecisionTreeClassifier
dt=DecisionTreeClassifier()
dt.fit(train_x_sc,train_y)
train_pred_dt=dt.predict(train_x_sc)
test_pred_dt=dt.predict(test_x)

/usr/local/lib/python3.7/dist-packages/sklearn/base.py:444: UserWarning: X has feature names, but DecisionTreeClassifier was fitted
f"X has feature names, but {self.__class__.__name__} was fitted without"

print("Training",accuracy_score(train_pred_dt,train_y))
print("testing",accuracy_score(test_pred_dt,test_y))

Training 1.0
testing 0.6866666666666666

dt1=DecisionTreeClassifier(criterion='entropy')
dt1.fit(train_x_sc,train_y)
train_pred_dt1=dt1.predict(train_x_sc)
test_pred_dt1=dt1.predict(test_x)

print("Training value",accuracy_score(train_pred_dt1,train_y))
print("testing value",accuracy_score(test_pred_dt1,test_y))

Training value 1.0
testing value 0.6866666666666666

from sklearn.ensemble import RandomForestClassifier
rf=RandomForestClassifier()
rf.fit(train_x_sc,train_y)
train_pred_rf=rf.predict(train_x_sc)
test_pred_rf=rf.predict(test_x)

/usr/local/lib/python3.7/dist-packages/sklearn/base.py:444: UserWarning: X has feature names, but RandomForestClassifier was fitted
f"X has feature names, but {self.__class__.__name__} was fitted without"

print("Training",accuracy_score(train_pred_rf,train_y))
print("Testing",accuracy_score(test_pred_rf,test_y))

Training 1.0
Testing 0.6866666666666666

```

```
from sklearn.ensemble import VotingClassifier
```

```
p=Perceptron()
```

```
svm=SVC()
```

```
knn=KNeighborsClassifier()
```

```
vc=VotingClassifier(estimators=[('perceptron',p),('svm',svm),('knn',knn)],voting='hard', weights=[3,1,1])
```

```
vc.fit(train_x_sc,train_y)
```

```
train_pred_vc=vc.predict(train_x_sc)
```

```
test_pred_vc=vc.predict(test_x)
```

```
/usr/local/lib/python3.7/dist-packages/sklearn/base.py:444: UserWarning: X has feature names, but Perceptron was fitted without fea
f"X has feature names, but {self.__class__.__name__} was fitted without"
/usr/local/lib/python3.7/dist-packages/sklearn/base.py:444: UserWarning: X has feature names, but SVC was fitted without feature na
f"X has feature names, but {self.__class__.__name__} was fitted without"
/usr/local/lib/python3.7/dist-packages/sklearn/base.py:444: UserWarning: X has feature names, but KNeighborsClassifier was fitted w
f"X has feature names, but {self.__class__.__name__} was fitted without"
```

```
print("Training",accuracy_score(train_pred_vc,train_y))
```

```
print("Testing",accuracy_score(test_pred_vc,test_y))
```

```
Training 0.7449856733524355
```

```
Testing 0.6866666666666666
```

```
from sklearn.ensemble import BaggingClassifier
```

```
bag=BaggingClassifier(base_estimator=knn,n_estimators=20)
```

```
bag.fit(train_x_sc,train_y)
```

```
train_pred_bag=bag.predict(train_x_sc)
```

```
test_pred_bag=bag.predict(test_x)
```

```
/usr/local/lib/python3.7/dist-packages/sklearn/base.py:444: UserWarning: X has feature names, but BaggingClassifier was fitted with
f"X has feature names, but {self.__class__.__name__} was fitted without"
```

```
print("Training",accuracy_score(train_pred_bag,train_y))
```

```
print("Testing",accuracy_score(test_pred_bag,test_y))
```

```
Training 0.9140401146131805
```

```
Testing 0.6866666666666666
```

```
cols=df.columns
```

```
cols
```

```
Index(['Gender', 'AGE', 'HbA1c', 'HDL', 'LDL', 'BMI', 'CLASS'], dtype='object')
```

```
sns.pairplot(df)
```

```
plt.show()
```

