```
import numpy as np
import pandas as pd
test1 = pd.read csv("C:\\Users\\HP\\Desktop\\phy\\health care
diabetes.csv")
test1.head()
   Pregnancies Glucose BloodPressure SkinThickness
                                                         Insulin
BMI \
             6
                    148
                                     72
                                                     35
                                                                  33.6
1
             1
                     85
                                     66
                                                     29
                                                               0
                                                                   26.6
2
             8
                    183
                                     64
                                                      0
                                                               0
                                                                  23.3
3
             1
                     89
                                     66
                                                     23
                                                              94 28.1
4
             0
                    137
                                     40
                                                     35
                                                             168 43.1
   DiabetesPedigreeFunction
                                   Outcome
                              Age
                       0.627
0
                               50
                                         1
                       0.351
                                         0
1
                               31
2
                       0.672
                               32
                                         1
3
                       0.167
                                         0
                               21
                       2.288
                               33
                                         1
df = test1.replace(0,'NAN')
df.head()
  Pregnancies Glucose BloodPressure SkinThickness Insulin
                                                              BMI \
0
            6
                  148
                                  72
                                                             33.6
                                                 35
                                                        NAN
1
            1
                   85
                                  66
                                                 29
                                                        NAN
                                                             26.6
2
            8
                  183
                                  64
                                                NAN
                                                        NAN
                                                             23.3
3
            1
                   89
                                  66
                                                 23
                                                         94
                                                             28.1
4
          NAN
                  137
                                                        168 43.1
                                  40
                                                 35
   DiabetesPedigreeFunction
                              Age Outcome
0
                       0.627
                               50
1
                       0.351
                               31
                                      NAN
2
                       0.672
                               32
                                        1
3
                       0.167
                               21
                                      NAN
                       2.288
                               33
                                        1
df1 = df.replace('NAN',np.nan)
df1.head()
   Pregnancies Glucose BloodPressure SkinThickness
                                                         Insulin
BMI \
0
           6.0
                  148.0
                                   72.0
                                                   35.0
                                                             NaN 33.6
```

1	1.0	85.0		66.0	29.0	NaN	26.6
2	8.0	183.0		64.0	NaN	NaN	23.3
3	1.0	89.0		66.0	23.0	94.0	28.1
4	NaN	137.0		40.0	35.0	168.0	43.1
Dia 0 1 2 3 4	abetesPedig	reeFunction 0.627 0.351 0.672 0.167 2.288	50 31 32 21	Outcome 1.0 NaN 1.0 NaN	9 N 9 N		
df2= d df2.he	df1.interpo ead()	late()					
Pre BMI \	-	Glucose Bl	oodPres	ssure S	SkinThickness	Insulin	
0	6.0	148.0		72.0	35.0	NaN	33.6
1	1.0	85.0		66.0	29.0	NaN	26.6
2	8.0	183.0		64.0	26.0	NaN	23.3
3	1.0	89.0		66.0	23.0	94.0	28.1
4	3.0	137.0		40.0	35.0	168.0	43.1
DiabetesPedigreeFunction Age Outcome 0 0.627 50 1.0 1 0.351 31 1.0 2 0.672 32 1.0 3 0.167 21 1.0 4 2.288 33 1.0							
<pre>x = df2['Insulin'].mean() df2['Insulin'].fillna(x,inplace = True) print(df2)</pre>							
F BMI \	Pregnancies	Glucose	BloodPr	ressure	SkinThicknes	s Ins	ulin
0 33.6	6.0	148.0		72.0	35.	0 159.04	5098
1 26.6	1.0	85.0		66.0	29.	0 159.04	5098
2	8.0	183.0		64.0	26.	0 159.04	5098

23.3						
3	1.0	89.0		66.0	23.0	94.000000
28.1						
4	3.0	137.0		40.0	35.0	168.000000
43.1						
• •						
762	10.0	101 0		76 0	40.0	100 000000
763 32.9	10.0	101.0		76.0	48.0	180.000000
764	2.0	122.0		70.0	27.0	146.000000
36.8	2.0	122.0		70.0	27.0	140.000000
765	5.0	121.0		72.0	23.0	112.000000
26.2				7 = 1 0		
766	1.0	126.0		60.0	27.0	112.000000
30.1						
767	1.0	93.0		70.0	31.0	112.000000
30.4						
	DiabetesPedi	.greeFunction	Age	Outcome		
0		0.627	50	1.0		
1		0.351	31	1.0		

	 	J -	
0	0.627	50	1.0
1	0.351	31	1.0
2	0.672	32	1.0
3	0.167	21	1.0
4	2.288	33	1.0
763	0.171	63	1.0
764	0.340	27	1.0
765	0.245	30	1.0
766	0.349	47	1.0
767	0.315	23	1.0

[768 rows x 9 columns]

df2.head()

	regnancies	Glucose	BloodPressure	SkinThickness	Insulin
BMI 0	6.0	148.0	72.0	35.0	159.045098
33.6	1.0	85.0	66.0	29.0	159.045098
26.6	8.0	183.0	64.0	26.0	159.045098
23.3	1.0	89.0	66.0	23.0	94.000000
28.1 4 43.1	3.0	137.0	40.0	35.0	168.000000

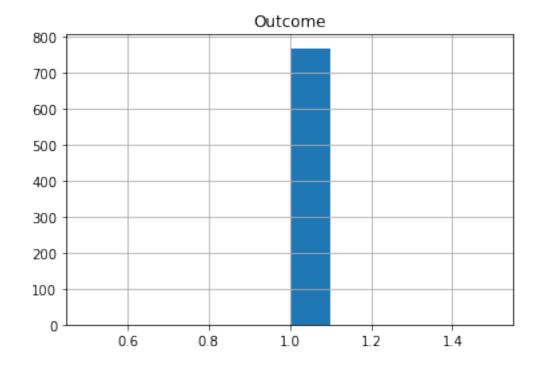
DiabetesPedigreeFunction Age Outcome 0 0.627 50 1.0

```
0.351
1
                                  31
                                           1.0
2
                         0.672
                                  32
                                           1.0
3
                         0.167
                                  21
                                           1.0
4
                         2.288
                                  33
                                           1.0
```

import matplotlib.pyplot as plt
from matplotlib import style
%matplotlib inline

df2.hist(column = 'Outcome')

array([[<AxesSubplot:title={'center':'Outcome'}>]], dtype=object)

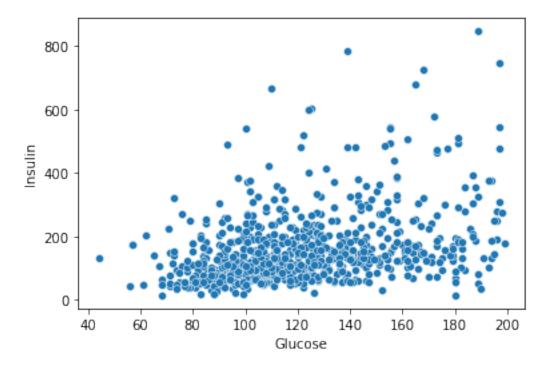


import seaborn as sns

sns.scatterplot(df2.Glucose, df2.Insulin);

C:\ANACONDA\lib\site-packages\seaborn\\_decorators.py:36: FutureWarning: Pass the following variables as keyword args: x, y. From version 0.12, the only valid positional argument will be `data`, and passing other arguments without an explicit keyword will result in an error or misinterpretation.

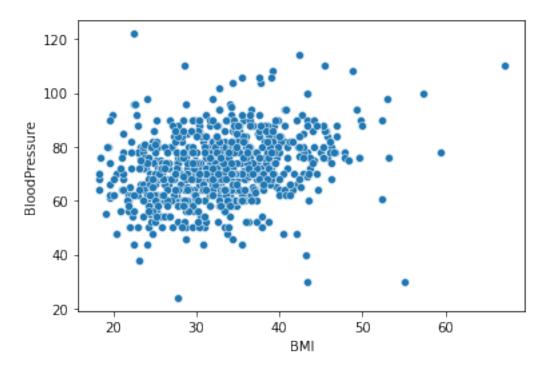
warnings.warn(



sns.scatterplot(df2.BMI, df2.BloodPressure);

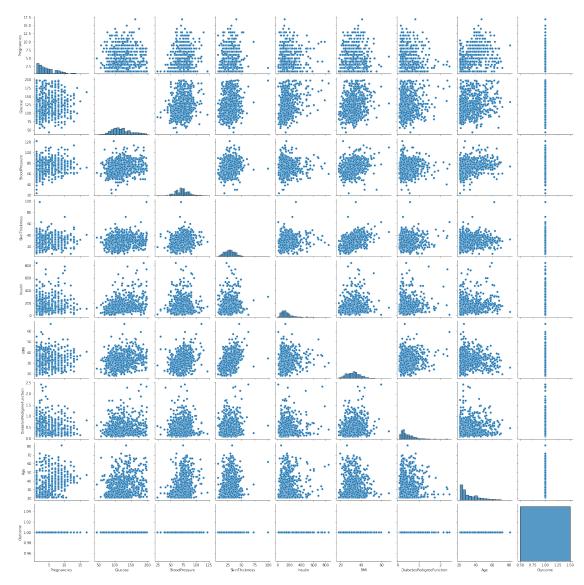
C:\ANACONDA\lib\site-packages\seaborn\\_decorators.py:36: FutureWarning: Pass the following variables as keyword args: x, y. From version 0.12, the only valid positional argument will be `data`, and passing other arguments without an explicit keyword will result in an error or misinterpretation.

warnings.warn(



## sns.pairplot(df2)

## <seaborn.axisgrid.PairGrid at 0x19db391b340>



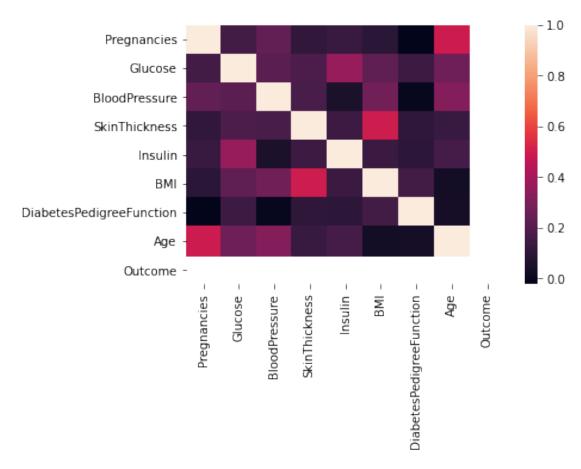
## ##CORRELATION ANALYSIS

SkinThickness \	Pregnancie	es Glucose	e BloodPressure
Pregnancies	1.00000	0.158436	0.233739
0.110458 Glucose	0.15843	30 1.000000	0.216927
0.183719 BloodPressure	0.23373	39 0.216927	1.00000
0.175081 SkinThickness	0.11045	8 0.183719	0.175081
1.000000 Insulin	0.13049	0.362336	0.050028
0.139029 BMI	0.09029	06 0.231684	0.273978
0.494883 DiabetesPedigreeFunction	-0.02381	19 0.138814	-0.005157
0.103014 Age	0.49139	99 0.268057	0.318358
0.130788 Outcome	Na	aN NaN	I NaN
NaN			
\	Insulin	BMI D	DiabetesPedigreeFunction
Pregnancies	0.130491	0.090296	-0.023819
Glucose	0.362336	0.231684	0.138814
BloodPressure	0.050028	0.273978	-0.005157
SkinThickness	0.139029	0.494883	0.103014
Insulin	1.000000	0.136063	0.100111
BMI	0.136063	1.000000	0.157147
DiabetesPedigreeFunction	0.100111	0.157147	1.000000
Age	0.161294	0.025975	0.033561
Outcome	NaN	NaN	NaN
Pregnancies Glucose BloodPressure SkinThickness Insulin BMI	Age 0.491399 0.268057 0.318358 0.130788 0.161294 0.025975	Outcome NaN NaN NaN NaN NaN	

DiabetesPedigreeFunction 0.033561 NaN Age 1.000000 NaN Outcome NaN NaN

sns.heatmap(df2.corr())

## <AxesSubplot:>



train = pd.read\_csv("C:\\Users\\HP\\Desktop\\phy\\train.csv")
train.head()

Age Ge	nder A	AppointmentRegistration	ApointmentData
DayOfTheW	eek \	\	
0 38	F	2015-10-20T08:33:56Z	2015-10-23T00:00:00Z
Friday			
1 56	F	2014-02-03T10:05:26Z	2014-02-20T00:00:00Z
Thursday			
2 27	F	2014-04-29T07:57:32Z	2014-05-20T00:00:00Z
Tuesday			
3 24	М	2014-04-02T13:53:37Z	2014-05-06T00:00:00Z
Tuesday			
4 48	F	2014-01-07T10:07:17Z	2014-01-30T00:00:00Z
Thursday			
That Saay			

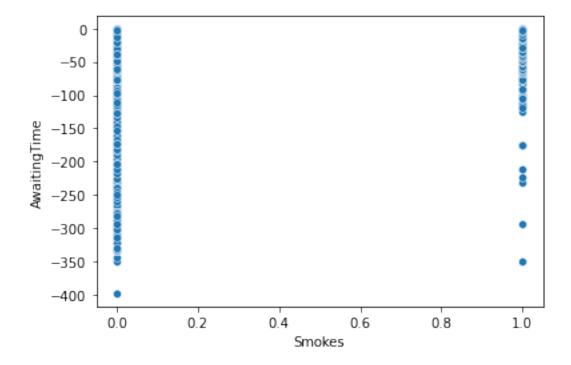
Sc	Status holarship		Alcoolism	HiperTension	Handcap	Smokes
0	No-Show	0	0	0	0	1
1 0	No-Show	1	0	1	Θ	0
2	Show-Up	0	0	Θ	Θ	0
3	Show-Up	Θ	0	0	0	0
4	Show-Up	0	0	Θ	0	0
0 1 2 3 4	Tubercul	osis Sms_ 0 0 0 0 0	Reminder 0 1 0 0 1 0 1	AwaitingTime -3 -17 -21 -34 -23		
ap	<pre>test = pd.read_csv("C:\\Users\\HP\\Desktop\\phy\\healthcare appointment data.csv") test.head()</pre>					
Da	Age Gendo		mentRegist	ration /	Apointment	Data
0	19 dnesday		-12-16T14:	46:25Z 2015-01	1-14T00:00	:00Z
1	24 dnesday	F 2015	6-08-18T07:	91:26Z 2015-08	B-19T00:00	:00Z
2	4 esday	F 2014	-02-17T12:	53:46Z 2014-02	2-18T00:00	:00Z
3	5 ursday	M 2014	-07-23T17:	92:11Z 2014-08	3-07T00:00	:00Z
4	38 esday	M 2015	5-10-21T15:	20:09Z 2015-10	9-27T00:00	:00Z
Sc	Status holarship	Diabetes	Alcoolism	HiperTension	Handcap	Smokes
0 0	Show-Up	0	Θ	Θ	Θ	0
1	Show-Up	0	0	Θ	Θ	0
0 2	Show-Up	0	0	0	0	0
0	Show-Up	0	0	0	0	0
0 4 0	Show-Up	0	0	0	0	0

Tub 0 1 2 3 4	erculosis Sms_ 0 0 0 0 0	Reminder Await 0 0 0 1 1	ingTime -29 -1 -1 -15 -6		
	describe()	1	-0		
count mean std min 25% 50% 75% max	Age 210000.000000 37.761824 22.794334 -1.000000 19.000000 38.000000 56.000000 113.000000	Diabetes 210000.000000 0.077290 0.267052 0.000000 0.000000 0.000000 1.000000	Alcoolism 210000.000000 0.024676 0.155137 0.000000 0.000000 0.000000 0.000000 1.000000	HiperTension 210000.000000 0.214862 0.410727 0.000000 0.000000 0.000000 0.000000 1.000000	\
count mean std min 25% 50% 75% max	Handcap 210000.000000 0.020471 0.155854 0.000000 0.000000 0.000000 4.000000	Smokes 210000.000000 0.052033 0.222095 0.000000 0.000000 0.000000 1.000000	Scholarship 210000.000000 0.097738 0.296961 0.000000 0.000000 0.000000 1.000000	Tuberculosis 210000.000000 0.000429 0.020698 0.000000 0.000000 0.000000 1.000000	\
count mean std min 25% 50% 75% max	Sms_Reminder 210000.000000 0.574238 0.499776 0.000000 1.000000 1.000000 2.000000	AwaitingTime 210000.000000 -13.833538 15.685272 -398.000000 -20.000000 -8.000000 -4.000000 -1.000000			
test.d	escribe()				
count mean std min 25% 50% 75% max	Age 300000.000000 37.808017 22.809014 -2.000000 19.000000 38.000000 56.000000 113.000000	Diabetes 300000.000000 0.077967 0.268120 0.000000 0.000000 0.000000 0.000000	Alcoolism 300000.000000 0.025010 0.156156 0.000000 0.000000 0.000000 0.000000	HiperTension 300000.0000000 0.215890 0.411439 0.000000 0.000000 0.000000 0.000000 1.000000	\

```
Smokes
                                         Scholarship
                                                        Tuberculosis
             Handcap
                                                                       \
       300000.000000
                       300000.000000
                                       300000.000000
                                                       300000.000000
count
                                            0.096897
mean
            0.020523
                            0.052370
                                                            0.000450
std
            0.155934
                            0.222772
                                            0.295818
                                                            0.021208
min
            0.000000
                            0.000000
                                            0.000000
                                                            0.000000
25%
            0.000000
                            0.000000
                                            0.000000
                                                            0.000000
50%
            0.000000
                            0.000000
                                            0.000000
                                                            0.000000
75%
            0.000000
                            0.000000
                                            0.000000
                                                            0.000000
            4.000000
                            1.000000
                                            1.000000
                                                            1.000000
max
        Sms Reminder
                        AwaitingTime
       3000\overline{0}0.000000
                       300000.000000
count
            0.574173
                          -13.841813
mean
std
            0.499826
                           15.687697
min
            0.000000
                         -398.000000
                          -20,000000
25%
            0.000000
50%
            1.000000
                           -8.000000
75%
            1.000000
                           -4.000000
            2.000000
                           -1.000000
max
train.shape
(210000, 15)
test.shape
(300000, 15)
sns.scatterplot(train.Smokes,train.AwaitingTime)
C:\ANACONDA\lib\site-packages\seaborn\ decorators.py:36:
FutureWarning: Pass the following variables as keyword args: x, y.
From version 0.12, the only valid positional argument will be `data`
and passing other arguments without an explicit keyword will result in
an error or misinterpretation.
```

<AxesSubplot:xlabel='Smokes', ylabel='AwaitingTime'>

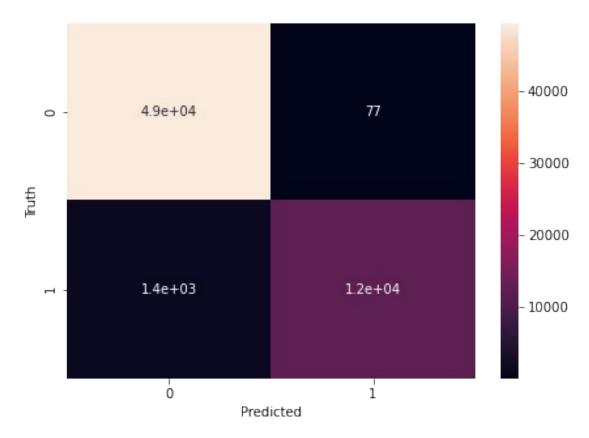
warnings.warn(



from sklearn.model\_selection import train\_test\_split

```
train.drop(['Gender','AppointmentRegistration','ApointmentData','DayOf
TheWeek','Status'], axis = 'columns')
y = train.HiperTension
x_train,x_test,y_train,y_test =
train_test_split(x,y,random_state=42,test size=0.3)
x train.shape,x test.shape,y train.shape,y test.shape
((147000, 10), (63000, 10), (147000,), (63000,))
##KNN CLASSIFIER
from sklearn.neighbors import KNeighborsClassifier
knn = KNeighborsClassifier(n neighbors = 10)
knn.fit(x_train, y_train)
KNeighborsClassifier(n neighbors=10)
knn.score(x_test,y_test)
0.9763333333333334
from sklearn.metrics import confusion_matrix
y pred = knn.predict(x test)
cm = confusion_matrix(y_test, y_pred)
\mathsf{cm}
```

Text(42.0, 0.5, 'Truth')



from sklearn.metrics import classification\_report
print(classification\_report(y\_test, y\_pred))

	precision	recall	fl-score	support
0 1	0.97 0.99	1.00 0.90	0.99 0.94	49443 13557
accuracy macro avg weighted avg	0.98 0.98	0.95 0.98	0.98 0.96 0.98	63000 63000 63000