ad0v1h4qd

March 18, 2024

[1]:
```
import numpy as np
import pandas as pd

#import visualization libraries
import matplotlib.pyplot as plt
import seaborn as sns

#import sklearn libraries
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import MinMaxScaler
from sklearn.preprocessing import StandardScaler
from sklearn.linear_model import LogisticRegression
```

[2]:
```
## Remove warnings
import warnings
warnings.filterwarnings('ignore')
```

[3]:
```
# Load the data sets

df = pd.read_csv("diabetes.csv")
pd.set_option('display.max_columns',None)
df.head()
```

[3]:

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	DiabetesPedigreeFunction	Age	Outcome
0	6	148	72	35	0	33.6	0.627	50	1
1	1	85	66	29	0	26.6	0.351	31	0
2	8	183	64	0	0	23.3	0.672	32	1
3	1	89	66	23	94	28.1	0.167	21	0
4	0	137	40	35	168	43.1	2.288	33	1

[4]:
```
df.tail()
```

[4]:

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	DiabetesPedigreeFunction	Age	Outcome
763	10	101	76	48	180	32.9	0.171	63	0
764	2	122	70	27	0	36.8	0.340	27	0
765	5	121	72	23	112	26.2	0.245	30	0
766	1	126	60	0	0	30.1	0.349	47	1
767	1	93	70	31	0	30.4	0.315	23	0

[5]:
```
# checking shape of dataset
df.shape
```

[5]:
```
(768, 9)
```

[6]:
```
df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 768 entries, 0 to 767
Data columns (total 9 columns):
 #   Column                    Non-Null Count  Dtype  
---  ------                    --------------  -----  
 0   Pregnancies               768 non-null    int64  
 1   Glucose                   768 non-null    int64  
 2   BloodPressure             768 non-null    int64  
 3   SkinThickness             768 non-null    int64  
 4   Insulin                   768 non-null    int64  
 5   BMI                       768 non-null    float64
 6   DiabetesPedigreeFunction  768 non-null    float64
 7   Age                       768 non-null    int64  
 8   Outcome                   768 non-null    int64  
dtypes: float64(2), int64(7)
memory usage: 54.1 KB
```

[7]:
```
# checking missing values

df.isnull().mean()*100
```

[7]:
```
Pregnancies                 0.0
Glucose                     0.0
BloodPressure               0.0
```

```
SkinThickness               0.0
Insulin                     0.0
BMI                         0.0
DiabetesPedigreeFunction    0.0
Age                         0.0
Outcome                     0.0
dtype: float64
```

[8]:
```
df.columns
```

[8]:
```
Index(['Pregnancies', 'Glucose', 'BloodPressure', 'SkinThickness', 'Insulin',
       'BMI', 'DiabetesPedigreeFunction', 'Age', 'Outcome'],
      dtype='object')
```

[9]:
```
df.describe()  # checking summary of dataset
```

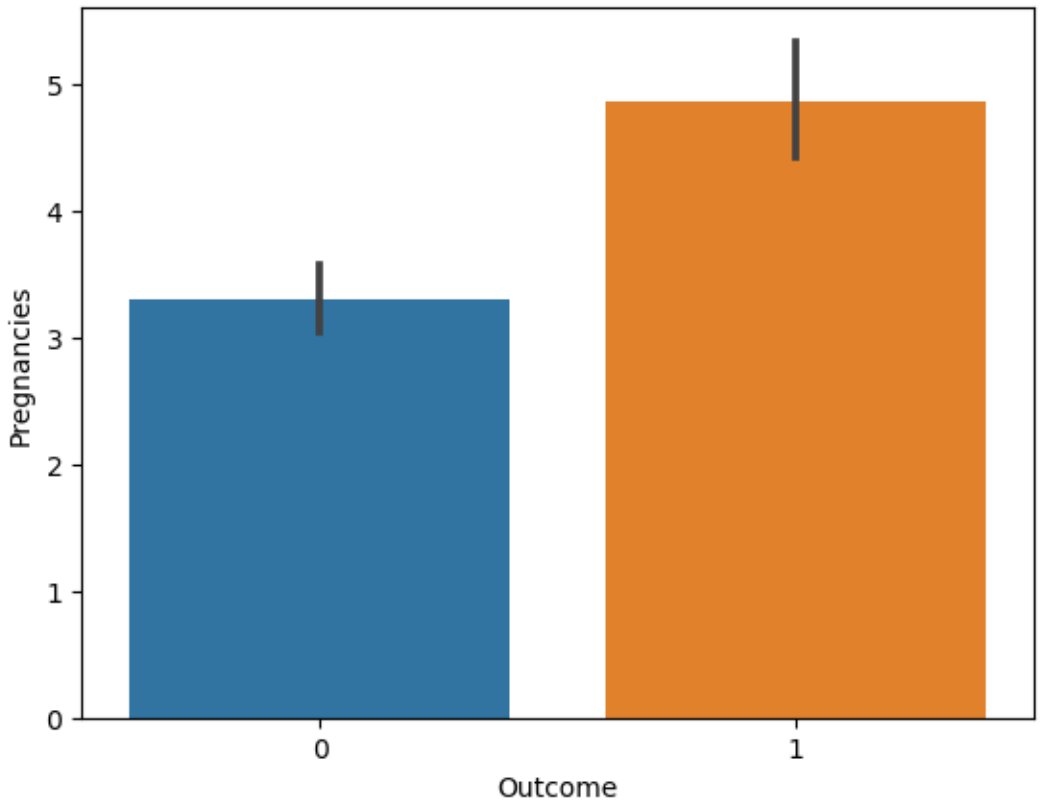
[9]:
```
       Pregnancies     Glucose  BloodPressure  SkinThickness     Insulin  \
count   768.000000  768.000000     768.000000     768.000000  768.000000
mean      3.845052  120.894531      69.105469      20.536458   79.799479
std       3.369578   31.972618      19.355807      15.952218  115.244002
min       0.000000    0.000000       0.000000       0.000000    0.000000
25%       1.000000   99.000000      62.000000       0.000000    0.000000
50%       3.000000  117.000000      72.000000      23.000000   30.500000
75%       6.000000  140.250000      80.000000      32.000000  127.250000
max      17.000000  199.000000     122.000000      99.000000  846.000000

              BMI  DiabetesPedigreeFunction         Age     Outcome
count  768.000000                768.000000  768.000000  768.000000
mean    31.992578                  0.471876   33.240885    0.348958
std      7.884160                  0.331329   11.760232    0.476951
min      0.000000                  0.078000   21.000000    0.000000
25%     27.300000                  0.243750   24.000000    0.000000
50%     32.000000                  0.372500   29.000000    0.000000
75%     36.600000                  0.626250   41.000000    1.000000
max     67.100000                  2.420000   81.000000    1.000000
```

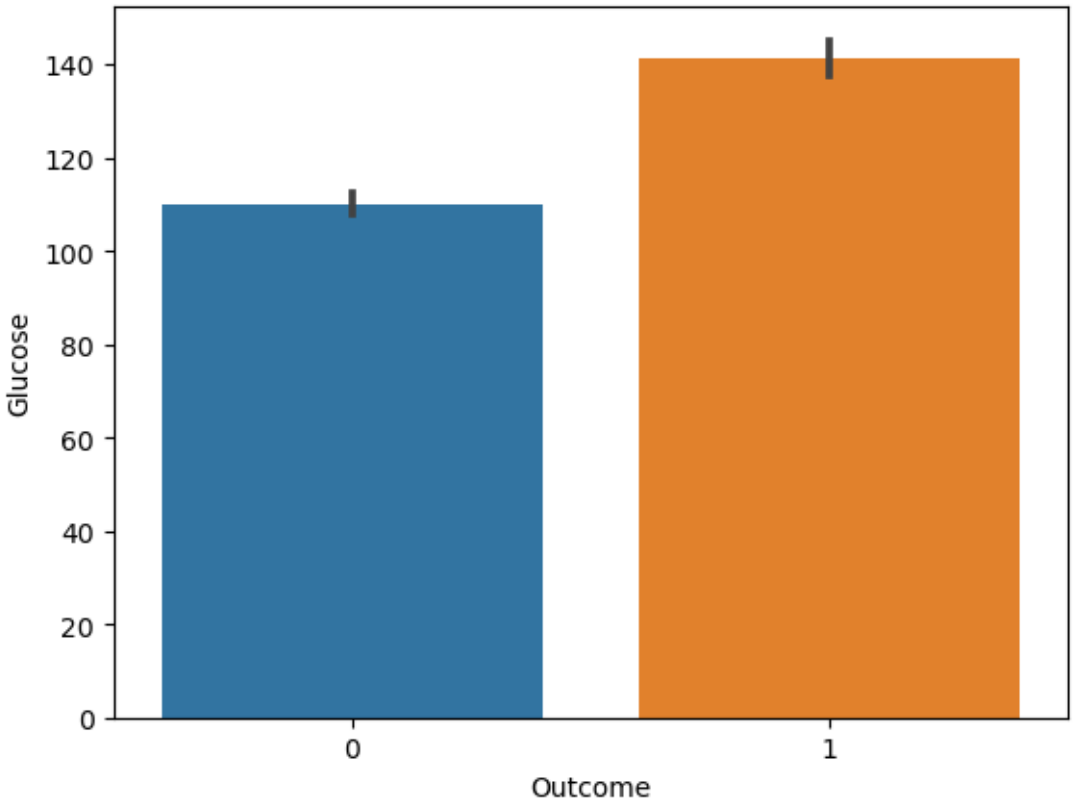
[10]:
```
# calculating probability
df['Outcome'].value_counts()/len(df)*100
```

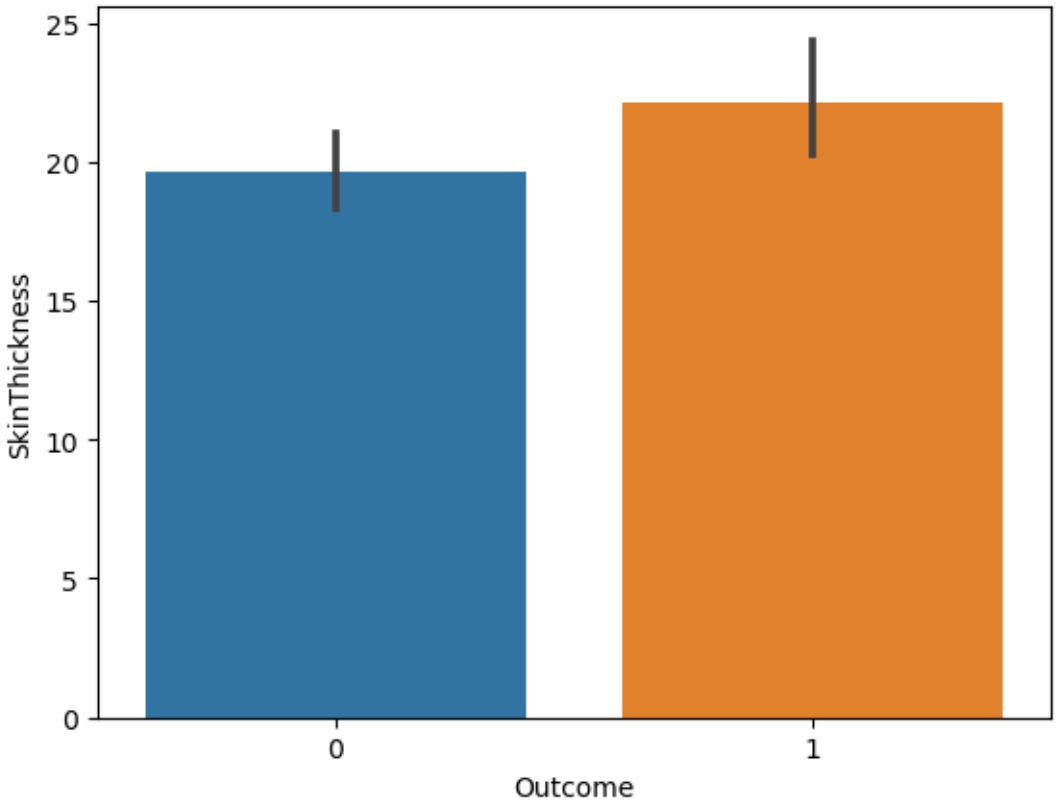
[10]:
```
Outcome
0    65.104167
1    34.895833
Name: count, dtype: float64
```

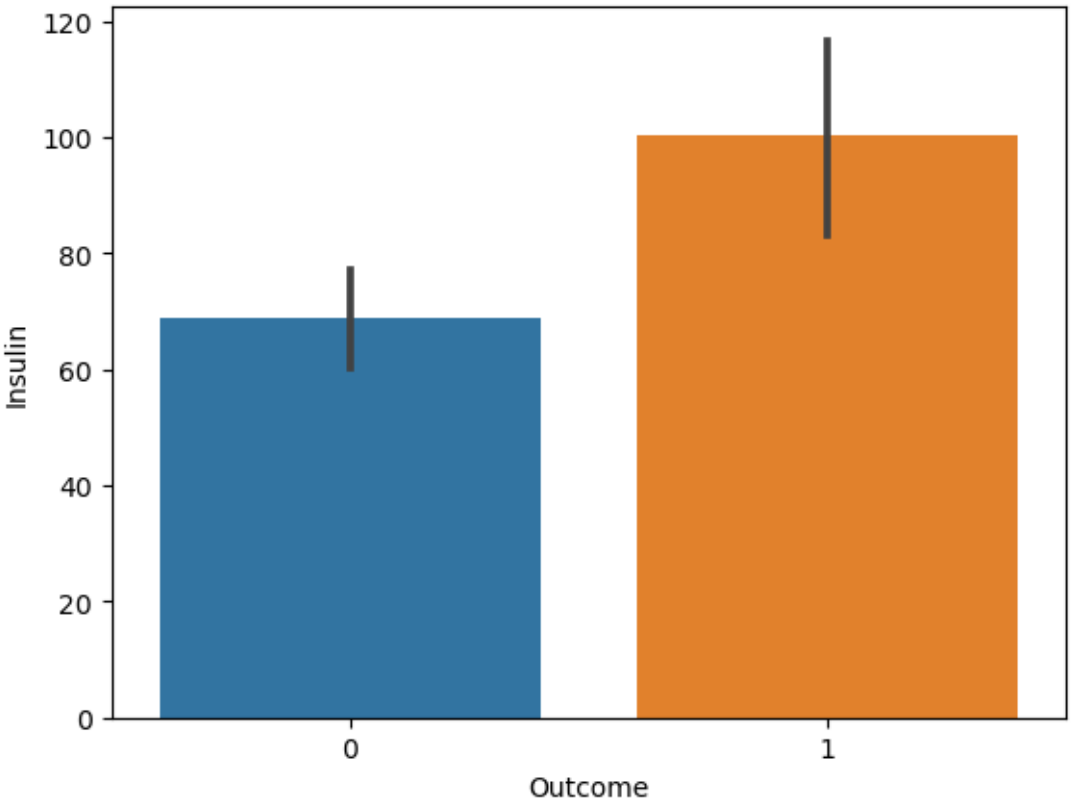
[11]:
```
# EDA
plt.figure()
sns.barplot(x='Outcome',y='Pregnancies',data=df)
```

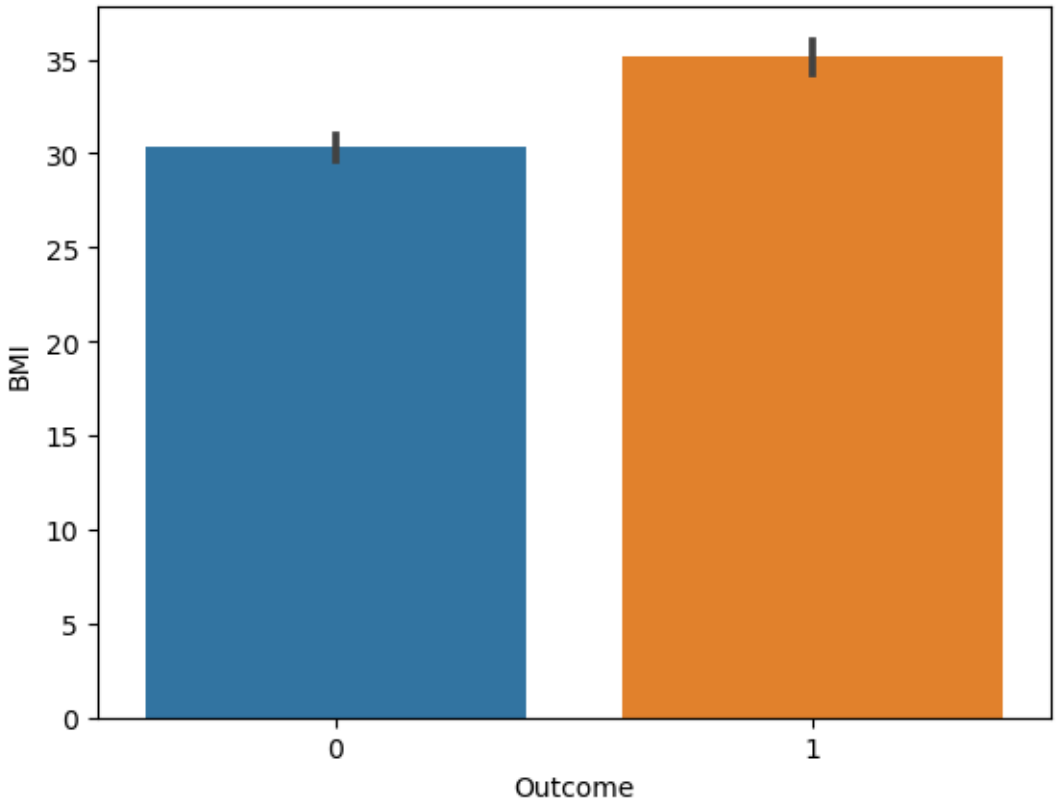
```
plt.show()
plt.figure()
sns.barplot(x='Outcome' , y='Glucose' , data=df)
plt.show()
plt.figure()
sns.barplot(x='Outcome', y='SkinThickness' , data=df)
plt.show()
plt.figure()
sns.barplot(x='Outcome', y='Insulin' , data=df)
plt.show()
plt.figure()
sns.barplot(x='Outcome', y='BMI' , data=df)
plt.show()
plt.figure()
sns.barplot(x='Outcome', y='DiabetesPedigreeFunction' , data=df)
plt.figure()
sns.barplot(x='Outcome', y='Age' , data=df)
plt.show()
```

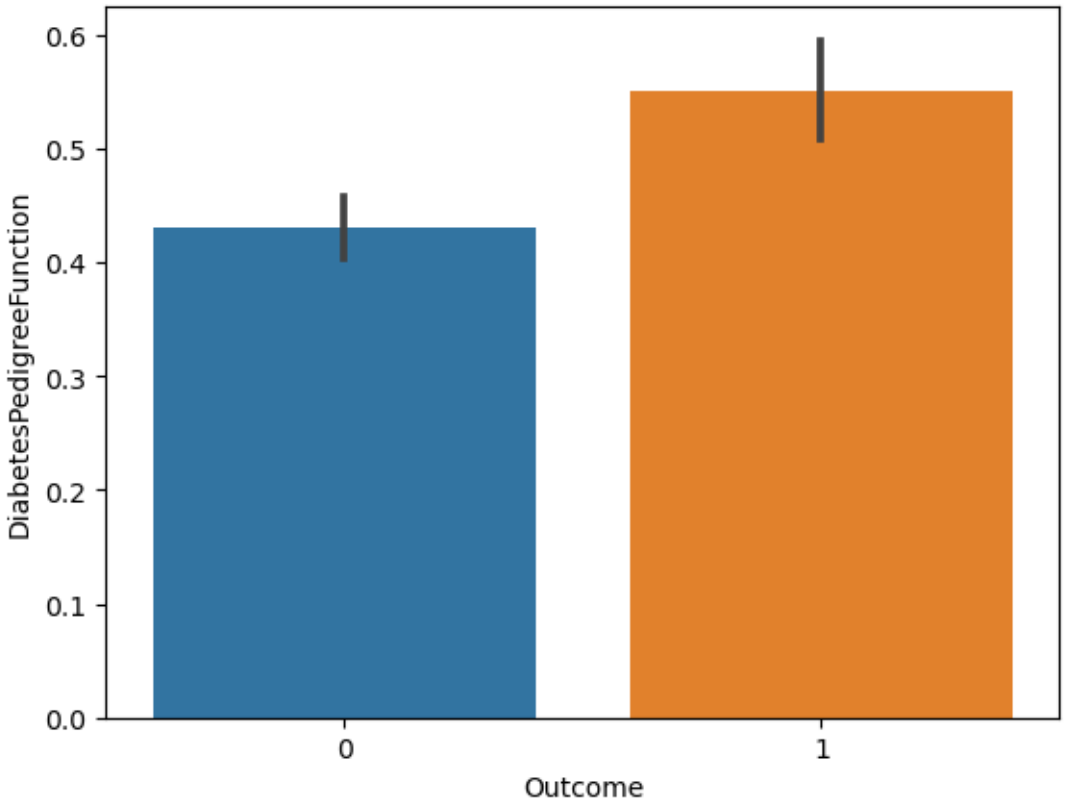


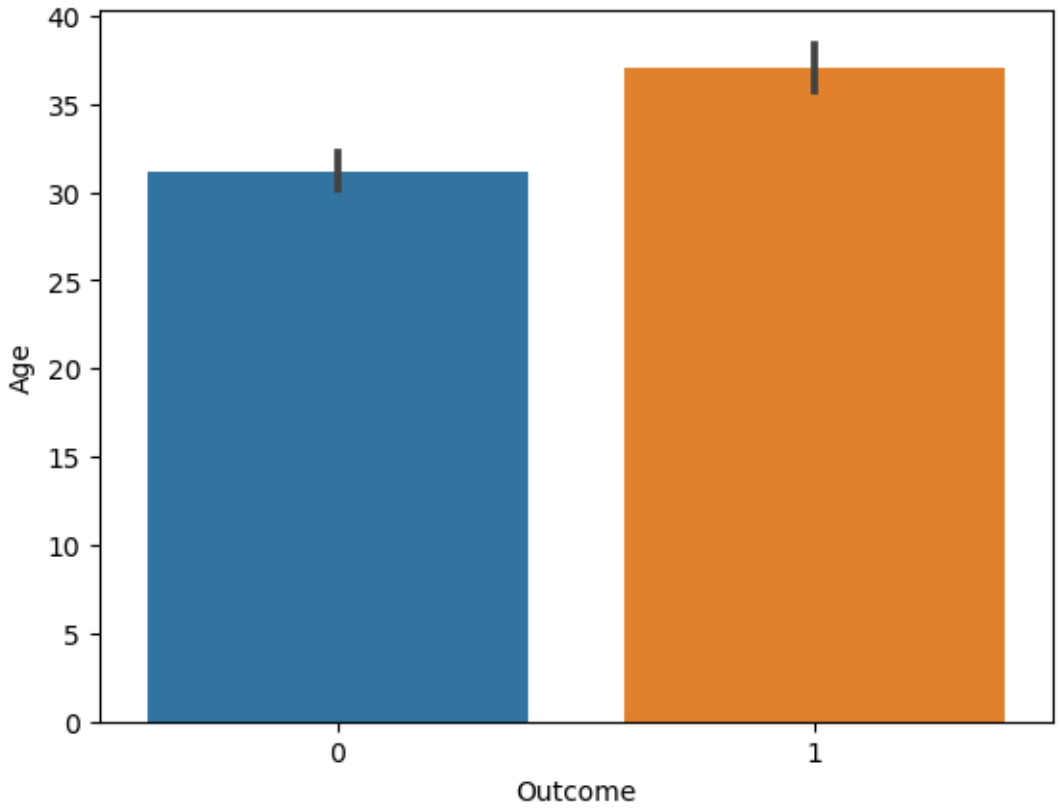

140
120
100
80
60
40
20
0
Glucose
0
1
Outcome


25
20
15
10
5
0
SkinThickness
0
1
Outcome

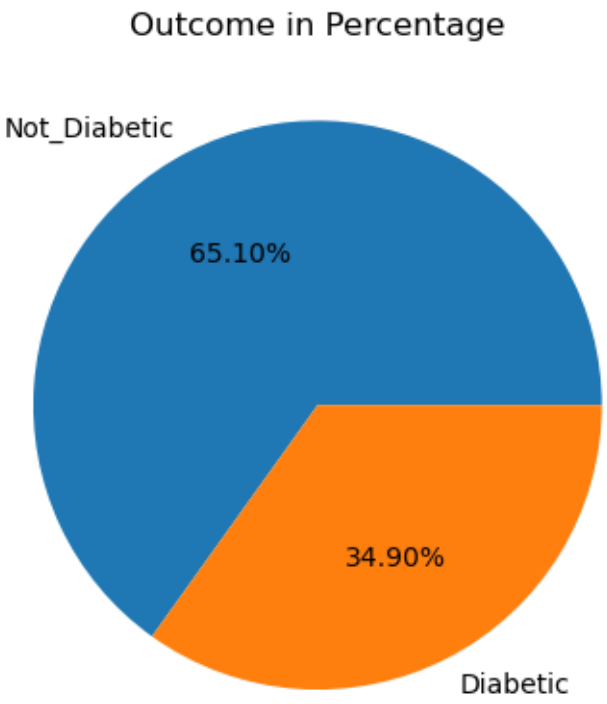

120
100
80
60
40
20
0
Insulin
0
1
Outcome


35
30
25
20
15
10
5
0
BMI
0
1
Outcome


0.6
0.5
0.4
0.3
0.2
0.1
0.0
DiabetesPedigreeFunction
0
1
Outcome

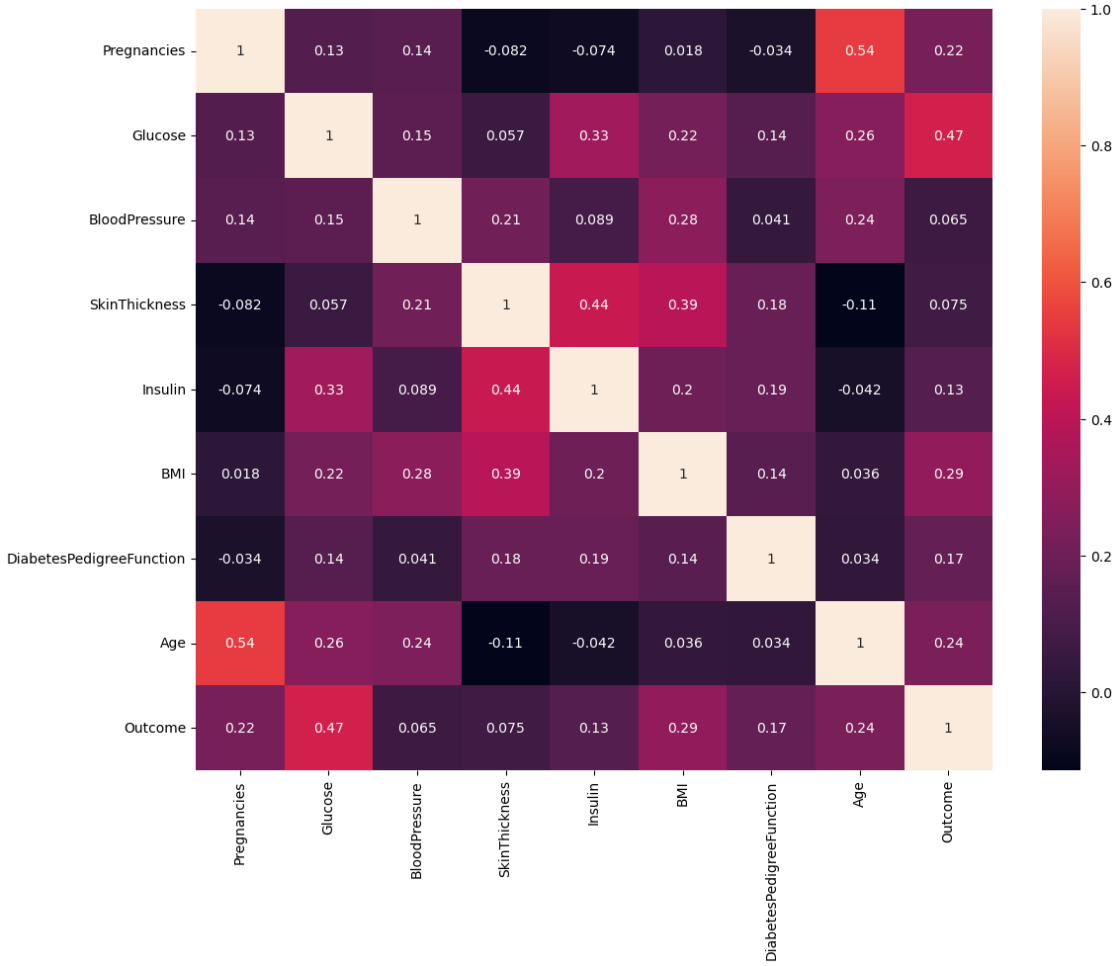


[12]:
```
plt.title('Outcome in Percentage')
plt.pie(df['Outcome'].
  ↪value_counts(),labels=['Not_Diabetic','Diabetic'],autopct='%1.2f%%')
plt.show()
```



[13]:
```
plt.figure(figsize=(13,10))
sns.heatmap(df.corr(),annot=True)
```

[13]: <Axes: >



[14]:
```
# Data preprocessing
 # Replacing Nan with mean values
df["Glucose"].fillna(df["Glucose"].mean(), inplace = True)
df["BloodPressure"].fillna(df["BloodPressure"].mean(), inplace = True)
df["SkinThickness"].fillna(df["SkinThickness"].mean(), inplace = True)
df["Insulin"].fillna(df["Insulin"].mean(), inplace = True)
df["BMI"].fillna(df["BMI"].mean(), inplace = True)
df["Pregnancies"].fillna(df["Pregnancies"].mean(), inplace = True)
df["DiabetesPedigreeFunction"].fillna(df["DiabetesPedigreeFunction"].mean(),␣
 ↪inplace = True)
```

[15]:
```
df.describe().T
```

[15]:

	count	mean	std	min	25%	\
Pregnancies	768.0	3.845052	3.369578	0.000	1.00000	
Glucose	768.0	120.894531	31.972618	0.000	99.00000	
BloodPressure	768.0	69.105469	19.355807	0.000	62.00000	

```
SkinThickness              768.0   20.536458   15.952218   0.000   0.00000
Insulin                    768.0   79.799479  115.244002   0.000   0.00000
BMI                        768.0   31.992578    7.884160   0.000  27.30000
DiabetesPedigreeFunction   768.0    0.471876    0.331329   0.078   0.24375
Age                        768.0   33.240885   11.760232  21.000  24.00000
Outcome                    768.0    0.348958    0.476951   0.000   0.00000

                             50%        75%     max
Pregnancies                3.0000    6.00000   17.00
Glucose                  117.0000  140.25000  199.00
BloodPressure             72.0000   80.00000  122.00
SkinThickness             23.0000   32.00000   99.00
Insulin                   30.5000  127.25000  846.00
BMI                       32.0000   36.60000   67.10
DiabetesPedigreeFunction   0.3725    0.62625    2.42
Age                       29.0000   41.00000   81.00
Outcome                    0.0000    1.00000    1.00
```

0.0.1 Prediction using Logistic Regression

[16]:
```
#Splitting Data
X = df.drop(columns='Outcome',axis=1)
Y = df['Outcome']
```

[17]:
```
#Splitting the data
X_train,X_test,Y_train,Y_test = train_test_split(X,Y,test_size=0.
 ↪2,stratify=Y,random_state=2)
print(X.shape)
print(X_train.shape)
print(X_test.shape)
```

```
(768, 8)
(614, 8)
(154, 8)
```

[18]:
```
model = LogisticRegression()
model.fit(X_train, Y_train)
y_predict = model.predict(X_test)
```

[19]:
```
y_predict
```

[19]:
```
array([0, 0, 1, 1, 0, 0, 0, 1, 0, 1, 1, 0, 0, 1, 0, 1, 0, 0, 0, 0, 0, 0,
       1, 0, 0, 0, 0, 1, 0, 1, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 1, 0, 0, 1,
       0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0,
       0, 1, 1, 0, 1, 0, 1, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0,
       1, 0, 0, 0, 1, 0, 0, 1, 0, 0, 1, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
       1, 0, 0, 0, 0, 1, 0, 1, 0, 1, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 1,
```

```
 1, 1, 0, 0, 0, 0, 0, 1, 0, 1, 0, 0, 0, 1, 0, 0, 0, 0, 1, 0, 0, 1],
dtype=int64)
```

[20]:
```
#confusion matrix
from sklearn.metrics import confusion_matrix
confusion_matrix(y_predict,Y_test)
```

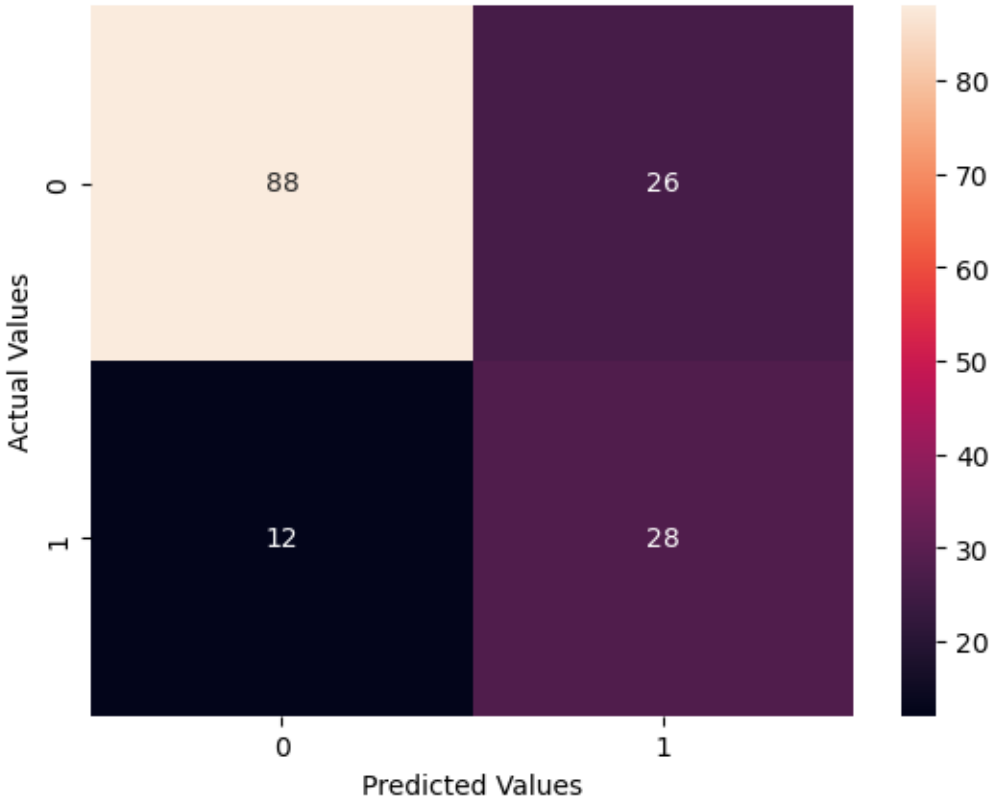
[20]:
```
array([[88, 26],
       [12, 28]], dtype=int64)
```

[21]:
```
from sklearn.metrics import accuracy_score ,classification_report,f1_score
```

[22]:
```
Accuracy=accuracy_score(Y_test,y_predict)
print("Accuracy Score of test data : ",Accuracy)
```

```
Accuracy Score of test data :  0.7532467532467533
```

[23]:
```
#Evaluating model
cf_matrix = confusion_matrix(y_predict,Y_test)
sns.heatmap(cf_matrix,annot=True,fmt=".0f")
plt.xlabel("Predicted Values")
plt.ylabel("Actual Values")
plt.show()
```



[24]:
```
print(classification_report(Y_test,y_predict))
```

```
              precision    recall  f1-score   support

           0       0.77      0.88      0.82       100
           1       0.70      0.52      0.60        54

    accuracy                           0.75       154
   macro avg       0.74      0.70      0.71       154
weighted avg       0.75      0.75      0.74       154
```

[25]:
```
#Model Testing

y_predict = model.predict([[4,130,65,30,90,33.6,0.627,50]])
print(y_predict)
if y_predict==1:
    print("You are Diabetic")
else:
    print("You are not Diabetic")
```

```
[1]
You are Diabetic
```

[26]:
```
#Model Testing

y_predict = model.predict([[4,110,40,20,150,30.6,0.527,30]])
print(y_predict)
if y_predict==1:
    print("You are Diabetic")
else:
    print("You are not Diabetic")
```

```
[0]
You are not Diabetic
```

0.0.2 Prediction Random Forest Classifier

[27]:
```
from sklearn.ensemble import RandomForestClassifier
rf = RandomForestClassifier()
```

[28]:
```
rf.fit(X_train,Y_train)
```

[28]:
```
RandomForestClassifier()
```

[29]:
```
Y_pred1 = rf.predict(X_test)
Y_pred1
```

[29]:
```
array([0, 0, 0, 1, 0, 0, 0, 1, 0, 1, 1, 0, 0, 1, 0, 1, 0, 0, 0, 0, 0, 0,
       1, 0, 0, 0, 0, 1, 1, 1, 1, 0, 0, 1, 0, 0, 0, 0, 0, 0, 1, 0, 0, 1,
       1, 0, 1, 0, 0, 0, 0, 0, 1, 0, 1, 1, 0, 0, 0, 0, 0, 0, 1, 0, 1, 0,
       0, 1, 1, 0, 1, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0,
       1, 0, 0, 0, 1, 0, 0, 1, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
       1, 0, 0, 0, 0, 1, 0, 1, 0, 1, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 1,
       1, 1, 0, 0, 0, 0, 0, 1, 0, 1, 0, 0, 0, 1, 0, 0, 0, 0, 1, 0, 0, 0],
      dtype=int64)
```

[30]:
```
confusion_matrix(Y_pred1,Y_test)
```

[30]:
```
array([[87, 25],
       [13, 29]], dtype=int64)
```

[31]:
```
accuracy=accuracy_score(Y_test,Y_pred1)
print("Accuracy Score of test data : ",accuracy)
```

```
Accuracy Score of test data :  0.7532467532467533
```

[32]:
```
print(classification_report(Y_test,Y_pred1))
```

```
              precision    recall  f1-score   support

           0       0.78      0.87      0.82       100
           1       0.69      0.54      0.60        54

    accuracy                           0.75       154
   macro avg       0.73      0.70      0.71       154
weighted avg       0.75      0.75      0.74       154
```

[33]:
```
predict1 = rf.predict([[4,110,40,20,150,30.6,0.527,30]])
print(predict1)
if predict1==1:
    print("You are Diabetic")
else:
    print("You are not Diabetic")
```

```
[0]
You are not Diabetic
```

Thank You