```
In [1]: import pandas as pd
         import numpy as np
         import matplotlib.pyplot as plt
         import seaborn as sns
         %matplotlib inline
         df=pd.read_csv('after_week1.csv')
         df.head()
In [2]: df=pd.read_csv('after_week1.csv')
         df.head()
Out[2]:
                                                            Insulin BMI DiabetesPedigreeFunction Age Outcome
            Pregnancies Glucose BloodPressure SkinThickness
          0
                          148.0
                                        72.0
                                                35.000000
                                                          79.799479 33.6
                                                                                        0.627
                                                                                              50
                                                          79.799479 26.6
          1
                          85.0
                                        66.0
                                                29.000000
                                                                                        0.351
                                                                                                        0
          2
                          183.0
                                        64.0
                                                20.536458
                                                          79.799479 23.3
                                                                                        0.672
                                                                                              32
```

23.000000

Countplot

3

```
In [8]: sns.set_style("darkgrid")
    sns.countplot(df['Outcome'])
    plt.title("Countplot of Outcome")
    plt.xlabel('Outcome')
    plt.ylabel('count')
    print('count of class is:\n',df['Outcome'].value_counts())
```

94.000000 28.1

35.000000 168.000000 43.1

0.167

2.288

33

0

1

count of class is: 0 500 1 268

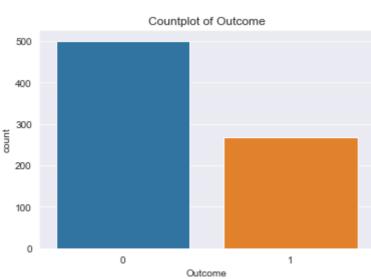
Name: Outcome, dtype: int64

89.0

137.0

66.0

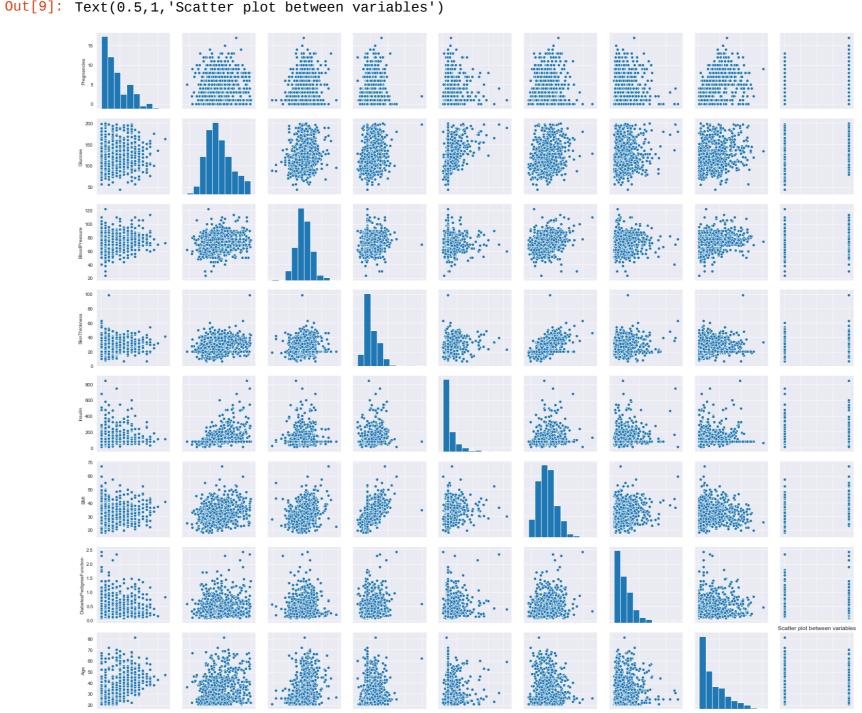
40.0



We can see that both class is balanced so we need not to perform any sampling method to maintain the balance between both classes. Therefor i will be directly using this data in training and testing purpose without performing any sampling method.

Scatter Plot

```
In [9]: sns.pairplot(df)
plt.title('Scatter plot between variables')
```



We can see from scatter plot that there is no strong multicolinearity among features, but between skin thickness and BMI, Pregnancies and age it looks like there is small chance of positive correlation..i will explore more when analyzing correlation

Correlation Analysis

In [10]: df.corr()

Out[10]:

Pregnancies Glucose BloodPressure SkinThickness Insulin BMI DiabetesPedigreeFunction

	i regnanoies	Ciuoosc	Biodui iessuie	Citini monness	mounn	D.VII.	Diabetesi edigicei dilotto
Pregnancies	1.000000	0.127964	0.208984	0.013376	-0.018082	0.021546	-0.03352
Glucose	0.127964	1.000000	0.219666	0.160766	0.396597	0.231478	0.13710
BloodPressure	0.208984	0.219666	1.000000	0.134155	0.010926	0.281231	0.00037
SkinThickness	0.013376	0.160766	0.134155	1.000000	0.240361	0.535703	0.15496
Insulin	-0.018082	0.396597	0.010926	0.240361	1.000000	0.189856	0.15780
ВМІ	0.021546	0.231478	0.281231	0.535703	0.189856	1.000000	0.15350
DiabetesPedigreeFunction	-0.033523	0.137106	0.000371	0.154961	0.157806	0.153508	1.00000
Age	0.544341	0.266600	0.326740	0.026423	0.038652	0.025748	0.03356
Outcome	0.221898	0.492908	0.162986	0.175026	0.179185	0.312254	0.17384

In [13]: plt.figure(dpi=80)
sns.heatmap(df.corr(),cmap= 'coolwarm')

