

# Foundations of Data Science Project - Diabetes Analysis

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## Context

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Diabetes is one of the most frequent diseases worldwide and the number of diabetic patients are growing over the years. The main cause of diabetes remains unknown, yet scientists believe that both genetic factors and environmental lifestyle play a major role in diabetes.

A few years ago research was done on a tribe in America which is called the Pima tribe (also known as the Pima Indians). In this tribe, it was found that the ladies are prone to diabetes very early. Several constraints were placed on the selection of these instances from a larger database. In particular, all patients were females at least 21 years old of Pima Indian heritage.

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## Objective

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Here, we are analyzing different aspects of Diabetes in the Pima Indians tribe by doing Exploratory Data Analysis.

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## Data Dictionary

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The dataset has the following information:

- Pregnancies: Number of times pregnant
- Glucose: Plasma glucose concentration over 2 hours in an oral glucose tolerance test
- BloodPressure: Diastolic blood pressure (mm Hg)
- SkinThickness: Triceps skin fold thickness (mm)
- Insulin: 2-Hour serum insulin ( $\mu$ U/ml)
- BMI: Body mass index ( $\text{weight in kg}/(\text{height in m})^2$ )
- DiabetesPedigreeFunction: A function that scores the likelihood of diabetes based on family history.
- Age: Age in years
- Outcome: Class variable (0: a person is not diabetic or 1: a person is diabetic)

## Q 1: Import the necessary libraries and briefly explain the use of each library (3 Marks)

In [7]: # remove \_\_\_\_\_ & write the appropriate library name

```
import numpy as np
import pandas as pd

import seaborn as sns
import matplotlib as plt
%matplotlib inline
```

Write your Answer here:

Ans 1:numpy is used for high function mathematical problems that have multideminsions. matplotlibis use for static visuabls. pandas is used for satistics and data science. seaborn is also a visual libery that makes attractive visuals for data

## Q 2: Read the given dataset (1 Mark)

In [8]: # remove \_\_\_\_\_ & write the appropriate function name

```
pima = pd.read_csv('diabetes.csv')
```

## Q3. Show the last 10 records of the dataset. How many columns are there? (1 Mark)

In [9]: # remove \_\_\_\_\_ and write the appropriate number in the function

```
pima.tail(10)
```

Out[9]:

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	DiabetesPedigre
758	1	106	76	20	79	37.5	
759	6	190	92	20	79	35.5	
760	2	88	58	26	16	28.4	
761	9	170	74	31	79	44.0	
762	9	89	62	20	79	22.5	
763	10	101	76	48	180	32.9	
764	2	122	70	27	79	36.8	
765	5	121	72	23	112	26.2	
766	1	126	60	20	79	30.1	
767	1	93	70	31	79	30.4	

Write your Answer here:

Ans 3:9

Q4. Show the first 10 records of the dataset (1 Mark)

```
In [... # remove ____ & write the appropriate function name and the number o  
pima.head(10)
```

```
Out[10]:
```

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	DiabetesPer
0	6	148	72	35	79	33.600000	
1	1	85	66	29	79	26.600000	
2	8	183	64	20	79	23.300000	
3	1	89	66	23	94	28.100000	
4	0	137	40	35	168	43.100000	
5	5	116	74	20	79	25.600000	
6	3	78	50	32	88	31.000000	
7	10	115	69	20	79	35.300000	
8	2	197	70	45	543	30.500000	
9	8	125	96	20	79	31.992578	

Q5. What do you understand by the dimension of the dataset? Find the dimension of the `pima` dataframe. (1 Mark)

```
In [11]: # remove ____ & write the appropriate function name  
pima.shape  
Out[11]: (768, 9)
```

Write your Answer here:

Ans 5:the dataframe is 768 rows x 9 columns.

Q6. What do you understand by the size of the dataset? Find the size of the `pima` dataframe. (1 Mark)

```
In [12]: # remove _____ & write the appropriate function name

        pima.size

Out[12]: 6912
```

Write your Answer here:

Ans 6:there is 6912 variables

Q7. What are the data types of all the variables in the data set? (2 Marks)

**Hint: Use the `info()` function to get all the information about the dataset.**

```
In [13]: # remove _____ & write the appropriate function name

        pima.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
```

Write your Answer here:

Ans 7:pregnancies, glucose, blood pressure, skin thickness, insulin, bmi, diabetes pedigree function, age outcome.

Q8. What do we mean by missing values? Are there any missing values in the `pima` dataframe? (2 Marks)

```
In [14]: # remove ____ & write the appropriate function name

        pima.isnull().values.any()

Out[14]: False
```

Write your Answer here:

Ans 8: no missing values

Q9. What do the summary statistics of the data represent? Find the summary statistics for all variables except 'Outcome' in the `pima` data. Take one column/variable from the output table and explain all its statistical measures. (3 Marks)

```
In [15]: # remove ____ & write the appropriate function name

        pima.iloc[:,0:8].info()

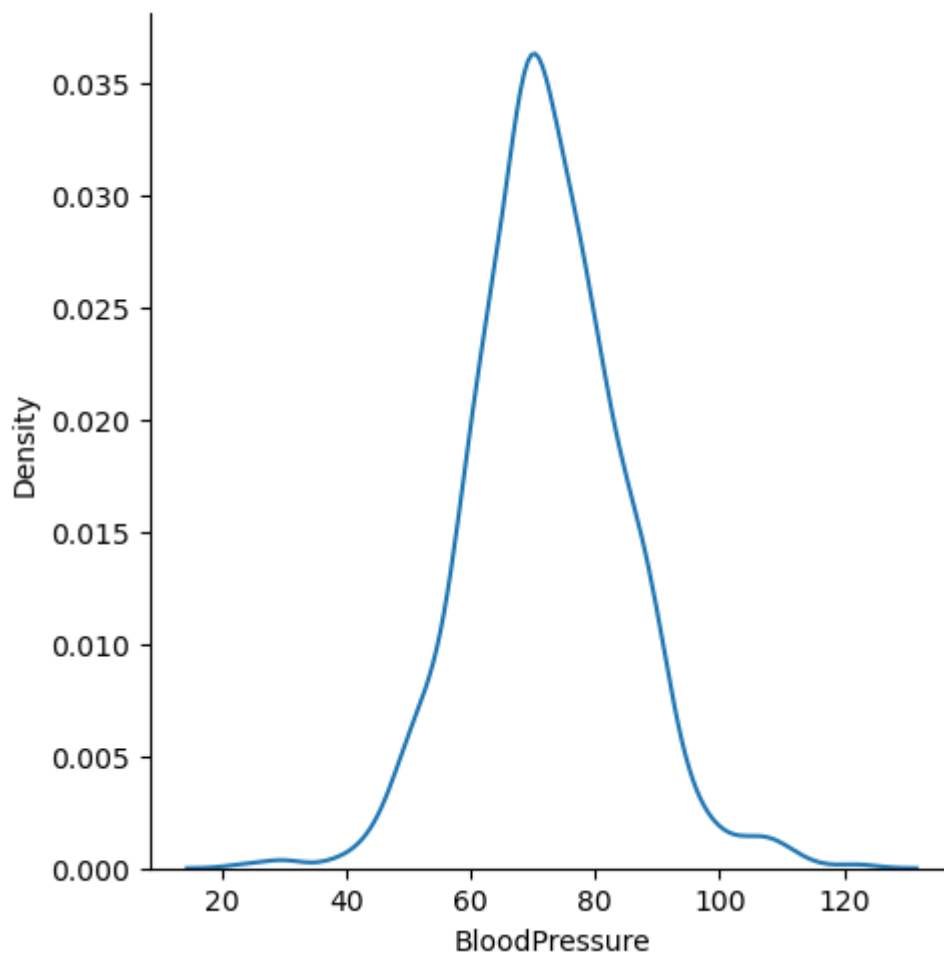
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 768 entries, 0 to 767
Data columns (total 8 columns):
 #   Column                Non-Null Count  Dtype
---  -
 0   Pregnancies           768 non-null    int64
 1   Glucose               768 non-null    int64
 2   BloodPressure         768 non-null    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
dtypes: float64(2), int64(6)
memory usage: 48.1 KB
```

Write your Answer here:

Ans 9: pregnancies, glucose, blood pressure, skin thickness, age, and insulin have 64 whole number values. bmi, diabetes pedigree function have decimal number. each of which has 64 values.

Q 10. Plot the distribution plot for the variable 'BloodPressure'. Write detailed observations from the plot. (2 Marks)

```
In [16]: # remove ____ & write the appropriate library name
import seaborn as sns
import matplotlib.pyplot as plt
sns.displot(pima['BloodPressure'], kind='kde')
plt.show()
```



Write your Answer here:

Ans 10: the majority have a blood pressure in the 70s. Very little people have it in the 40s and under and 100 and above.

Q 11. What is the 'BMI' of the person having the highest 'Glucose'? (1 Mark)

```
In [17]: # remove _____ & write the appropriate function name
```

```
pima[pima['Glucose']==pima['Glucose'].max()][ 'BMI' ]
```

```
Out[17]: 661    42.9  
         Name: BMI, dtype: float64
```

Write your Answer here:

Ans 11:the person with the highest glucose level has a marker of 42.9

Q12.

12.1 What is the mean of the variable 'BMI'?

12.2 What is the median of the variable 'BMI'?

12.3 What is the mode of the variable 'BMI'?

12.4 Are the three measures of central tendency equal?

(3 Marks)

```
In [18]: # remove _____ & write the appropriate function name
```

```
m1 = pima['BMI'].mean() # mean  
print(m1)  
m2 = pima['BMI'].median() # median  
print(m2)  
m3 = pima['BMI'].mode()[0] # mode  
print(m3)
```

```
32.45080515543617  
32.0  
32.0
```

Write your Answer here:

Ans 12:the 3 measurement are not all the same.the mean is off by 0.4.

Q13. How many women's 'Glucose' levels are above the mean level of 'Glucose'? (1 Mark)

```
In [19]: # remove _____ & write the appropriate function name
```

```
pima[pima['Glucose']>pima['Glucose'].mean()].shape[0]
```

```
Out[19]: 343
```

Write your Answer here:

Ans 13:343

Q14. How many women have their 'BloodPressure' equal to the median of 'BloodPressure' and their 'BMI' less than the median of 'BMI'? (2 Marks)

```
In [... # remove _____ & write the appropriate column name
```

```
pima[(pima['BloodPressure']==pima['BloodPressure'].median()) & (pima[
```



```
Out[20]:
```

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	DiabetesPedigr
14	5	166	72	19	175	25.8	
93	4	134	72	20	79	23.8	
103	1	81	72	18	40	26.6	
205	5	111	72	28	79	23.9	
299	8	112	72	20	79	23.6	
325	1	157	72	21	168	25.6	
330	8	118	72	19	79	23.1	
366	6	124	72	20	79	27.6	
380	1	107	72	30	82	30.8	
393	4	116	72	12	87	22.1	
406	4	115	72	20	79	28.9	
446	1	100	72	12	70	25.3	
460	9	120	72	22	56	20.8	
488	4	99	72	17	79	25.6	
497	2	81	72	15	76	30.1	
510	12	84	72	31	79	29.7	
568	4	154	72	29	126	31.3	
615	3	106	72	20	79	25.8	
635	13	104	72	20	79	31.2	
644	3	103	72	30	152	27.6	
717	10	94	72	18	79	23.1	
765	5	121	72	23	112	26.2	

Write your Answer here:

Ans 14 there are 22 women

Q15. Create a pairplot for the variables 'Glucose', 'SkinThickness', and 'DiabetesPedigreeFunction'. Write your observations from the plot. (4 Marks)

In [... # remove \_\_\_\_\_ & write the appropriate function name

```
sns.pairplot(data=pima,vars=['Glucose', 'SkinThickness', 'DiabetesPed:
plt.show()
```

```

-----
AttributeError                                Traceback (most recent call last)
~\AppData\Local\Temp\ipykernel_21692\3835635000.py in <module>
      1 # remove ____ & write the appropriate function name
      2
----> 3 sns.pairpolr(data=pima,vars=['Glucose', 'SkinThickness', 'Diabetes
PedigreeFunction'], hue='Outcome')
      4 plt.show()

```

**AttributeError:** module 'seaborn' has no attribute 'pairpolr'

Write your Answer here:

Ans 15: there is not negative or positive correlation, however there is a range where central tendency is and a cut from both tails.

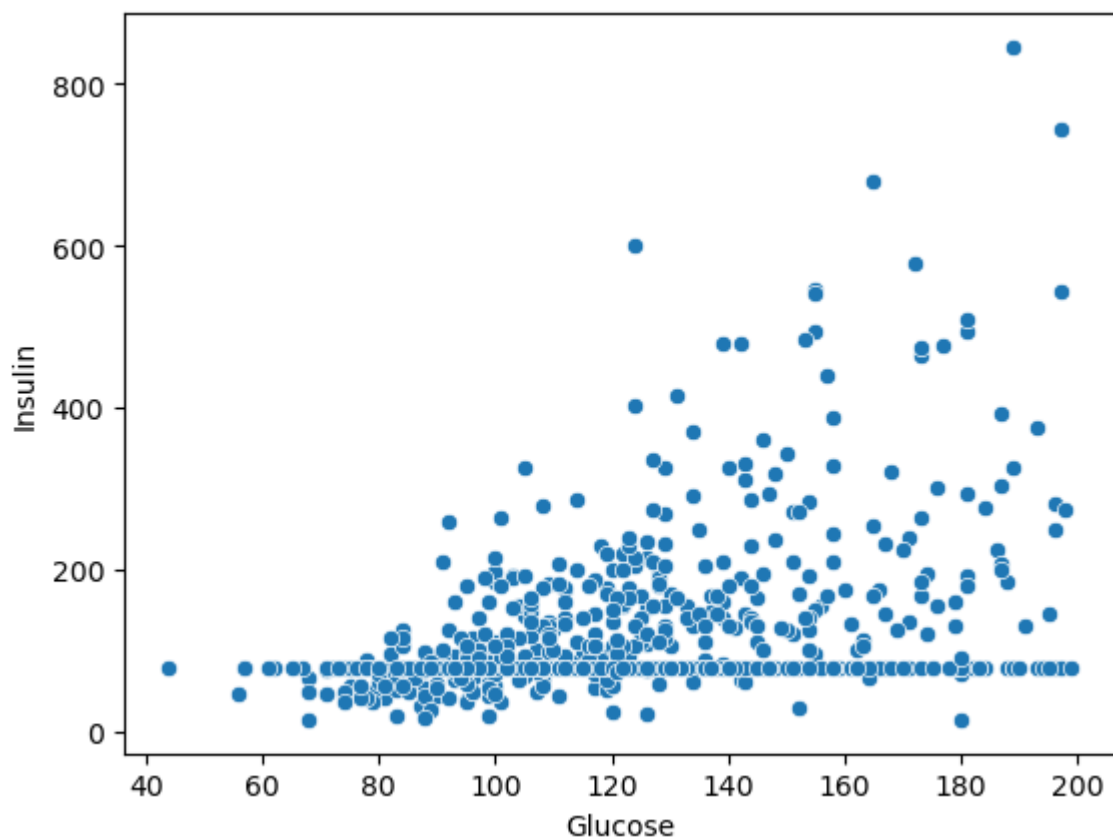
**Q16.** Plot the scatterplot between 'Glucose' and 'Insulin'. Write your observations from the plot. (2 Marks)

```
In [40]: # remove ____ & write the appropriate function name
```

```

sns.scatterplot(x='Glucose',y='Insulin',data=pima)
plt.pyplot.show()

```



Write your Answer here:

Ans 16: there is a positive correlation with glucose and insulin. the higher glucose goes up the higher insulin goes up.

Q 17. Plot the boxplot for the 'Age' variable. Are there outliers? (2 Marks)

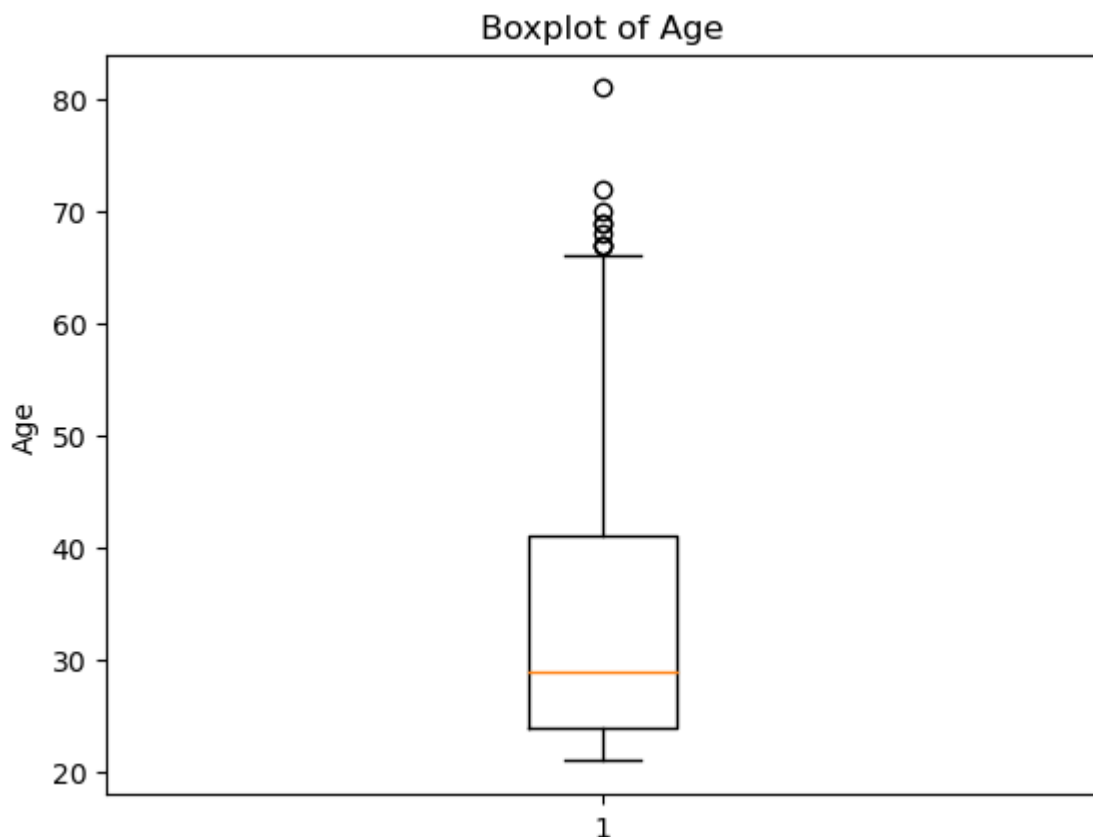
```
In [24]: import numpy as np
import pandas as pd

import seaborn as sns
import matplotlib as plt
%matplotlib inline
```

```
In [37]: # remove ____ & write the appropriate function and column name
```

```
plt.pyplot.boxplot(pima['Age'])

plt.pyplot.title('Boxplot of Age')
plt.pyplot.ylabel('Age')
plt.pyplot.show()
```



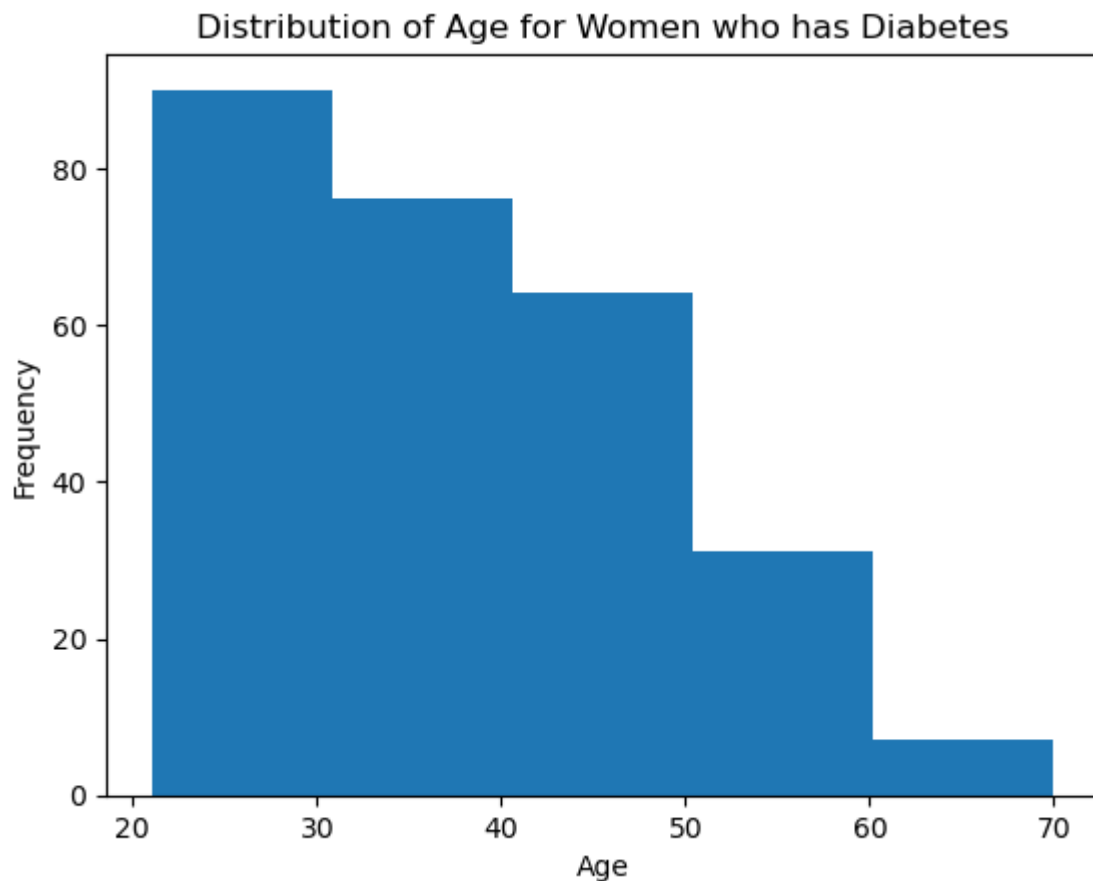
Write your Answer here:

Ans 17:there are 6 outliers.

Q18. Plot histograms for the 'Age' variable to understand the number of women in different age groups given whether they have diabetes or not. Explain both histograms and compare them. (3 Marks)

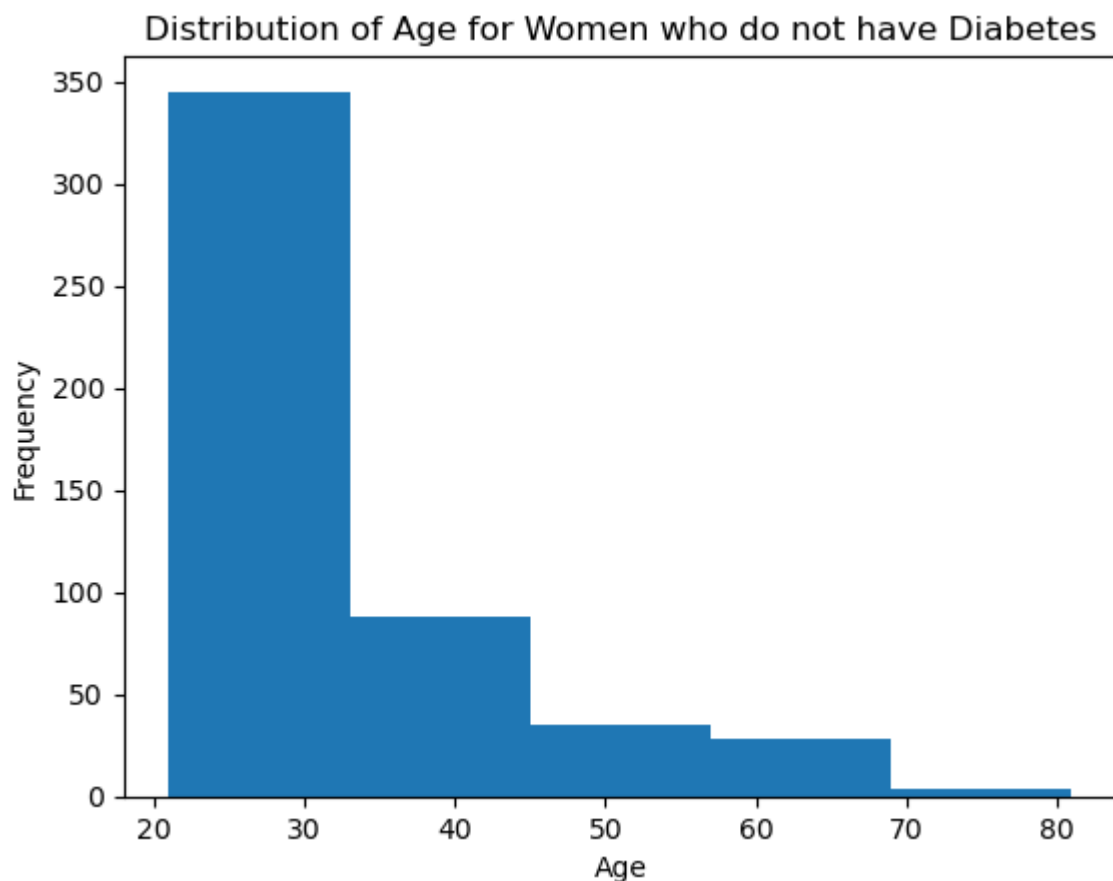
In [36... # remove \_\_\_\_ & write the appropriate function and column name

```
plt.pyplot.hist(pima[pima['Outcome']==1]['Age'], bins = 5)
plt.pyplot.title('Distribution of Age for Women who has Diabetes')
plt.pyplot.xlabel('Age')
plt.pyplot.ylabel('Frequency')
plt.pyplot.show()
```



In [41... # remove \_\_\_\_ & write the appropriate function and column name

```
plt.pyplot.hist(pima[pima['Outcome']==0]['Age'], bins = 5)
plt.pyplot.title('Distribution of Age for Women who do not have Dial')
plt.pyplot.xlabel('Age')
plt.pyplot.ylabel('Frequency')
plt.pyplot.show()
```



Write your Answer here:

Ans 18: women who have diabetes have more frequency in ages 20-60 compared to women who do not have diabetes, which has a significant drop-off after age 30.

**Q 19. What is the Interquartile Range of all the variables? Why is this used? Which plot visualizes the same? (2 Marks)**

In [42]: # remove \_\_\_\_ & write the appropriate variable name

```
Q1 = pima.quantile(0.25)
Q3 = pima.quantile(0.75)
IQR = Q3 - Q1
print(IQR)
```

Pregnancies	5.0000
Glucose	40.5000
BloodPressure	16.0000
SkinThickness	12.0000
Insulin	48.2500
BMI	9.1000
DiabetesPedigreeFunction	0.3825
Age	17.0000
Outcome	1.0000
dtype: float64	

Write your Answer here:

Ans 19: its used to see the first Q1 and see the difference between the Q1 and Q3. the box plot visualizes the same information as the interquartile range. It shows the minimum, first quartile, median, third quartile, and maximum of a dataset, which are the same values as the interquartile range.

## Q 20. Find and visualize the correlation matrix. Write your observations from the plot. (3 Marks)

In [43]: # remove \_\_\_\_\_ & write the appropriate function name and run the co

```
corr_matrix = pima.iloc[:,0:8].corr()
```

```
corr_matrix
```

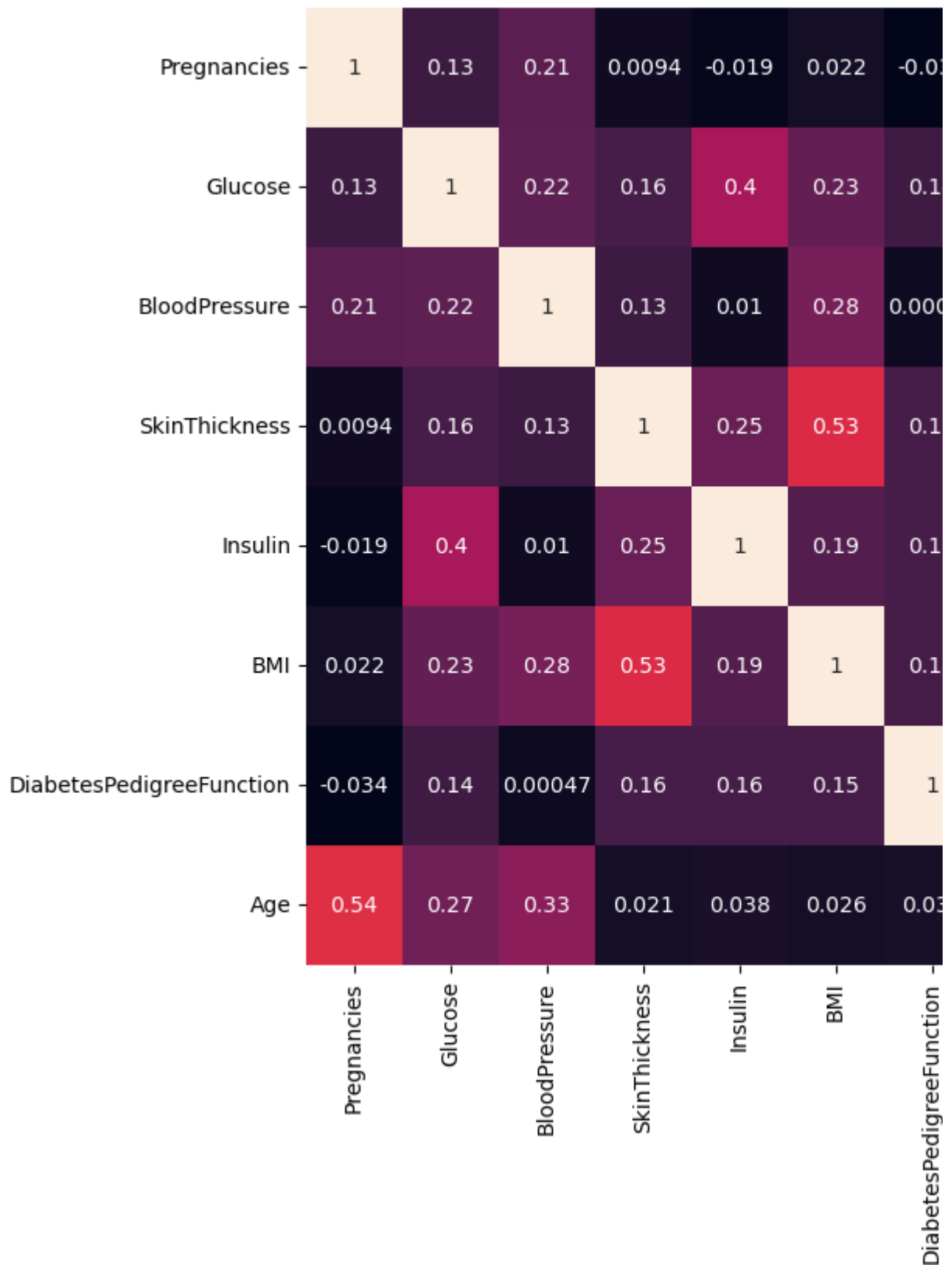
Out[43]:

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insu
Pregnancies	1.000000	0.128022	0.208987	0.009393	-0.0187
Glucose	0.128022	1.000000	0.219765	0.158060	0.3961
BloodPressure	0.208987	0.219765	1.000000	0.130403	0.0104
SkinThickness	0.009393	0.158060	0.130403	1.000000	0.2454
Insulin	-0.018780	0.396137	0.010492	0.245410	1.0000
BMI	0.021546	0.231464	0.281222	0.532552	0.1899
DiabetesPedigreeFunction	-0.033523	0.137158	0.000471	0.157196	0.1582
Age	0.544341	0.266673	0.326791	0.020582	0.0376

In [46]: # remove \_\_\_\_\_ & write the appropriate function name

```
plt.pyplot.figure(figsize=(8,8))
sns.heatmap(corr_matrix, annot = True)
```

```
# display the plot
plt.pyplot.show()
```



Write your Answer here:

Ans 20: age has a strong correlation with pregnancy. diabetes pedigree overall has a weak correlation. bmi has a strong correlation with skin thickness. insulin has a moderate correlation with glucose.

