Problem statement

Data collection

Importing libraries

```
In [1]:
```

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
```

Importing dataset

```
In [2]: data-nd noad csy(n"C:)Usons)uson
```

```
data=pd.read_csv(r"C:\Users\user\Downloads\4_drug200 - 4_drug200.csv")
data
```

Out[2]:		Age	Sex	ВР	Cholesterol	Na_to_K	Drug
	0	23	F	HIGH	HIGH	25.355	drugY
	1	47	М	LOW	HIGH	13.093	drugC
	2	47	М	LOW	HIGH	10.114	drugC
	3	28	F	NORMAL	HIGH	7.798	drugX
	4	61	F	LOW	HIGH	18.043	drugY
	•••			•••		•••	
	195	56	F	LOW	HIGH	11.567	drugC
	196	16	М	LOW	HIGH	12.006	drugC
	197	52	М	NORMAL	HIGH	9.894	drugX
	198	23	М	NORMAL	NORMAL	14.020	drugX

LOW

NORMAL

200 rows × 6 columns

40

head

199

```
In [3]:
```

```
# to display first 8 dataset values
da=data.head(8)
da
```

11.349 drugX

Out[3]:		Age	Sex	ВР	Cholesterol	Na_to_K	Drug
	0	23	F	HIGH	HIGH	25.355	drugY
	1	47	М	LOW	HIGH	13.093	drugC
	2	47	М	LOW	HIGH	10.114	drugC
	3	28	F	NORMAL	HIGH	7.798	drugX
	4	61	F	LOW	HIGH	18.043	drugY
	5	22	F	NORMAL	HIGH	8.607	drugX
	6	49	F	NORMAL	HIGH	16.275	drugY
	7	41	М	LOW	HIGH	11.037	drugC

info

```
In [4]:
         # to identify missing values
         data.info()
        <class 'pandas.core.frame.DataFrame'>
        RangeIndex: 200 entries, 0 to 199
        Data columns (total 6 columns):
             Column
                         Non-Null Count Dtype
         #
                         -----
                                         int64
         0
                         200 non-null
             Age
         1
                         200 non-null
                                         object
             Sex
         2
                         200 non-null
                                         object
             BP
         3
             Cholesterol 200 non-null
                                         object
             Na_to_K
                         200 non-null
                                         float64
         5
                         200 non-null
                                         object
             Drug
        dtypes: float64(1), int64(1), object(4)
        memory usage: 9.5+ KB
```

describe

count	200.000000	200.000000
mean	44.315000	16.084485
std	16.544315	7.223956
min	15.000000	6.269000
25%	31.000000	10.445500
50%	45.000000	13.936500
75%	58.000000	19.380000

```
        Age
        Na_to_K

        max
        74.000000
        38.247000
```

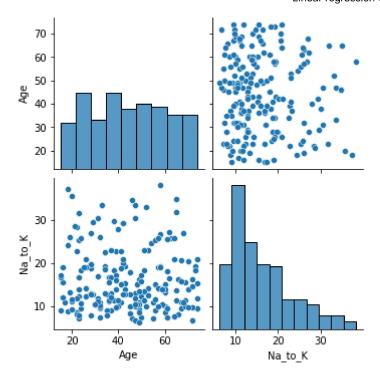
columns

```
In [6]:
         # to display headings of the dataset
         data.columns
Out[6]: Index(['Age', 'Sex', 'BP', 'Cholesterol', 'Na_to_K', 'Drug'], dtype='object')
In [7]:
         a=data.dropna(axis=1)
Out[7]:
             Age Sex
                            BP Cholesterol Na_to_K Drug
          0
               23
                    F
                          HIGH
                                     HIGH
                                            25.355 drugY
           1
               47
                    Μ
                          LOW
                                     HIGH
                                            13.093 drugC
           2
                                     HIGH
               47
                          LOW
                                            10.114 drugC
                    Μ
           3
               28
                    F NORMAL
                                     HIGH
                                            7.798 drugX
               61
                    F
                          LOW
                                     HIGH
                                            18.043 drugY
        195
               56
                    F
                          LOW
                                     HIGH
                                            11.567 drugC
        196
               16
                          LOW
                                     HIGH
                                            12.006 drugC
                    М
        197
               52
                    M NORMAL
                                     HIGH
                                            9.894 drugX
        198
               23
                       NORMAL
                                  NORMAL
                                            14.020 drugX
        199
               40
                          LOW
                                  NORMAL
                                            11.349 drugX
       200 rows × 6 columns
In [8]:
         a.columns
Out[8]: Index(['Age', 'Sex', 'BP', 'Cholesterol', 'Na_to_K', 'Drug'], dtype='object')
```

EDA and Visualization

```
In [9]: sns.pairplot(a)
```

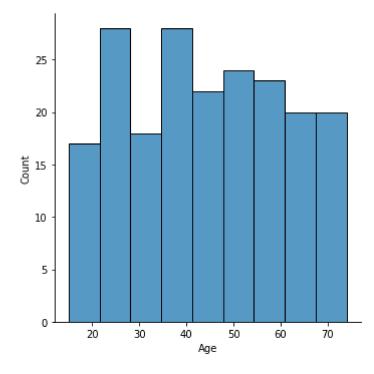
Out[9]: <seaborn.axisgrid.PairGrid at 0x246963d7ac0>



distribution plot

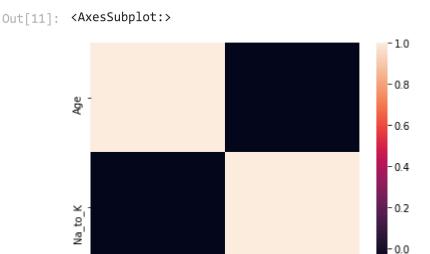
```
In [10]: sns.displot(a["Age"])
```

Out[10]: <seaborn.axisgrid.FacetGrid at 0x24696e0c430>



correlation

```
In [11]:
    dat=data[['Age', 'Sex', 'BP', 'Cholesterol', 'Na_to_K', 'Drug']]
    sns.heatmap(dat.corr())
```



Age

To train the model-Model Building

Na to K

```
In [12]:
          x=a[['Age']]
          y=a['Age']
In [13]:
          # to split my dataset into training and test data
          from sklearn.model_selection import train_test_split
          x_train,x_test,y_train,y_test=train_test_split(x,y,test_size=0.3)
In [14]:
          from sklearn.linear model import LinearRegression
          lr= LinearRegression()
          lr.fit(x_train,y_train)
Out[14]: LinearRegression()
In [15]:
          print(lr.intercept_)
          -7.105427357601002e-15
In [16]:
          coeff=pd.DataFrame(lr.coef_,x.columns,columns=['Co-efficient'])
          coeff
Out[16]:
              Co-efficient
                      1.0
         Age
In [17]:
          prediction=lr.predict(x_test)
          plt.scatter(y_test,prediction)
Out[17]: <matplotlib.collections.PathCollection at 0x24698788850>
```

