SUMESH R -20104169

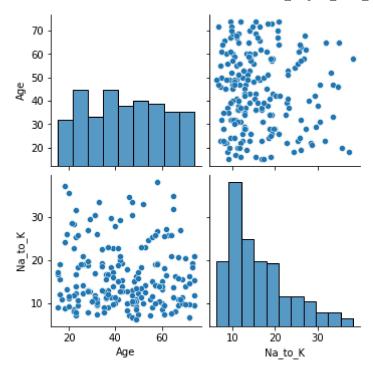
```
In [1]:
          import numpy as np
          import pandas as pd
          import matplotlib.pyplot as plt
          import seaborn as sns
         from sklearn.model_selection import train_test_split
          from sklearn.linear_model import LinearRegression
In [2]:
          df = pd.read_csv("4_drug200.csv")
          # .dropna(axis="columns")
          df
Out[2]:
              Age
                  Sex
                            BP Cholesterol Na_to_K
                                                     Drug
           0
               23
                     F
                          HIGH
                                      HIGH
                                             25.355
                                                    drugY
           1
               47
                           LOW
                                      HIGH
                                             13.093 drugC
                    Μ
           2
               47
                           LOW
                                      HIGH
                                             10.114 drugC
                    Μ
           3
               28
                     F NORMAL
                                      HIGH
                                                    drugX
                                              7.798
               61
                     F
                           LOW
                                      HIGH
                                                   drugY
                                             18.043
         195
                     F
                           LOW
               56
                                      HIGH
                                             11.567 drugC
         196
                           LOW
               16
                    Μ
                                      HIGH
                                             12.006 drugC
         197
                    Μ
                       NORMAL
                                      HIGH
                                              9.894 drugX
               52
         198
               23
                       NORMAL
                                   NORMAL
                                             14.020 drugX
                    M
         199
               40
                           LOW
                                   NORMAL
                                             11.349 drugX
        200 rows × 6 columns
In [3]:
          df.head()
Out[3]:
            Age Sex
                          BP Cholesterol Na_to_K
                                                   Drug
                   F
         0
             23
                        HIGH
                                    HIGH
                                           25.355 drugY
         1
                         LOW
             47
                                    HIGH
                                           13.093 drugC
                  Μ
         2
             47
                         LOW
                                    HIGH
                                           10.114 drugC
                  М
             28
                   F NORMAL
                                    HIGH
                                            7.798 drugX
                         LOW
                                           18.043 drugY
             61
                                    HIGH
```

Data cleaning and pre processing

```
In [4]:
         df.info()
         <class 'pandas.core.frame.DataFrame'>
         RangeIndex: 200 entries, 0 to 199
        Data columns (total 6 columns):
              Column
                           Non-Null Count Dtype
          0
              Age
                            200 non-null
                                             int64
          1
                           200 non-null
              Sex
                                             object
          2
              BP
                           200 non-null
                                             object
          3
              Cholesterol 200 non-null
                                             object
              Na to K
                           200 non-null
                                             float64
                           200 non-null
              Drug
                                             object
         dtypes: float64(1), int64(1), object(4)
        memory usage: 9.5+ KB
In [5]:
         df.describe()
Out[5]:
                     Age
                            Na_to_K
         count 200.000000 200.000000
                44.315000
                           16.084485
         mean
           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
                74.000000
                           38.247000
          max
In [6]:
          df.columns
Out[6]: Index(['Age', 'Sex', 'BP', 'Cholesterol', 'Na_to_K', 'Drug'], dtype='object')
```

EDA and VISUALIZATION

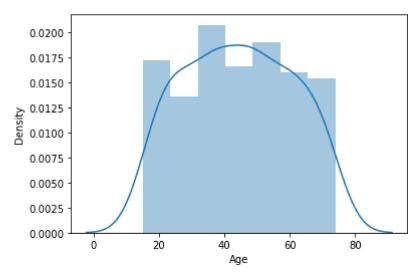
```
In [7]: sns.pairplot(df)
Out[7]: <seaborn.axisgrid.PairGrid at 0x2b10c766310>
```



In [8]: sns.distplot(df["Age"])

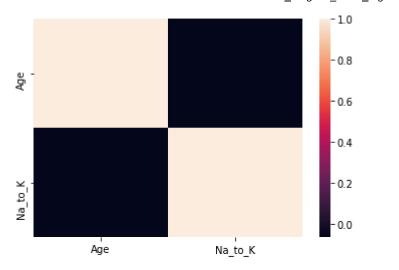
C:\ProgramData\Anaconda3\lib\site-packages\seaborn\distributions.py:2557: FutureWarning:
 distplot` is a deprecated function and will be removed in a future version. Please adap
 t your code to use either `displot` (a figure-level function with similar flexibility) o
 r `histplot` (an axes-level function for histograms).
 warnings.warn(msg, FutureWarning)

Out[8]: <AxesSubplot:xlabel='Age', ylabel='Density'>



```
In [9]: df1 = df[['Age', 'Sex', 'BP', 'Cholesterol', 'Na_to_K', 'Drug']]
In [10]: sns.heatmap(df1.corr())
```

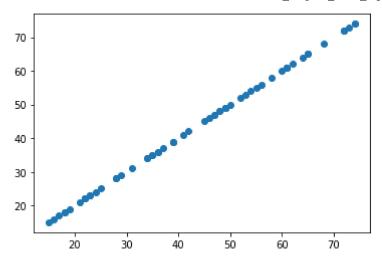
Out[10]: <AxesSubplot:>



split the data into training and test data

```
In [12]:
          x_train, x_test, y_train, y_test = train_test_split(x,y,test_size=0.3)
In [13]:
          lr = LinearRegression()
          lr.fit(x train, y train)
Out[13]: LinearRegression()
In [14]:
          lr.intercept
         -7.105427357601002e-15
Out[14]:
In [15]:
          coeff = pd.DataFrame(lr.coef_, x.columns, columns =['Co-efficient'])
          coeff
                   Co-efficient
Out[15]:
             Age 1.000000e+00
          Na to K 1.626388e-17
In [16]:
          prediction = lr.predict(x_test)
          plt.scatter(y_test, prediction)
```

Out[16]: <matplotlib.collections.PathCollection at 0x2b10c72a400>



In [17]: lr.score(x_test,y_test)

Out[17]: 1.0