```
In [1]:
          import numpy as np
          import pandas as pd
          import matplotlib.pyplot as plt
          import seaborn as sns
In [2]:
          df=pd.read csv("4 drug200.csv")
                                                      Drug
                             BP
                                 Cholesterol Na_to_K
Out[2]:
              Age Sex
           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
                                               7.798
                                                     drugX
           4
               61
                     F
                            LOW
                                       HIGH
                                              18.043 drugY
                             ...
                                         ...
           •••
                     ...
                                                  ...
                ...
         195
               56
                     F
                            LOW
                                       HIGH
                                              11.567 drugC
         196
                            LOW
                                              12.006 drugC
               16
                    Μ
                                       HIGH
         197
               52
                        NORMAL
                                       HIGH
                                               9.894 drugX
                    М
         198
               23
                    M NORMAL
                                    NORMAL
                                              14.020 drugX
         199
                     F
               40
                            LOW
                                    NORMAL
                                              11.349 drugX
        200 rows × 6 columns
In [3]:
          df.head()
Out[3]:
            Age Sex
                               Cholesterol Na_to_K
                                                    Drug
         0
             23
                   F
                         HIGH
                                     HIGH
                                            25.355 drugY
         1
             47
                  Μ
                         LOW
                                     HIGH
                                            13.093 drugC
         2
             47
                  М
                         LOW
                                     HIGH
                                            10.114 drugC
             28
                   F NORMAL
                                     HIGH
                                             7.798 drugX
```

DATA CLEANING AND DATA PREPROCESSING

```
In [4]:
    df.info()

(class 'pandas cono framo DataEnamo')
```

18.043 drugY

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 200 entries, 0 to 199
Data columns (total 6 columns):

LOW

HIGH

F

61

Out[5]:

In [6]:

In [7]:

Out[7]:

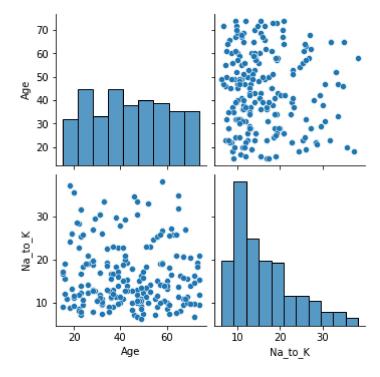
```
#
              Column
                            Non-Null Count Dtype
          0
              Age
                            200 non-null
                                              int64
              Sex
          1
                            200 non-null
                                              object
          2
              BP
                            200 non-null
                                              object
          3
              Cholesterol 200 non-null
                                              object
                            200 non-null
          4
              Na to K
                                              float64
          5
              Drug
                            200 non-null
                                              object
         dtypes: float64(1), int64(1), object(4)
         memory usage: 9.5+ KB
In [5]:
          df.describe()
                             Na_to_K
                     Age
         count 200.000000
                          200.000000
                44.315000
                            16.084485
         mean
                16.544315
                            7.223956
           std
                15.000000
                            6.269000
          min
          25%
                31.000000
                            10.445500
          50%
                45.000000
                            13.936500
          75%
                58.000000
                            19.380000
          max
                74.000000
                            38.247000
          df.columns
Out[6]: Index(['Age', 'Sex', 'BP', 'Cholesterol', 'Na_to_K', 'Drug'], dtype='object')
          df1=df.dropna(axis=1)
          df1
                             BP
                                 Cholesterol Na_to_K Drug
              Age Sex
               23
                     F
           0
                           HIGH
                                      HIGH
                                              25.355 drugY
           1
               47
                    Μ
                            LOW
                                      HIGH
                                              13.093 drugC
           2
               47
                    Μ
                            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
                            LOW
                                      HIGH
                                              12.006 drugC
               16
                    Μ
         197
               52
                        NORMAL
                                      HIGH
                                               9.894 drugX
                    М
         198
               23
                    Μ
                        NORMAL
                                   NORMAL
                                              14.020 drugX
         199
                     F
                                              11.349 drugX
               40
                            LOW
                                   NORMAL
```

200 rows × 6 columns

EDA AND VISUALIZATION

```
In [10]: sns.pairplot(df1)
```

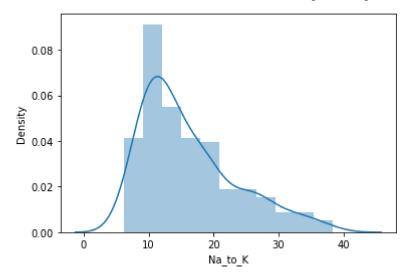
Out[10]: <seaborn.axisgrid.PairGrid at 0x1f83a5148b0>



```
In [11]: sns.distplot(df1['Na_to_K'])
```

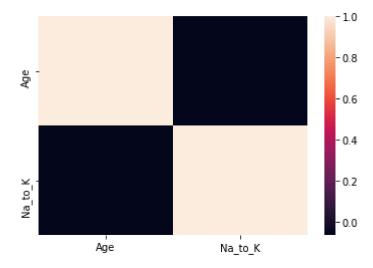
C:\ProgramData\Anaconda3\lib\site-packages\seaborn\distributions.py:2557: FutureWarning:
 `distplot` is a deprecated function and will be removed in a future version. Please adap
 t your code to use either `displot` (a figure-level function with similar flexibility) o
 r `histplot` (an axes-level function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

Out[11]: <AxesSubplot:xlabel='Na_to_K', ylabel='Density'>



```
In [12]: sns.heatmap(df1.corr())
```

Out[12]: <AxesSubplot:>



TO TRAIN THE MODEL AND MODEL BULDING

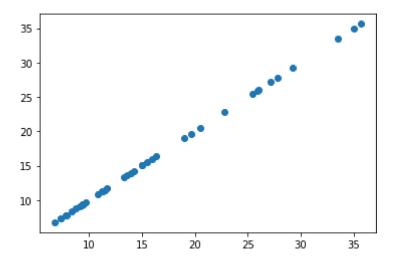
```
-7.105427357601002e-15
Out[16]:
```

```
In [17]:
          coeff=pd.DataFrame(lr.coef_,x.columns,columns=['Co-efficient'])
          coeff
```

Out[17]: Co-efficient 0.0 Age Na_to_K 1.0

```
In [18]:
          prediction =lr.predict(x_test)
          plt.scatter(y_test,prediction)
```

Out[18]: <matplotlib.collections.PathCollection at 0x1f83c7e0520>



ACCURACY

```
In [19]:
          lr.score(x_test,y_test)
Out[19]: 1.0
In [20]:
           lr.score(x_train,y_train)
Out[20]:
In [21]:
          from sklearn.linear_model import Ridge,Lasso
          rr=Ridge(alpha=10)
          rr.fit(x_train,y_train)
         Ridge(alpha=10)
Out[21]:
In [22]:
```

rr.score(x_test,y_test)

```
0.999998180981627
Out[22]:
In [23]:
          rr.score(x_train,y_train)
         0.99999826634159
Out[23]:
In [24]:
          la=Lasso(alpha=10)
          la.fit(x_train,y_train)
         Lasso(alpha=10)
Out[24]:
In [25]:
          la.score(x_test,y_test)
         0.9561160524705669
Out[25]:
In [26]:
          la.score(x_train,y_train)
         0.9561277317234831
Out[26]:
In [27]:
          from sklearn.linear_model import ElasticNet
          en=ElasticNet()
          en.fit(x train,y train)
Out[27]: ElasticNet()
In [28]:
          en.coef
         array([-0.
                            , 0.97927138])
Out[28]:
In [29]:
          en.intercept
         0.33284878190290357
Out[29]:
In [30]:
          prediction=en.predict(x_test)
In [31]:
          en.score(x_test,y_test)
         0.9995702099218393
Out[31]:
In [32]:
          from sklearn import metrics
          print(metrics.mean_absolute_error(y_test,prediction))
          print(metrics.mean_squared_error(y_test,prediction))
          print(np.sqrt(metrics.mean_squared_error(y_test,prediction)))
         0.14153108006342624
```

0.029500146144176297

0.1717560658147953