

```
In [1]: import numpy as np
import pandas as pd
import matplotlib.pyplot as py
import seaborn as sns
```

```
In [2]: d=pd.read_csv(r"C:\Users\user\Downloads\4_drug200 - 4_drug200.csv")
d
```

```
Out[2]:
```

	Age	Sex	BP	Cholesterol	Na_to_K	Drug
0	23	F	HIGH	HIGH	25.355	drugY
1	47	M	LOW	HIGH	13.093	drugC
2	47	M	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	M	LOW	HIGH	12.006	drugC
197	52	M	NORMAL	HIGH	9.894	drugX
198	23	M	NORMAL	NORMAL	14.020	drugX
199	40	F	LOW	NORMAL	11.349	drugX

200 rows × 6 columns

```
In [3]: d.head()
```

```
Out[3]:
```

	Age	Sex	BP	Cholesterol	Na_to_K	Drug
0	23	F	HIGH	HIGH	25.355	drugY
1	47	M	LOW	HIGH	13.093	drugC
2	47	M	LOW	HIGH	10.114	drugC
3	28	F	NORMAL	HIGH	7.798	drugX
4	61	F	LOW	HIGH	18.043	drugY

```
In [4]: d.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   Sex              200 non-null    object
```

```
2   BP          200 non-null    object
3   Cholesterol  200 non-null    object
4   Na_to_K     200 non-null    float64
5   Drug        200 non-null    object
dtypes: float64(1), int64(1), object(4)
memory usage: 9.5+ KB
```

```
In [5]: d.describe()
```

```
Out[5]:
```

	Age	Na_to_K
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
max	74.000000	38.247000

```
In [6]: d.columns
```

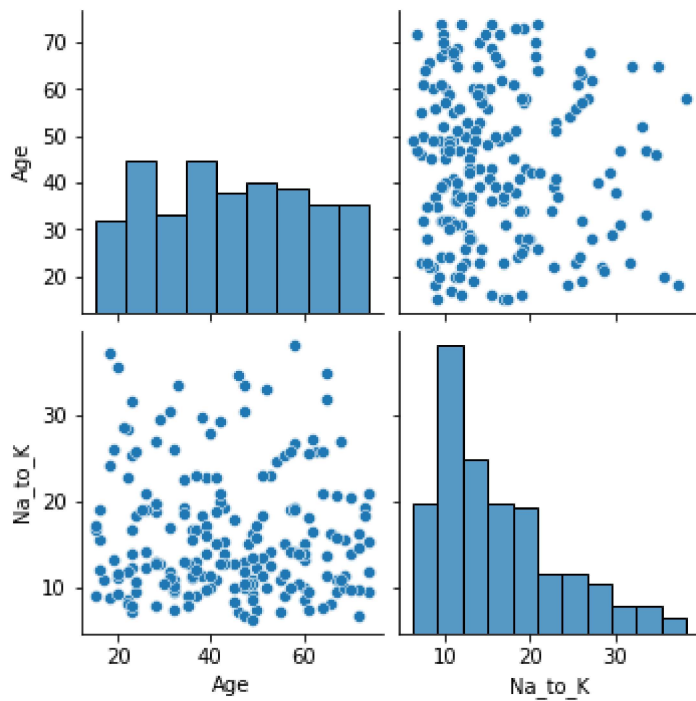
```
Out[6]: Index(['Age', 'Sex', 'BP', 'Cholesterol', 'Na_to_K', 'Drug'], dtype='object')
```

```
In [7]: d.index
```

```
Out[7]: RangeIndex(start=0, stop=200, step=1)
```

```
In [8]: sns.pairplot(d)
```

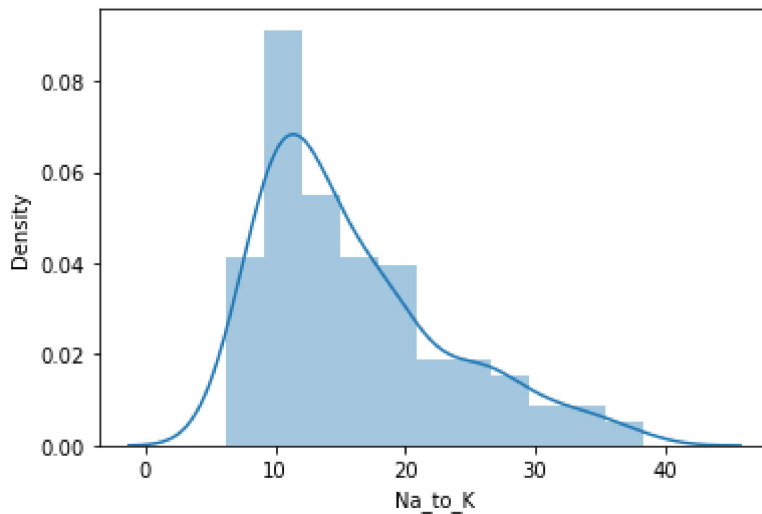
```
Out[8]: <seaborn.axisgrid.PairGrid at 0x1f5afb0a4f0>
```



```
In [9]: sns.distplot(d['Na_to_K'])
```

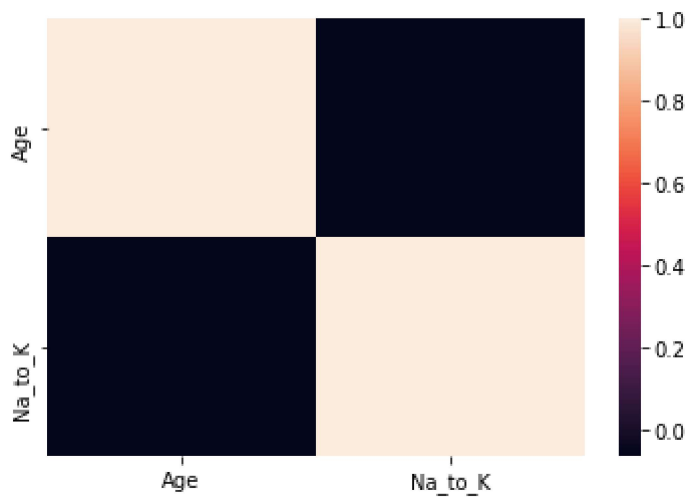
C:\ProgramData\Anaconda3\lib\site-packages\seaborn\distributions.py:2557: FutureWarning: `distplot` is a deprecated function and will be removed in a future version. Please adapt your code to use either `displot` (a figure-level function with similar flexibility) or `histplot` (an axes-level function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

```
Out[9]: <AxesSubplot:xlabel='Na_to_K', ylabel='Density'>
```



```
In [10]: d1=d[['Age', 'Sex', 'BP', 'Cholesterol', 'Na_to_K', 'Drug']]
sns.heatmap(d1.corr())
```

```
Out[10]: <AxesSubplot:>
```



```
In [11]: x=d1[['Age']]
         y=d1['Na_to_K']
```

```
In [12]: 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 [13]: from sklearn.linear_model import LinearRegression
```

```
In [14]: lr=LinearRegression()
         lr.fit(x_train,y_train)
```

```
Out[14]: LinearRegression()
```

```
In [15]: print(lr.intercept_)
```

```
17.17933073983918
```

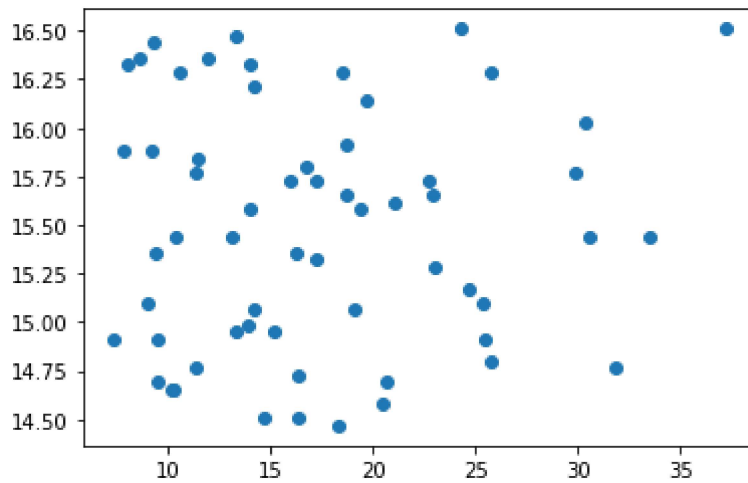
```
In [16]: coeff =pd.DataFrame(lr.coef_,x.columns,columns=["Co-efficient"])
         coeff
```

```
Out[16]:
```

	Co-efficient
Age	-0.037142

```
In [17]: prediction =lr.predict(x_test)
         py.scatter(y_test,prediction)
```

```
Out[17]: <matplotlib.collections.PathCollection at 0x1f5b1dc8cd0>
```



```
In [18]: print(lr.score(x_test,y_test))
```

```
-0.06699094296268249
```

```
In [19]: print(lr.score(x_train,y_train))
```

```
0.007313224337165747
```

```
In [20]: from sklearn.linear_model import Ridge,Lasso
```

```
In [21]: rr=Ridge(alpha=10)  
rr.fit(x_train,y_train)
```

```
Out[21]: Ridge(alpha=10)
```

```
In [22]: rr.score(x_test,y_test)
```

```
Out[22]: -0.06698719394941799
```

```
In [23]: la=Lasso(alpha=10)  
la.fit(x_train,y_train)
```

```
Out[23]: Lasso(alpha=10)
```

```
In [24]: la.score(x_test,y_test)
```

```
Out[24]: -0.0601250355138625
```

```
In [ ]:
```