

```
In [23]: import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
```

```
In [24]: data=pd.read_csv(r"C:\Users\user\Downloads\4_drug200.csv")
data
```

Out[24]:

	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 [25]: df=data.head(100)
df
```

Out[25]:

	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
...
95	36	M	LOW	NORMAL	11.424	drugX
96	58	F	LOW	HIGH	38.247	drugY
97	56	F	HIGH	HIGH	25.395	drugY
98	20	M	HIGH	NORMAL	35.639	drugY
99	15	F	HIGH	NORMAL	16.725	drugY

100 rows × 6 columns

```
In [26]: df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 100 entries, 0 to 99
Data columns (total 6 columns):
#   Column          Non-Null Count  Dtype
---  -
0   Age             100 non-null   int64
1   Sex             100 non-null   object
2   BP              100 non-null   object
3   Cholesterol      100 non-null   object
4   Na_to_K         100 non-null   float64
5   Drug            100 non-null   object
dtypes: float64(1), int64(1), object(4)
memory usage: 4.8+ KB
```

```
In [27]: df.describe()
```

```
Out[27]:
```

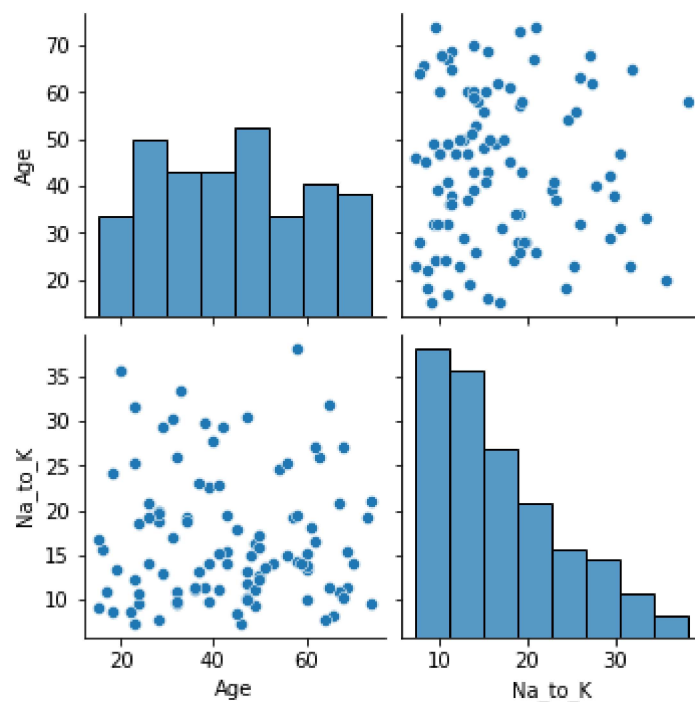
	Age	Na_to_K
count	100.000000	100.000000
mean	43.770000	16.823000
std	16.367531	7.257723
min	15.000000	7.285000
25%	30.500000	11.031250
50%	43.000000	15.025500
75%	58.000000	20.020250
max	74.000000	38.247000

```
In [28]: df.columns
```

```
Out[28]: Index(['Age', 'Sex', 'BP', 'Cholesterol', 'Na_to_K', 'Drug'], dtype='object')
```

```
In [29]: sns.pairplot(df)
```

```
Out[29]: <seaborn.axisgrid.PairGrid at 0x1ed590d4c70>
```



```
In [30]: da=df[['Age', 'Sex', 'BP', 'Cholesterol', 'Na_to_K', 'Drug']]
da
```

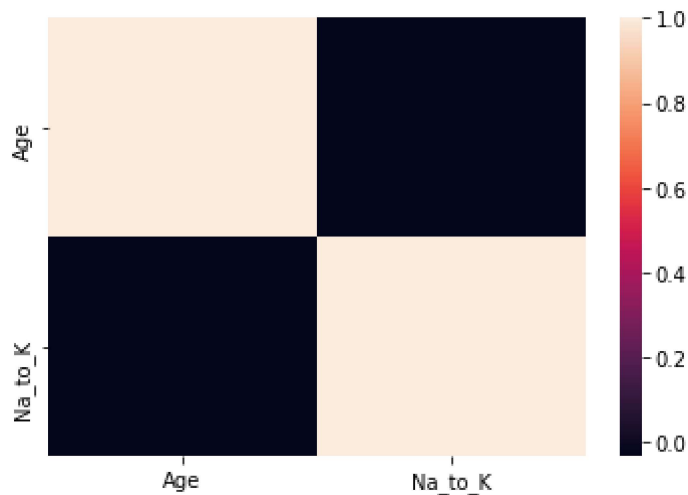
Out[30]:

	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
...
95	36	M	LOW	NORMAL	11.424	drugX
96	58	F	LOW	HIGH	38.247	drugY
97	56	F	HIGH	HIGH	25.395	drugY
98	20	M	HIGH	NORMAL	35.639	drugY
99	15	F	HIGH	NORMAL	16.725	drugY

100 rows × 6 columns

```
In [31]: sns.heatmap(da.corr())
```

Out[31]: <AxesSubplot:>



```
In [32]: x=df[['Na_to_K']]
y=df['Age']
```

```
In [33]: 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 [34]: from sklearn.linear_model import LinearRegression
```

```
lr=LinearRegression()
lr.fit(x_train,y_train)
```

```
Out[34]: LinearRegression()
```

```
In [35]: print(lr.intercept_)
```

```
48.23899261528304
```

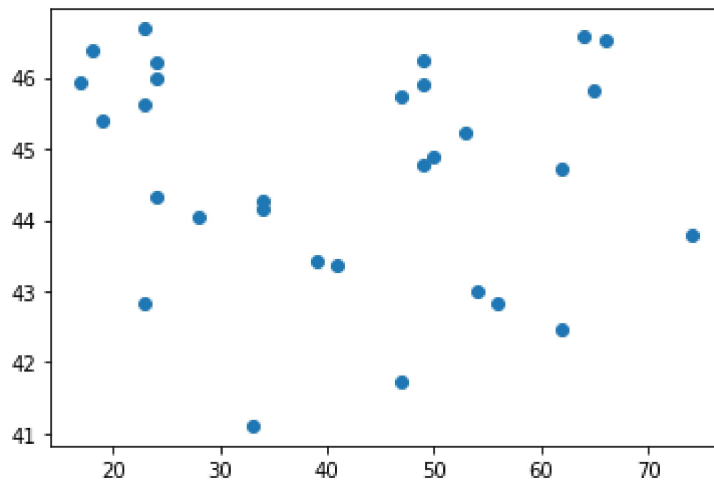
```
In [36]: coeff = pd.DataFrame(lr.coef_,x.columns,columns=['Co-efficient'])
coeff
```

```
Out[36]:
```

	Co-efficient
Na_to_K	-0.212774

```
In [37]: prediction=lr.predict(x_test)
plt.scatter(y_test,prediction)
```

```
Out[37]: <matplotlib.collections.PathCollection at 0x1ed5a35a7f0>
```



```
In [38]: print(lr.score(x_test,y_test))
```

```
-0.0613309781995135
```

```
In [39]: print(lr.score(x_train,y_train))
```

```
0.009292226156223271
```

```
In [40]: from sklearn.linear_model import Ridge,Lasso
```

```
In [41]: rr=Ridge(alpha=10)
         rr.fit(x_train,y_train)
```

Out[41]: Ridge(alpha=10)

```
In [42]: rr.score(x_test,y_test)
```

Out[42]: -0.0612282222195808

```
In [43]: la=Lasso(alpha=10)
         la.fit(x_train,y_train)
```

Out[43]: Lasso(alpha=10)

```
In [44]: la.score(x_test,y_test)
```

Out[44]: -0.033955212786694666

```
In [ ]:
```