

27-07-2023

```
In [ ]: # import libraries
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
```

```
In [104]: x=pd.read_csv(r"C:\Users\user\Downloads\4_drug200 - 4_drug200.csv")
```

Out[104]:

	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 [105]: x=x.head(100)
```

```
Out[105]:
```

	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 [106]:
```

```
<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 [107]:
```

```
Out[107]: Index(['Age', 'Sex', 'BP', 'Cholesterol', 'Na_to_K', 'Drug'], dtype='object')
```

```
In [109]: d=x[['Age', 'Sex', 'BP', 'Cholesterol', 'Na_to_K', 'Drug']]
```

Out[109]:

	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

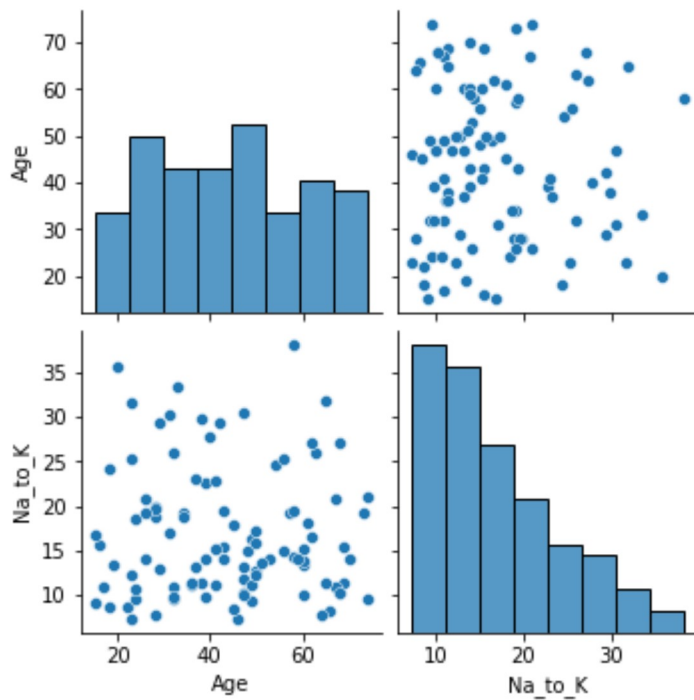
```
In [110]:
```

Out[110]:

	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 [111]:

Out[111]: <seaborn.axisgrid.PairGrid at 0x2602cb536a0>

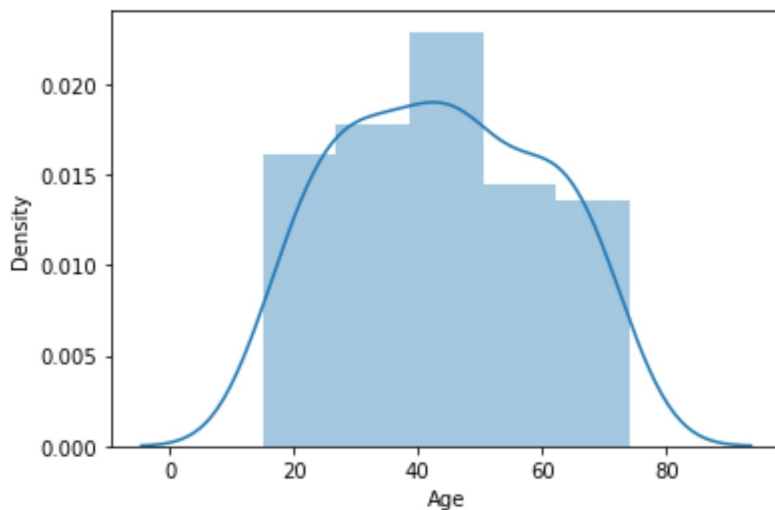


In [113]:

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)

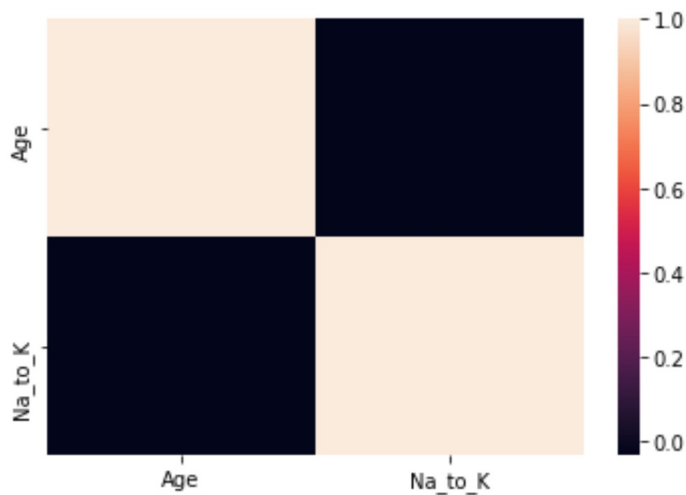
Out[113]: <AxesSubplot:xlabel='Age', ylabel='Density'>



In [114]: x1=x[['Age', 'Sex', 'BP', 'Cholesterol', 'Na_to_K', 'Drug']]

In [115]:

Out[115]: <AxesSubplot:>

In [131]: `x=x1[['Age','Na_to_K']]`In [132]: `# to split my dataset into training and test data``from sklearn.model_selection import train_test_split`In [133]: `from sklearn.linear_model import LinearRegression``lr=LinearRegression()`Out[133]: `LinearRegression()`

In [134]:

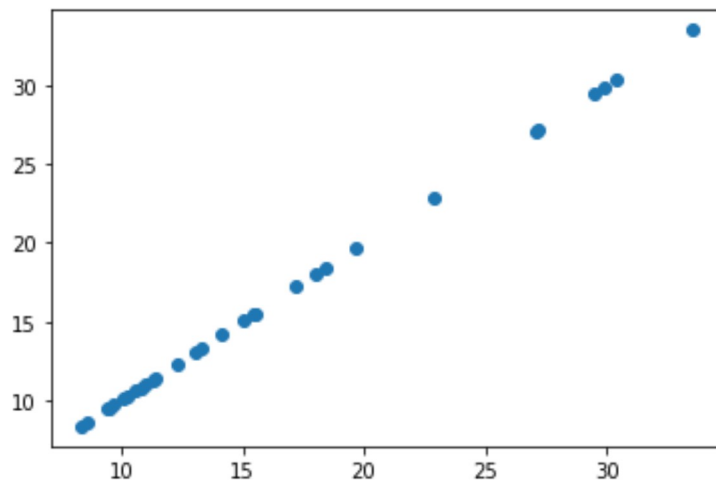
`1.0658141036401503e-14`In [135]: `coeff=pd.DataFrame(lr.coef_,x.columns,columns=['Co-efficient'])`

Out[135]:

	Co-efficient
Age	6.566793e-18
Na_to_K	1.000000e+00

```
In [136]: prediction=lr.predict(x_test)
```

```
Out[136]: <matplotlib.collections.PathCollection at 0x2602cb2a880>
```



```
In [137]:
```

```
Out[137]: 1.0
```

```
In [ ]:
```