D3

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

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

Out[2]:

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

200 rows × 6 columns

In [3]: 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	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				
<pre>dtypes: float64(1), int64(1), object(4)</pre>							

memory usage: 9.5+ KB

In [4]: df.describe()

Out[4]:

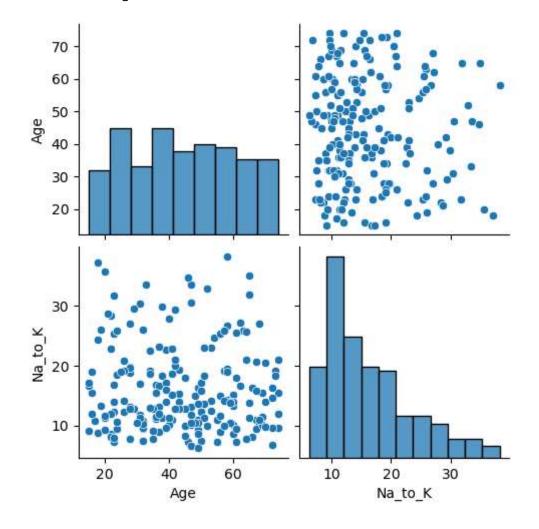
	Age	Na_to_K
count	200.000000	200.000000
mean	44.315000	16.084485
std	16.544315	7.223956
min	15.000000	6.269000
25%	31.000000	10.445500
50%	45.000000	13.936500
75%	58.000000	19.380000
max	74.000000	38.247000

In [5]: df.columns

Out[5]: Index(['Age', 'Sex', 'BP', 'Cholesterol', 'Na_to_K', 'Drug'], dtype='object')

In [6]: sns.pairplot(df)

Out[6]: <seaborn.axisgrid.PairGrid at 0x1eb144434d0>



In [7]: sns.distplot(df["Age"])

C:\Users\user\AppData\Local\Temp\ipykernel_7792\2732350774.py:1: UserWarning:

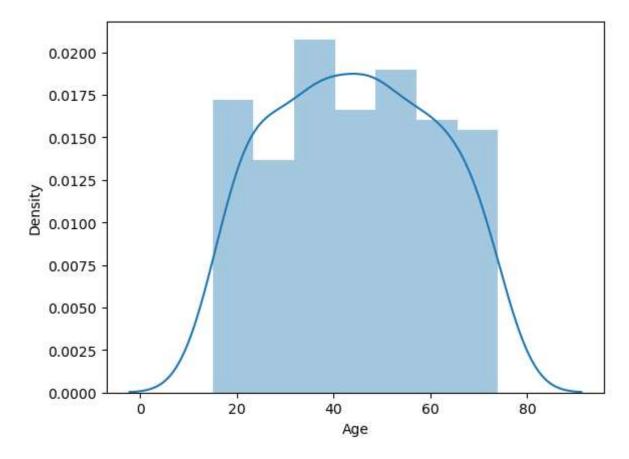
`distplot` is a deprecated function and will be removed in seaborn v0.14.0.

Please adapt your code to use either `displot` (a figure-level function with similar flexibility) or `histplot` (an axes-level function for histograms).

For a guide to updating your code to use the new functions, please see https://gist.github.com/mwaskom/de44147ed2974457ad6372750bbe5751 (https://gist.github.com/mwaskom/de44147ed2974457ad6372750bbe5751)

sns.distplot(df["Age"])

Out[7]: <Axes: xlabel='Age', ylabel='Density'>

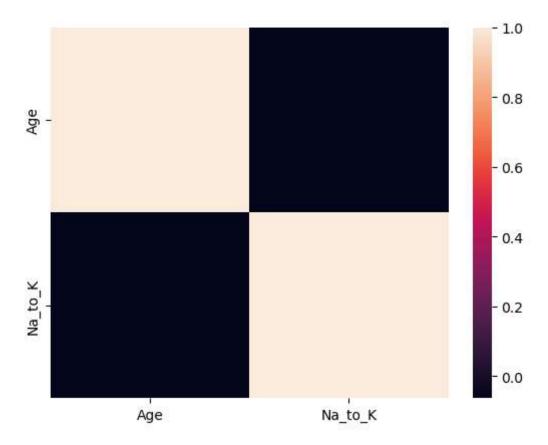


In [8]: df1=df[['Age', 'Sex', 'BP', 'Cholesterol', 'Na_to_K', 'Drug']]

```
In [9]: sns.heatmap(df1.corr())
```

C:\Users\user\AppData\Local\Temp\ipykernel_7792\781785195.py:1: FutureWarnin
g: The default value of numeric_only in DataFrame.corr is deprecated. In a fu
ture version, it will default to False. Select only valid columns or specify
the value of numeric_only to silence this warning.
 sns.heatmap(df1.corr())

Out[9]: <Axes: >



```
In [10]: x=df1[['Age']]
y=df1['Na_to_K']
```

In [11]: 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)

Out[12]: v LinearRegression LinearRegression()

```
In [13]: print(lr.intercept_)
```

15.435320212781702

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

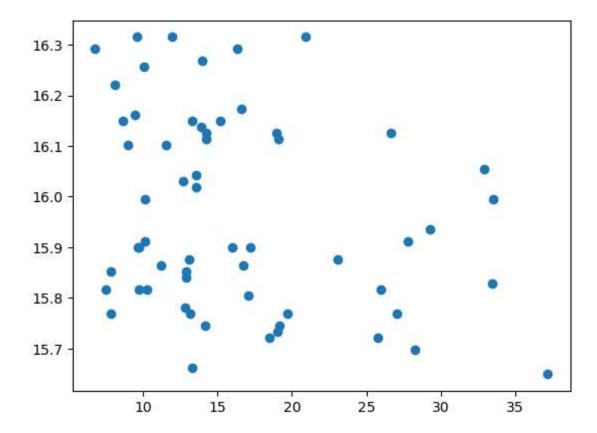
Out[14]:

Co-efficient

Age 0.011889

```
In [15]: prediction=lr.predict(x_test)
    plt.scatter(y_test, prediction)
```

Out[15]: <matplotlib.collections.PathCollection at 0x1eb0f056250>



```
In [16]: print(lr.score(x_test,y_test))
```

-0.017709603394363116

In [17]: from sklearn.linear_model import Ridge,Lasso

In [18]: rr=Ridge(alpha=10)
 rr.fit(x_train,y_train)

Out[18]: Ridge
Ridge(alpha=10)