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```
In [51]: #to import Libraries
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
```

```
In [54]: #to import dataset
data=pd.read_csv(r"C:\Users\user\Downloads\4_drug200 - 4_drug200.csv")
data
```

Out[54]:

	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 [55]: data.head()
```

Out[55]:

	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

DATA CLEANING AND PREPROCESSING

In [56]: data.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 [57]: data.describe()

Out[57]:

	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

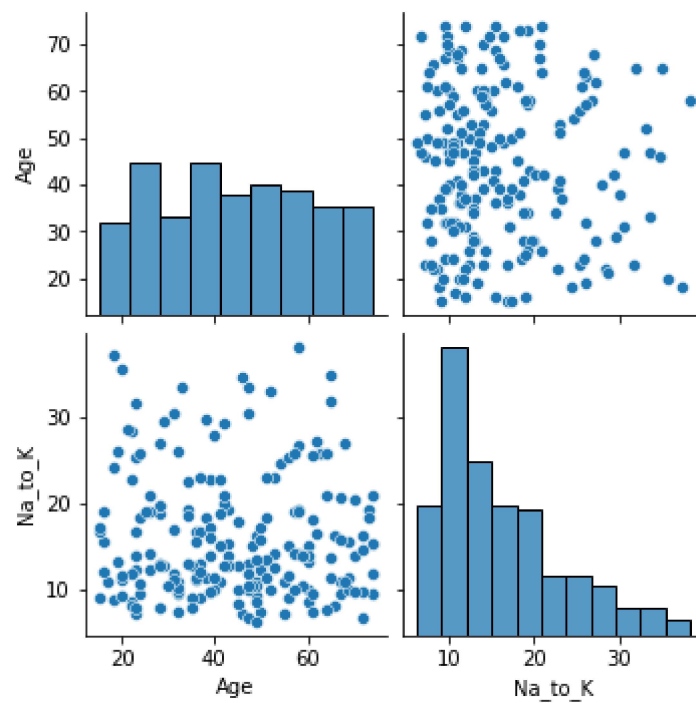
In [58]: data.columns

Out[58]: Index(['Age', 'Sex', 'BP', 'Cholesterol', 'Na_to_K', 'Drug'], dtype='object')

EDA and DATA VISUALIZATION

```
In [59]: sns.pairplot(data)
```

```
Out[59]: <seaborn.axisgrid.PairGrid at 0x241d536b910>
```

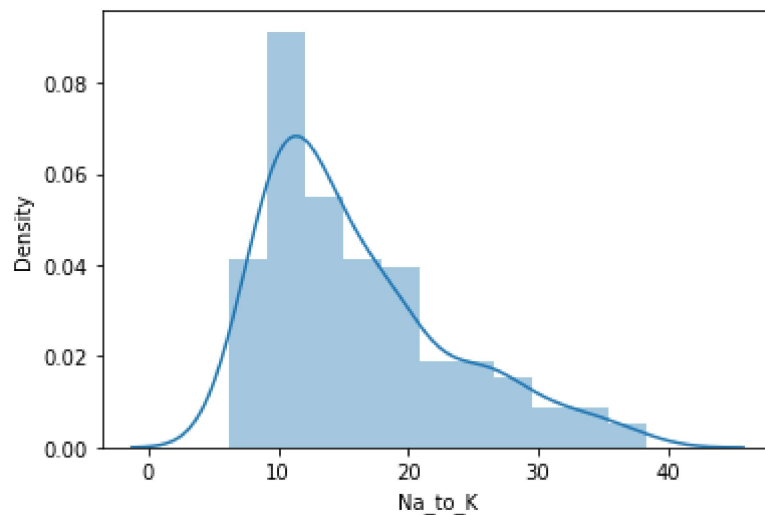


```
In [62]: sns.distplot(data['Na_to_K'])
```

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)
```

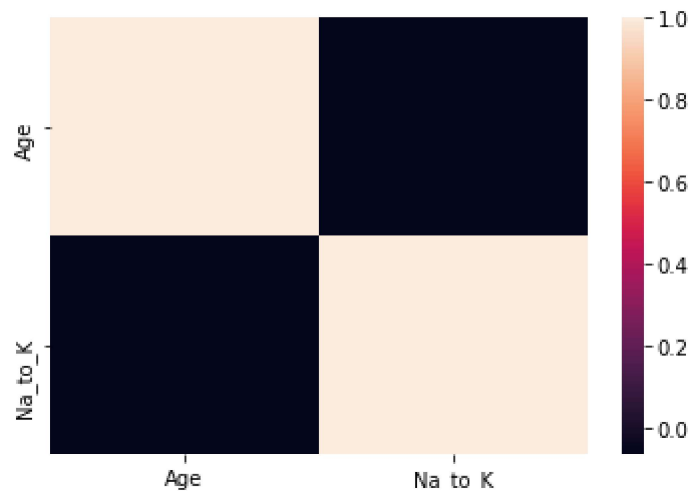
```
Out[62]: <AxesSubplot:xlabel='Na_to_K', ylabel='Density'>
```



```
In [63]: df=data[['Age', 'Sex', 'BP', 'Cholesterol', 'Na_to_K', 'Drug']]
```

```
In [64]: sns.heatmap(df.corr())
```

```
Out[64]: <AxesSubplot:>
```



TRAINING MODEL

```
In [69]: x=df[['Age']]  
y=df[['Na_to_K']]
```

```
In [70]: #to split my dataset into training and test  
  
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 [71]: from sklearn.linear_model import LinearRegression  
lr=LinearRegression()  
lr.fit(x_train,y_train)
```

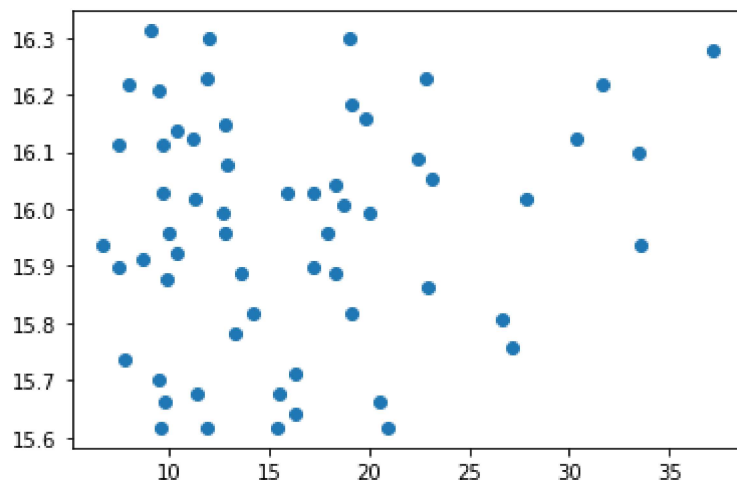
Out[71]: LinearRegression()

```
In [72]: #to find intercept  
print(lr.intercept_)
```

[16.48883429]

```
In [73]: prediction = lr.predict(x_test)  
plt.scatter(y_test,prediction)
```

Out[73]: <matplotlib.collections.PathCollection at 0x241d57f8880>



```
In [74]: print(lr.score(x_test,y_test))
```

0.0042465716188196945

RIDGE AND LASSO REGRESSION

```
In [75]: from sklearn.linear_model import Ridge,Lasso
```

```
In [76]: rr=Ridge(alpha=10)
         rr.fit(x_train,y_train)
```

Out[76]: Ridge(alpha=10)

```
In [77]: rr.score(x_test,y_test)
```

Out[77]: 0.0042448844866279645

```
In [78]: la=Lasso(alpha=10)
         la.fit(x_train,y_train)
```

Out[78]: Lasso(alpha=10)

```
In [79]: la.score(x_test,y_test)
```

Out[79]: -0.0027678708157250487