

Importing libraries

```
In [1]: import pandas as pd
import numpy as np
from numpy import linalg as la
from numpy import cov
```

Importing dataset

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

```
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

a.mean

```
In [3]: print(data.mean())
```

```
Age      44.315000
Na_to_K   16.084485
dtype: float64
```

median

```
In [4]: print(data.median())
```

```
Age          45.0000
Na_to_K      13.9365
dtype: float64
```

mode

In [5]:

print(data.mode())

```
   Age  Sex  BP Cholesterol  Na_to_K  Drug
0  47.0   M  HIGH          HIGH  12.006  drugY
1   NaN  NaN  NaN          NaN  18.295   NaN
```

describe

In [6]:

print(data.describe())

```
count      Age      Na_to_K
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
```

b.sum()

In [7]:

print(data.sum())

```
Age          8863
Sex          FMMFF...
BP          HIGHLOW...
Cholesterol  HIGHHIGH...
Na_to_K      3216.897
Drug        drugYdrugC...
dtype: object
```

cumsum()

In [8]:

print(data.cumsum())

```
   Age  Sex \
0    23   F
1    70  FM
2   117 FMM
3   145 FMMF
4   206 FMMFF
..   ...
195  8732 FMMFF...
196  8748 FMMFF...
197  8800 FMMFF...
198  8823 FMMFF...
199  8863 FMMFF...
```

		BP	\
0		HIGH	
1		HIGHLOW	
2		HIGHLOWLOW	
3		HIGHLOWLOWNORMAL	
4		HIGHLOWLOWNORMALLOW	
..		...	
195	HIGHLOWLOWNORMALLOWNORMALNORMALLOWNORMALLOWLOW...		
196	HIGHLOWLOWNORMALLOWNORMALNORMALLOWNORMALLOWLOW...		
197	HIGHLOWLOWNORMALLOWNORMALNORMALLOWNORMALLOWLOW...		
198	HIGHLOWLOWNORMALLOWNORMALNORMALLOWNORMALLOWLOW...		
199	HIGHLOWLOWNORMALLOWNORMALNORMALLOWNORMALLOWLOW...		
		Cholesterol	Na_to_K \
0		HIGH	25.355
1		HIGHHIGH	38.448
2		HIGHHIGHHIGH	48.562
3		HIGHHIGHHIGHHIGH	56.360
4		HIGHHIGHHIGHHIGHHIGH	74.403
..	
195	HIGHHIGHHIGHHIGHHIGHHIGHHIGHHIGHHIGHHIGHHIGHNORMALHIGH...		3169.628
196	HIGHHIGHHIGHHIGHHIGHHIGHHIGHHIGHHIGHHIGHHIGHNORMALHIGH...		3181.634
197	HIGHHIGHHIGHHIGHHIGHHIGHHIGHHIGHHIGHHIGHHIGHNORMALHIGH...		3191.528
198	HIGHHIGHHIGHHIGHHIGHHIGHHIGHHIGHHIGHHIGHHIGHNORMALHIGH...		3205.548
199	HIGHHIGHHIGHHIGHHIGHHIGHHIGHHIGHHIGHHIGHHIGHNORMALHIGH...		3216.897
		Drug	
0		drugY	
1		drugYdrugC	
2		drugYdrugCdrugC	
3		drugYdrugCdrugCdrugX	
4		drugYdrugCdrugCdrugXdrugY	
..		...	
195	drugYdrugCdrugCdrugXdrugYdrugXdrugYdrugCdrugYd...		
196	drugYdrugCdrugCdrugXdrugYdrugXdrugYdrugCdrugYd...		
197	drugYdrugCdrugCdrugXdrugYdrugXdrugYdrugCdrugYd...		
198	drugYdrugCdrugCdrugXdrugYdrugXdrugYdrugCdrugYd...		
199	drugYdrugCdrugCdrugXdrugYdrugXdrugYdrugCdrugYd...		

[200 rows x 6 columns]

count

In [9]:

print(data.count())

Age 200
Sex 200
BP 200
Cholesterol 200
Na_to_K 200
Drug 200
dtype: int64

min

In [10]:

print(data.min())

```
Age          15
Sex          F
BP           HIGH
Cholesterol  HIGH
Na_to_K      6.269
Drug         drugA
dtype: object
```

max()

```
In [11]: print(data.max())
```

```
Age          74
Sex          M
BP           NORMAL
Cholesterol  NORMAL
Na_to_K      38.247
Drug         drugY
dtype: object
```

covariance

```
In [12]: print(data.cov())
```

```
          Age  Na_to_K
Age    273.714347 -7.543752
Na_to_K -7.543752  52.185533
```

corelation

pearson corelation

```
In [14]: from scipy.stats import pearsonr
print(pearsonr(data["Age"],data["Na_to_K"]))

(-0.06311949726772592, 0.3745756399034559)
```

spearman corelation

```
In [15]: from scipy.stats import spearmanr
print(spearmanr(data["Age"],data["Na_to_K"]))

SpearmanrResult(correlation=-0.047273882688479915, pvalue=0.5062200581387418)
```

```
In [ ]:
```