

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

5 Data Sets

```
In [2]: a=pd.read_csv("drug.csv")
a
```

```
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) Find mean, median, mode and describe

```
In [3]: print(a.mean)
print("Median:")
print(a.median)
print("Mode:")
print(a.mode)
print("Describe")
print(a.describe())
```

```
<bound method NDFrame._add_numeric_operations.<locals>.mean of
Cholesterol Na_to_K Drug Age Sex BP
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 x 6 columns]>
Median:
<bound method NDFrame._add_numeric_operations.<locals>.median of          Age Sex      B
P 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 x 6 columns]>
Mode:
<bound method DataFrame.mode of          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 x 6 columns]>
Describe
      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
```

b) Find sum(), cumsum(), count, min and max values

In [4]:

print(a.sum())

Age 8863
Sex FMMFFFMFFMFFMFFMFFMFFMFFMFFMFFMFFMFFMFF...
BP HIGHLOWLOWNORMALLOWNORMALNORMALLOWNORMALLOW...
Cholesterol HIGHHHIGHHHIGHHHIGHHHIGHHHIGHHHIGHHIGNORMALHIGH...
Na_to_K 3216.897
Drug drugYdrugCdrugCdrugXdrugYdrugXdrugYdrugCdrugYd...
dtype: object

In [5]:

print(a.cumsum())

Age Sex \
0 23 F
1 70 FM
2 117 FMM
3 145 FMMF
4 206 FMFFF

```
[200 rows x 6 columns]
```

```
print(a.count())
print(a.min())
print(a.max())
```

[convert/html/DATASET 2.ipynb?download=false](#)

```

BP          NORMAL
Cholesterol NORMAL
Na_to_K     38.247
Drug        drugY
dtype: object

```

c) Find covariance and correlation (spearman and pearsons)

In [7]:

```
print(a.cov())
```

```

           Age  Na_to_K
Age    273.714347 -7.543752
Na_to_K -7.543752  52.185533

```

In [8]:

```
b=a.dropna(axis=1,how='any')
b
```

Out[8]:

	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 [11]:

```
from scipy.stats import pearsonr
print(pearsonr(a['Age'],a['Na_to_K']))
```

(-0.06311949726772592, 0.3745756399034559)

In [13]:

```
from scipy.stats import spearmanr
print(spearmanr(a['Age'],a['Na_to_K']))
```

SpearmanrResult(correlation=-0.047273882688479915, pvalue=0.5062200581387418)

In []: