1.IMPORTING LIBRARIES

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

2.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

3.head

```
In [3]:    data.head(8)
```

Out[3]:

	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

	Age	Sex	BP	Cholesterol	Na_to_K	Drug
5	22	F	NORMAL	HIGH	8.607	drugX
6	49	F	NORMAL	HIGH	16.275	drugY
7	41	M	LOW	HIGH	11.037	drugC

4.tail

In [4]:
```
data.tail(7)
```

Out[4]:

	Age	Sex	BP	Cholesterol	Na_to_K	Drug
193	72	M	LOW	HIGH	6.769	drugC
194	46	F	HIGH	HIGH	34.686	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

5.describe()

In [5]:
```
data.describe()
```

Out[5]:

	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

6.shape()

In [6]:
```
np.shape(data)
```

Out[6]: (200, 6)

7.size()

In [7]:
```python
np.size(data)
```

Out[7]: 1200

8.isna()

In [8]:
```python
data.isna()
```

Out[8]:

	Age	Sex	BP	Cholesterol	Na_to_K	Drug
0	False	False	False	False	False	False
1	False	False	False	False	False	False
2	False	False	False	False	False	False
3	False	False	False	False	False	False
4	False	False	False	False	False	False
...	...	...	...	...	...	...
195	False	False	False	False	False	False
196	False	False	False	False	False	False
197	False	False	False	False	False	False
198	False	False	False	False	False	False
199	False	False	False	False	False	False

200 rows × 6 columns

9.dropna

In [9]:
```python
data.dropna()
```

Out[9]:

	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

	Age	Sex	BP	Cholesterol	Na_to_K	Drug
...	...	...	...	...	...	...
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

10.selecting specific column

In [10]:
```python
da=data[["Age","Sex"]]
da
```

Out[10]:

	Age	Sex
0	23	F
1	47	M
2	47	M
3	28	F
4	61	F
...	...	...
195	56	F
196	16	M
197	52	M
198	23	M
199	40	F

200 rows × 2 columns

11.line plot

In [11]:
```python
data.plot.line()
```

Out[11]: <AxesSubplot:>
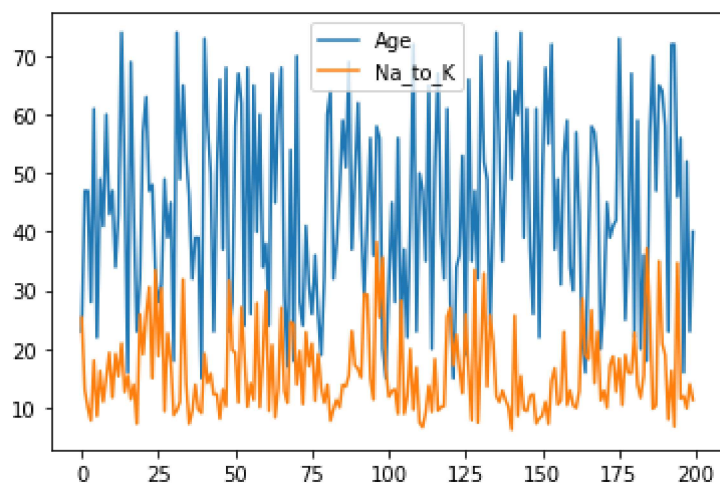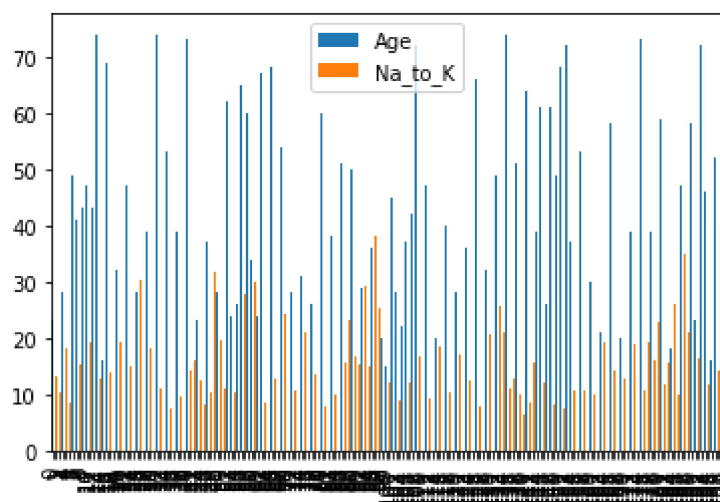
12.bar plot
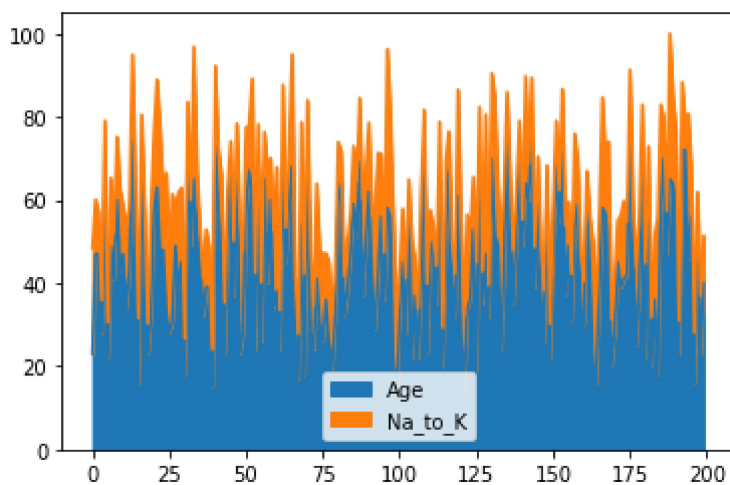
In [12]:
```
data.plot.bar()
```

Out[12]:   <AxesSubplot:>

13.area plot

In [13]:
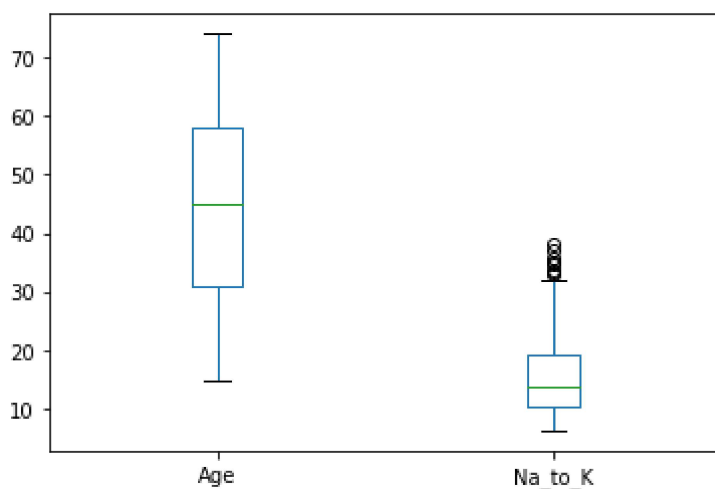```
data.plot.area()
```

Out[13]:    <AxesSubplot:>
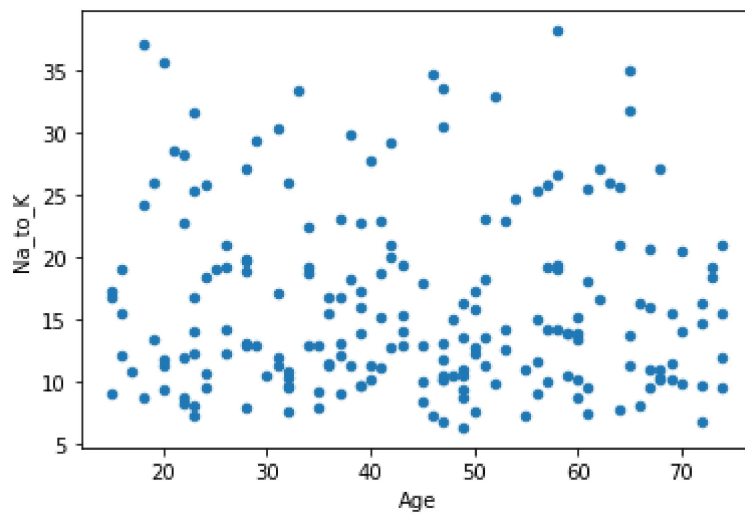
# 14.box plot

```
In [14]:   data.plot.box()
```

Out[14]:   <AxesSubplot:>

15.scatter plot
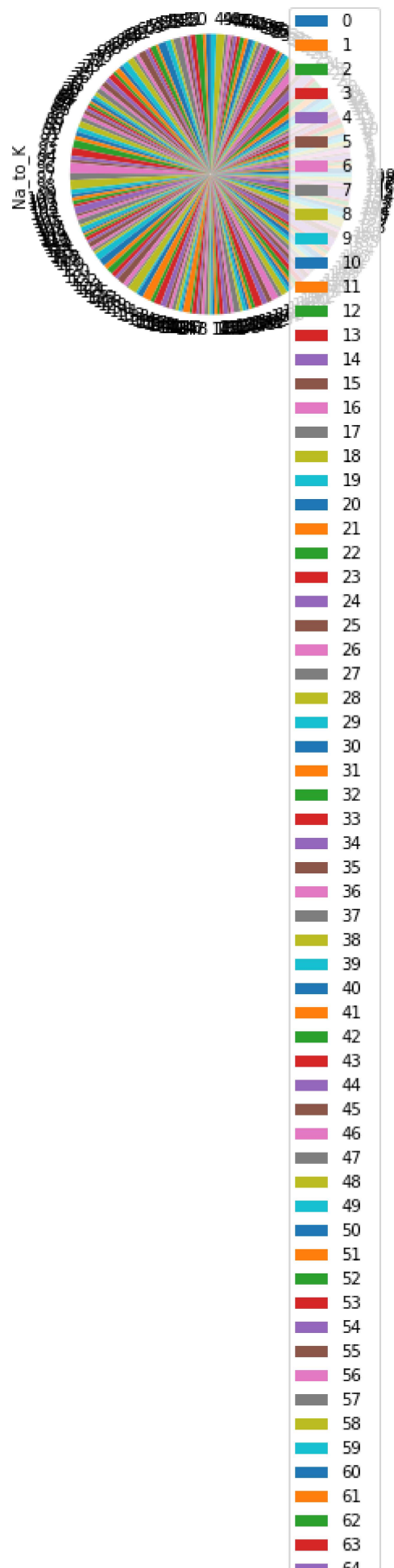
```
In [15]:   data.plot.scatter("Age","Na_to_K")
```
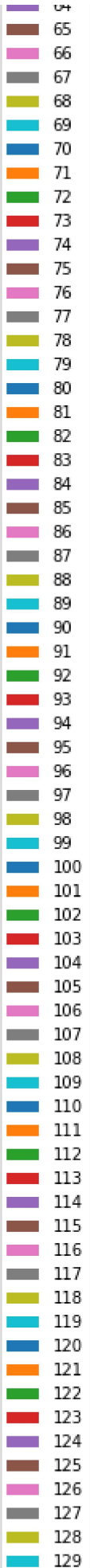
Out[15]:   <AxesSubplot:xlabel='Age', ylabel='Na_to_K'>

16.pie plot

In [17]:
```python
data.plot.pie(x="Saves",y="Na_to_K")
```

Out[17]: `<AxesSubplot:ylabel='Na_to_K'>`

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17.histogram

In [18]:
```python
data.plot.hist()
```

Out[18]: <AxesSubplot:ylabel='Frequency'>

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