

```
import pandas as pd
d=pd.read_csv('/content/lungcancer.csv')
print(d)
```

```
index Patient Id Age Gender Air Pollution Alcohol use Dust Allergy \
0 0 P1 33 1 2 4 5
1 1 P10 17 1 3 1 5
2 2 P100 35 1 4 5 6
3 3 P1000 37 1 7 7 7
4 4 P101 46 1 6 8 7
.. ...
995 995 P995 44 1 6 7 7
996 996 P996 37 2 6 8 7
997 997 P997 25 2 4 5 6
998 998 P998 18 2 6 8 7
999 999 P999 47 1 6 5 6

OccuPational Hazards Genetic Risk chronic Lung Disease ... Fatigue \
0 4 3 2 ... 3
1 3 4 2 ... 1
2 5 5 4 ... 8
3 7 6 7 ... 4
4 7 7 6 ... 3
.. ...
995 7 7 6 ... 5
996 7 7 6 ... 9
997 5 5 4 ... 8
998 7 7 6 ... 3
999 5 5 4 ... 8

Weight Loss Shortness of Breath Wheezing Swal0ing Difficulty \
0 4 2 2 3
1 3 7 8 6
2 7 9 2 1
3 2 3 1 4
4 2 4 1 4
.. ...
995 3 2 7 8
996 6 5 7 2
997 7 9 2 1
998 2 4 1 4
999 7 9 2 1

Clubbing of Finger Nails Frequent Cold Dry Cough Snoring Level
0 1 2 3 4 0
1 2 1 7 2 1
2 4 6 7 2 2
3 5 6 7 5 2
4 2 4 2 3 2
.. ...
995 2 4 5 3 2
996 4 3 1 4 2
997 4 6 7 2 2
998 2 4 2 3 2
999 4 6 7 2 2
```

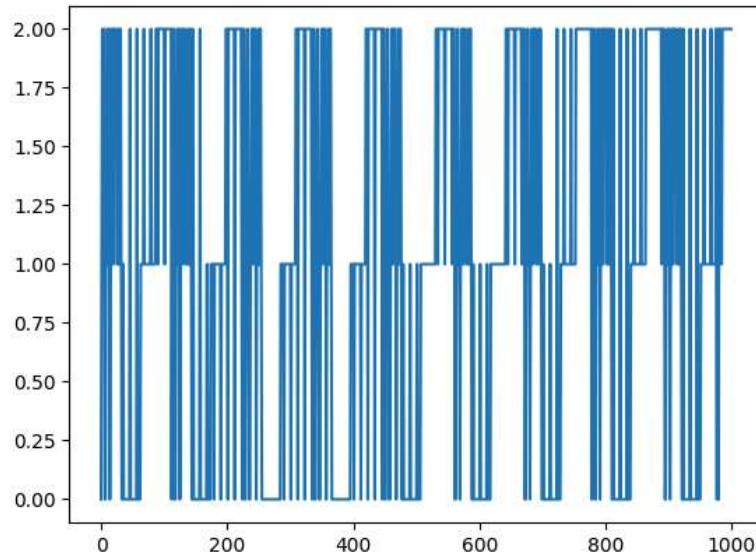
[1000 rows x 26 columns]

```
from google.colab import drive
drive.mount('/content/drive')

↳ Drive already mounted at /content/drive; to attempt to forcibly remount, call drive.mount("/content/drive", force_remount=True).
```

```
from matplotlib import pyplot as pt
x=d['index']
y=d['Level']
pt.plot(x,y)
```

↳ [`<matplotlib.lines.Line2D at 0x798c0fcfa2c20>`]

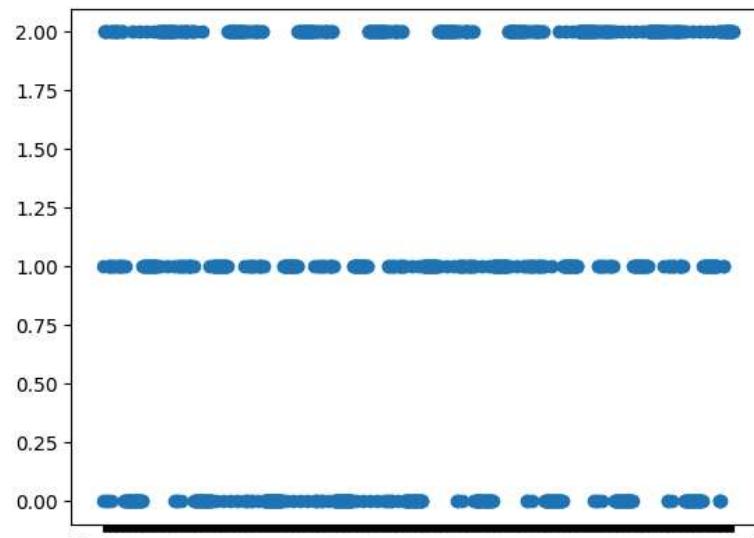


```
pt.scatter(x,y)
```

```
from matplotlib import pyplot as pt
x=d['Patient Id']
y=d['Level']
pt.plot(x,y)
```

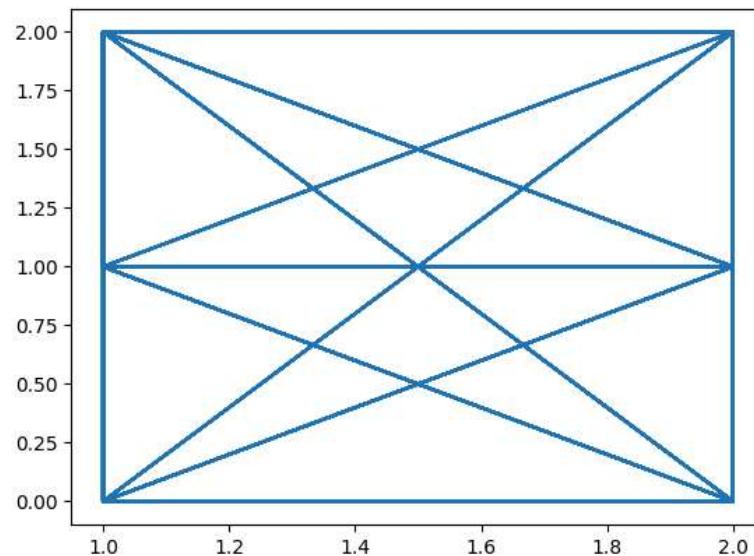
```
pt.scatter(x,y)
```

```
[ <matplotlib.collections.PathCollection at 0x798c245633d0>]
```



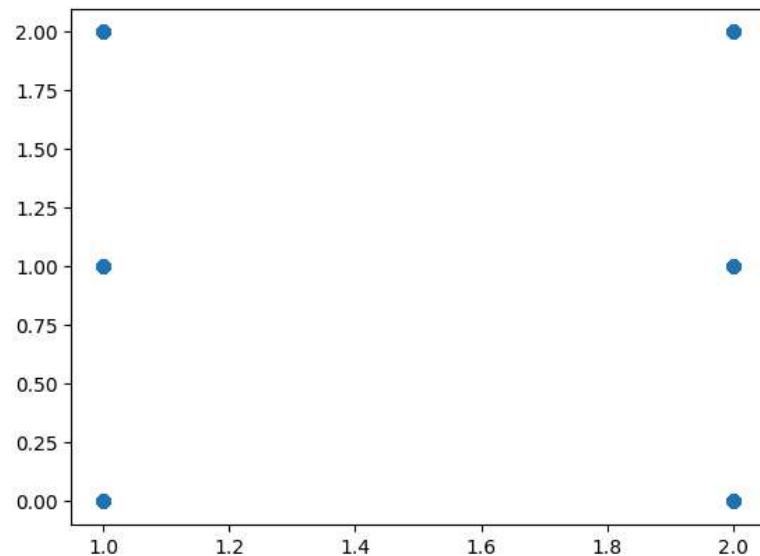
```
from matplotlib import pyplot as pt  
x=d['Gender']  
y=d['Level']  
pt.plot(x,y)
```

```
[ <matplotlib.lines.Line2D at 0x798c14fe2230>]
```



```
pt.scatter(x,y)
```

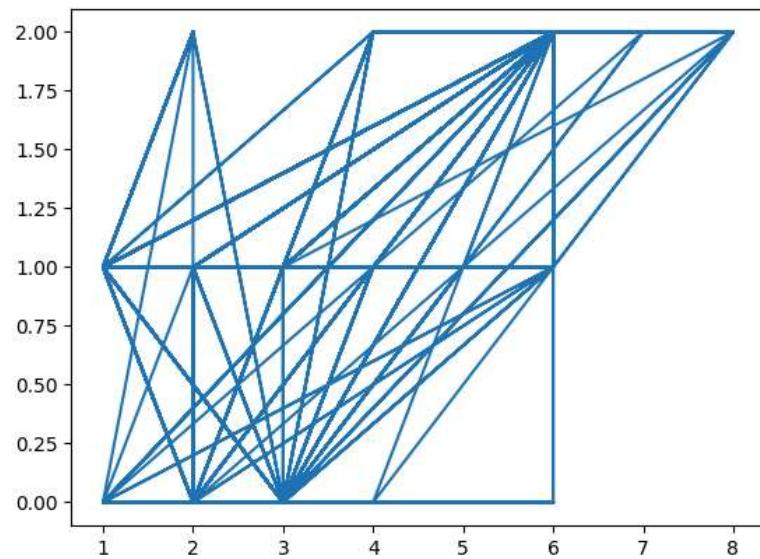
⤵ <matplotlib.collections.PathCollection at 0x798c0cc2cf0>



Start coding or [generate](#) with AI.

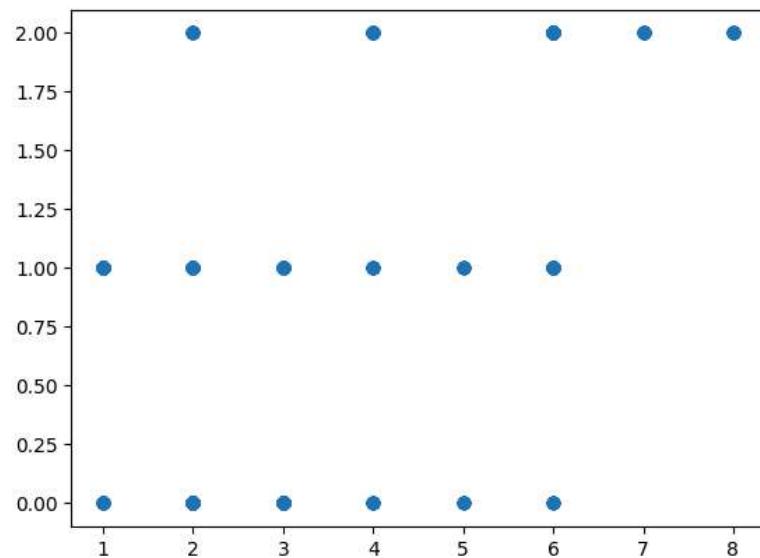
```
from matplotlib import pyplot as pt
x=d['Air Pollution']
y=d['Level']
pt.plot(x,y)
```

⤵ [<matplotlib.lines.Line2D at 0x798c0cab4a90>]



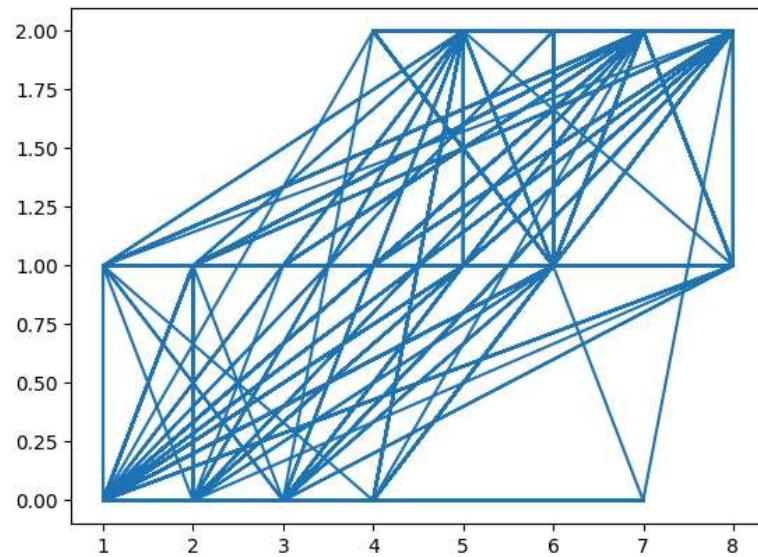
```
pt.scatter(x,y)
```

```
→ <matplotlib.collections.PathCollection at 0x798c0cb20a60>
```



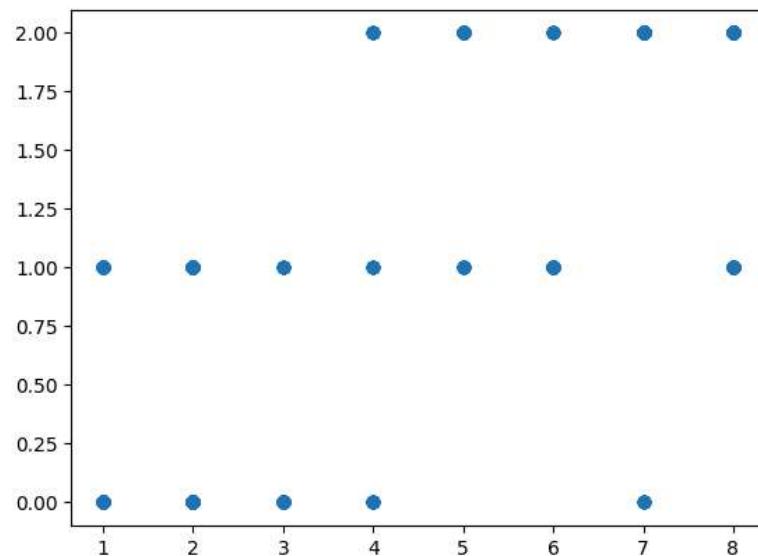
```
from matplotlib import pyplot as pt
x=d['Alcohol use']
y=d['Level']
pt.plot(x,y)
```

```
→ [matplotlib.lines.Line2D at 0x798c0c99ea10]
```



```
pt.scatter(x,y)
```

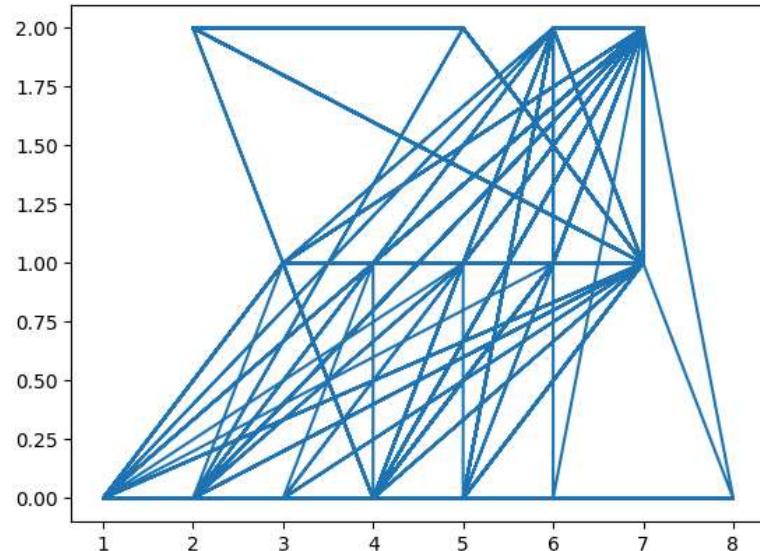
```
→ <matplotlib.collections.PathCollection at 0x798c0ca178e0>
```



```
from matplotlib import pyplot as pt  
x=d['Dust Allergy']
```

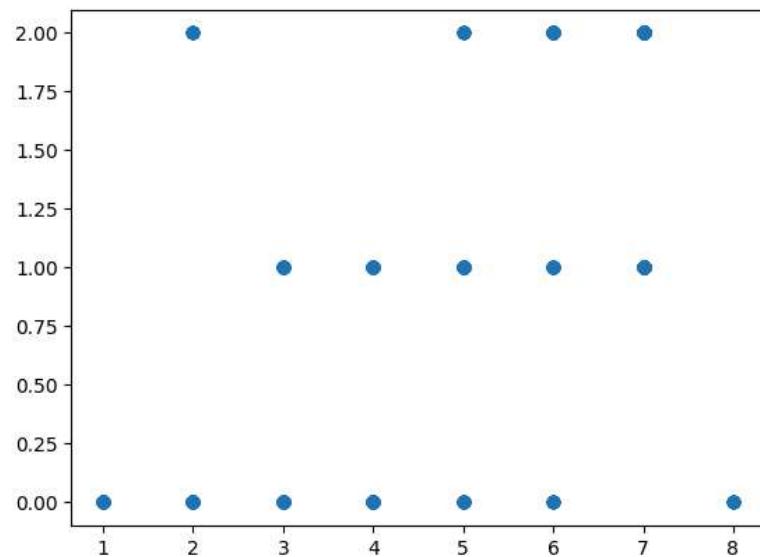
```
y=d['Level']  
pt.plot(x,y)
```

↳ [matplotlib.lines.Line2D at 0x798c0c8b8b20]



```
pt.scatter(x,y)
```

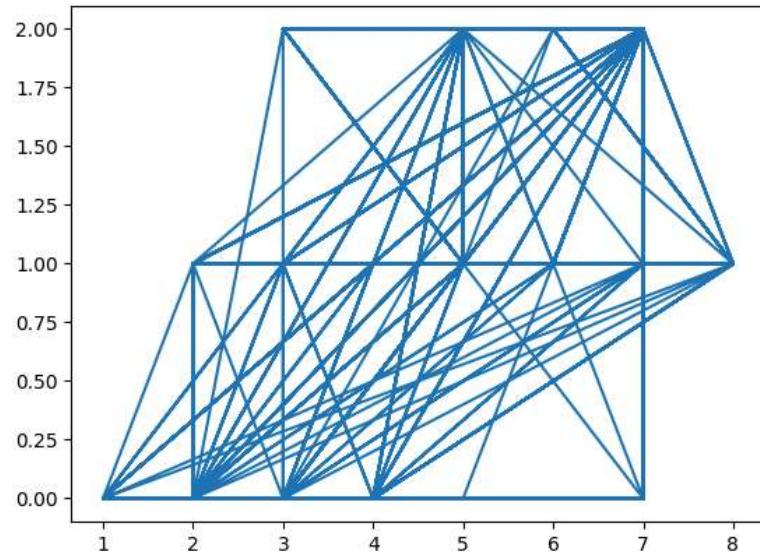
↳ [matplotlib.collections.PathCollection at 0x798c0c935330]



```
from matplotlib import pyplot as pt  
x=d['Occupational Hazards']
```

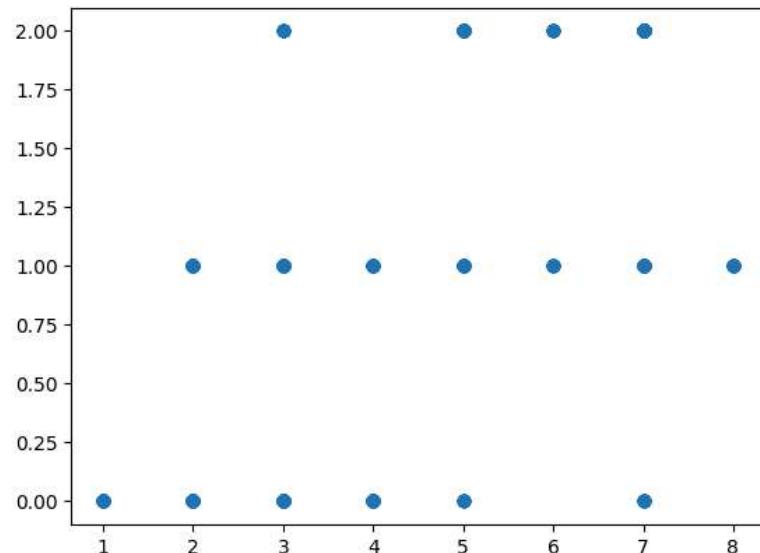
```
y=d['Level']  
pt.plot(x,y)
```

⤵ [matplotlib.lines.Line2D at 0x798c0c79eb30]



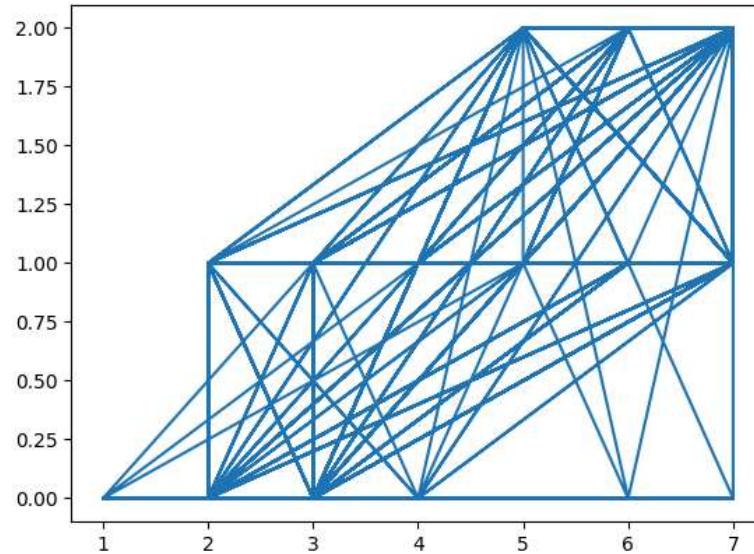
```
pt.scatter(x,y)
```

⤵ <matplotlib.collections.PathCollection at 0x798c0c81b160>



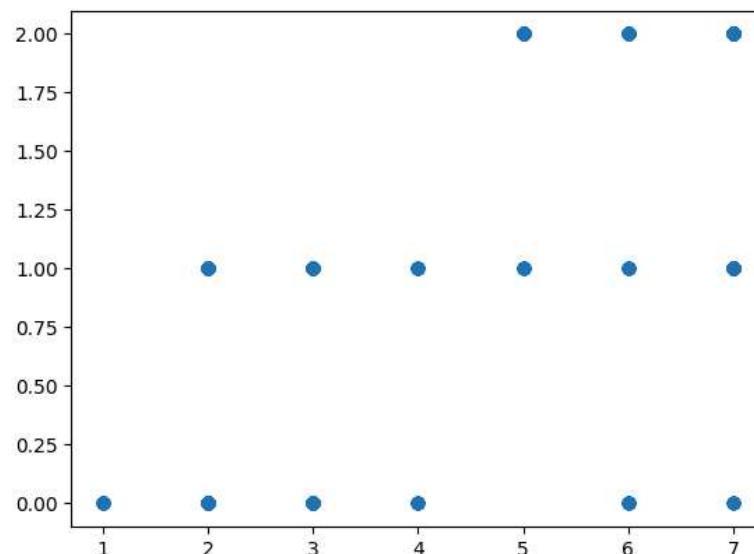
```
from matplotlib import pyplot as pt
x=d['Genetic Risk']
y=d['Level']
pt.plot(x,y)
```

→ [matplotlib.lines.Line2D at 0x798c0c6bc940]



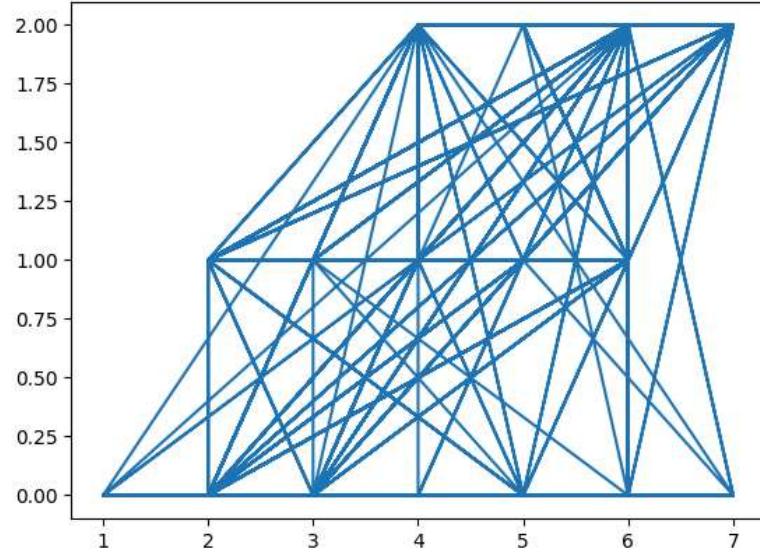
```
pt.scatter(x,y)
```

→ <matplotlib.collections.PathCollection at 0x798c0c53cc70>



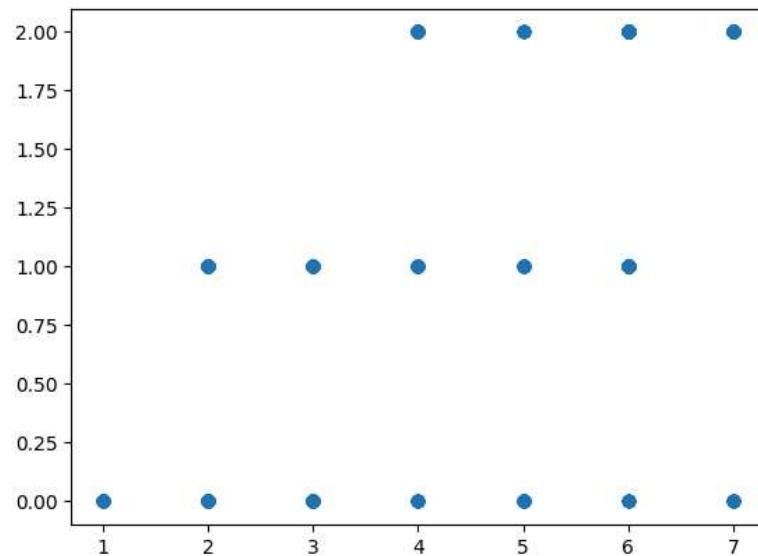
```
from matplotlib import pyplot as pt
x=d['chronic Lung Disease']
y=d['Level']
pt.plot(x,y)
```

→ [matplotlib.lines.Line2D at 0x798c0c6e5ea0]



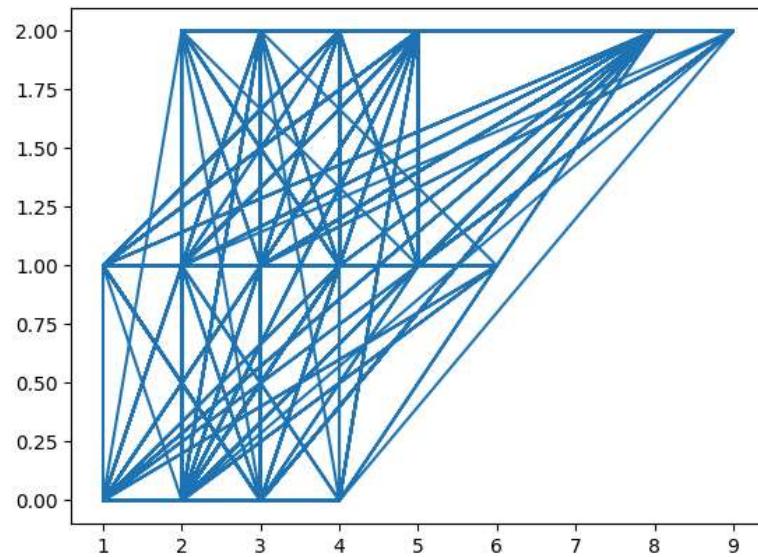
```
pt.scatter(x,y)
```

```
↳ <matplotlib.collections.PathCollection at 0x798c0c635e10>
```



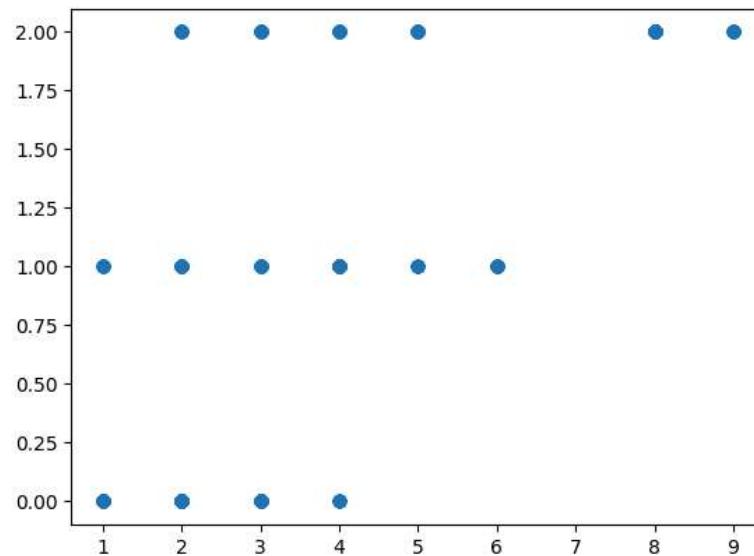
```
from matplotlib import pyplot as pt
x=d['Fatigue']
y=d['Level']
pt.plot(x,y)
```

```
↳ [<matplotlib.lines.Line2D at 0x798c0c4b2a70>]
```



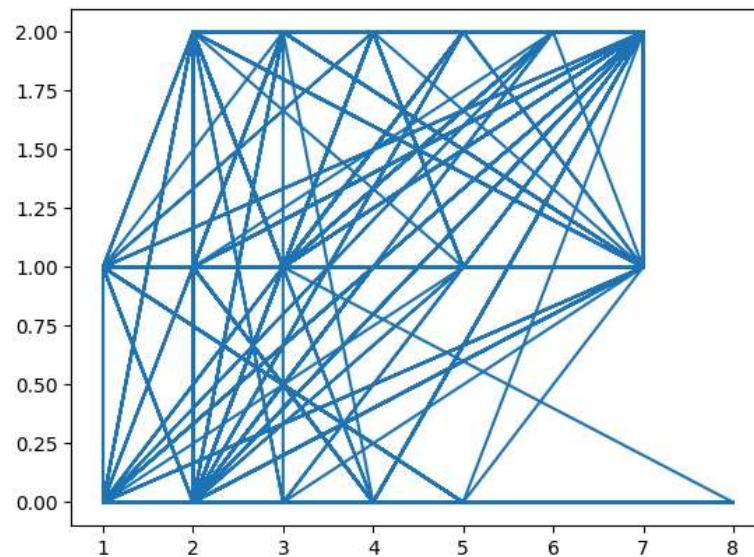
```
pt.scatter(x,y)
```

```
↳ <matplotlib.collections.PathCollection at 0x798c0c3542b0>
```



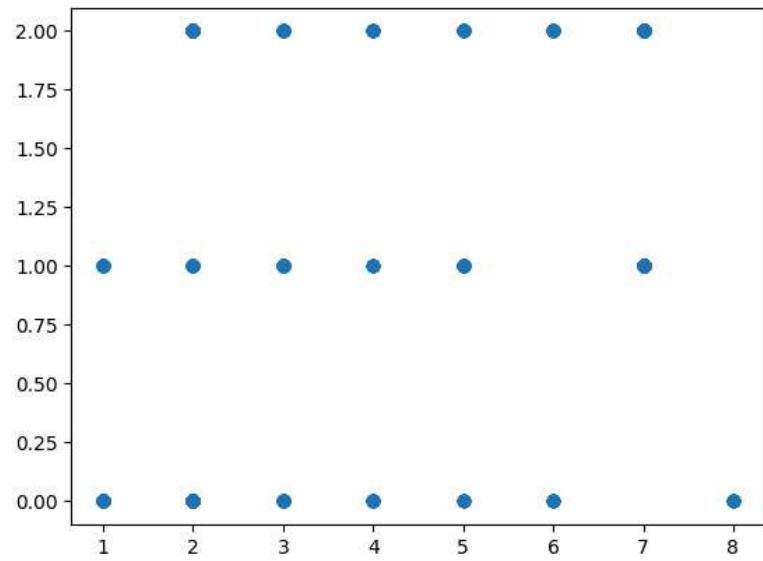
```
from matplotlib import pyplot as pt
x=d['Weight Loss']
y=d['Level']
pt.plot(x,y)
```

```
↳ [<matplotlib.lines.Line2D at 0x798c0c3c57b0>]
```



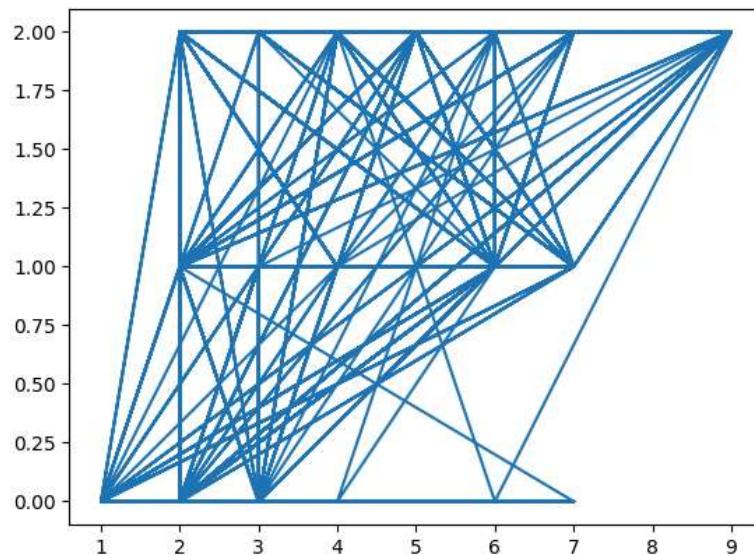
```
pt.scatter(x,y)
```

```
↳ <matplotlib.collections.PathCollection at 0x798c0c3c5930>
```



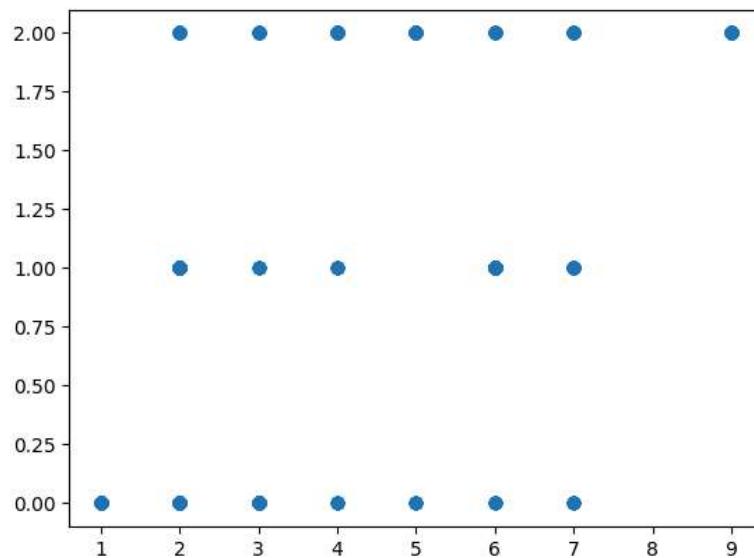
```
from matplotlib import pyplot as pt
x=d['Shortness of Breath']
y=d['Level']
pt.plot(x,y)
```

```
pt.plot(x,y)
```



```
pt.scatter(x,y)
```

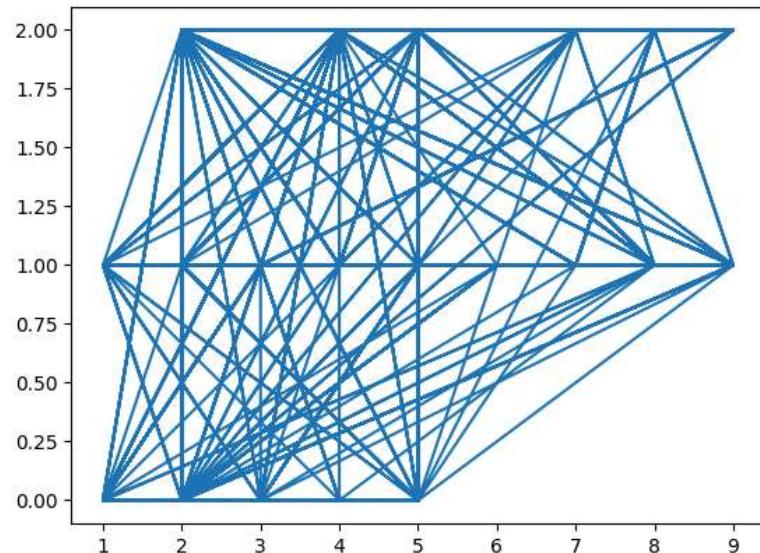
```
pt.plot(x,y)
```



```
from matplotlib import pyplot as pt  
x=d['Clubbing of Finger Nails']
```

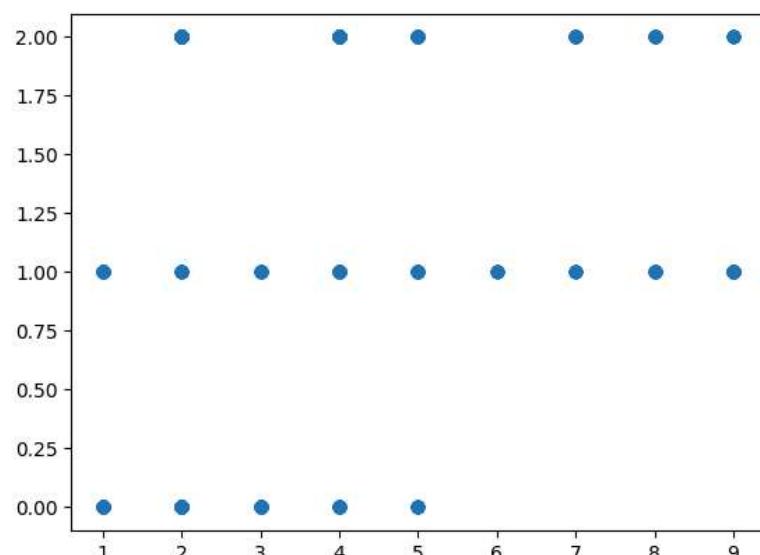
```
y=d['Level']  
pt.plot(x,y)
```

↳ [matplotlib.lines.Line2D at 0x798c0c1dba60]



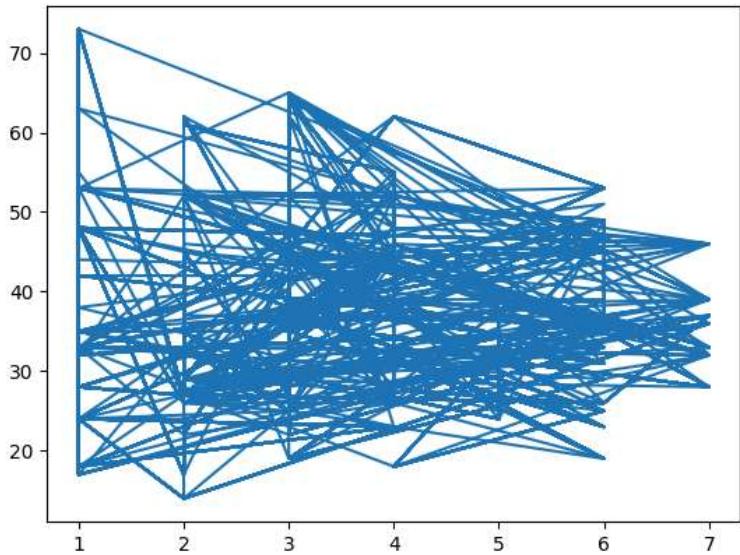
```
pt.scatter(x,y)
```

↳ <matplotlib.collections.PathCollection at 0x798c0c0847c0>



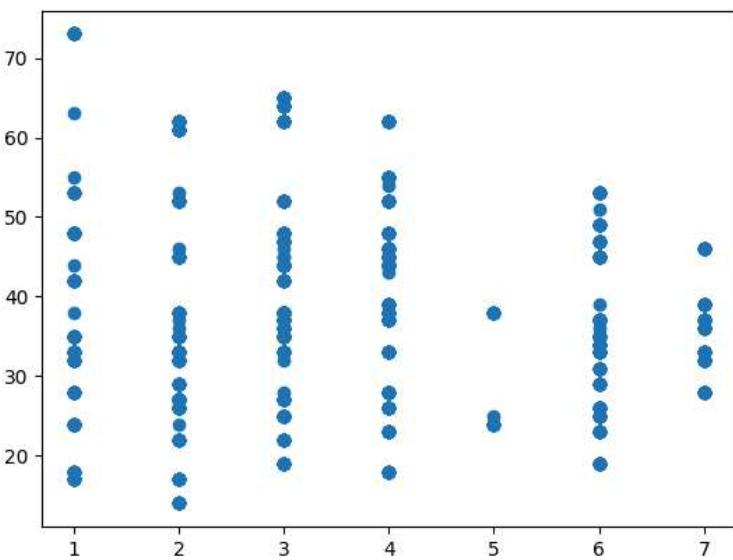
```
from matplotlib import pyplot as pt
x=d['Frequent Cold']
y=d['Age']
pt.plot(x,y)
```

[<matplotlib.lines.Line2D at 0x798c0c0fe140>]



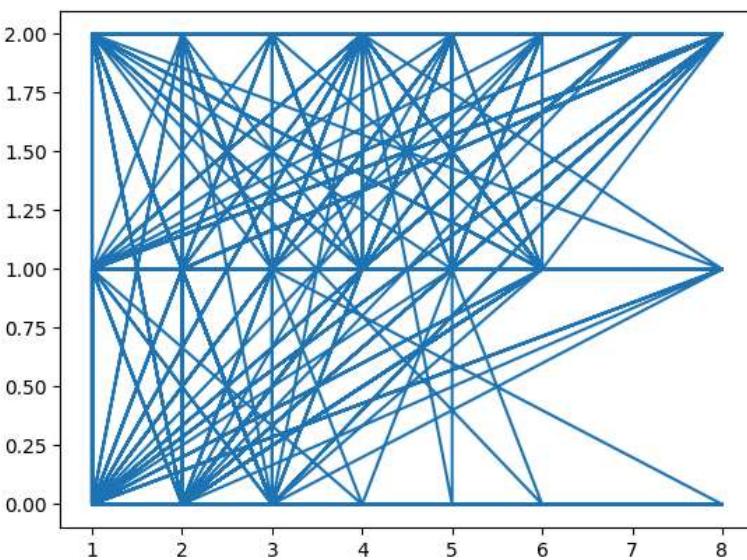
```
pt.scatter(x,y)
```

```
⤵ <matplotlib.collections.PathCollection at 0x798c07f3d840>
```



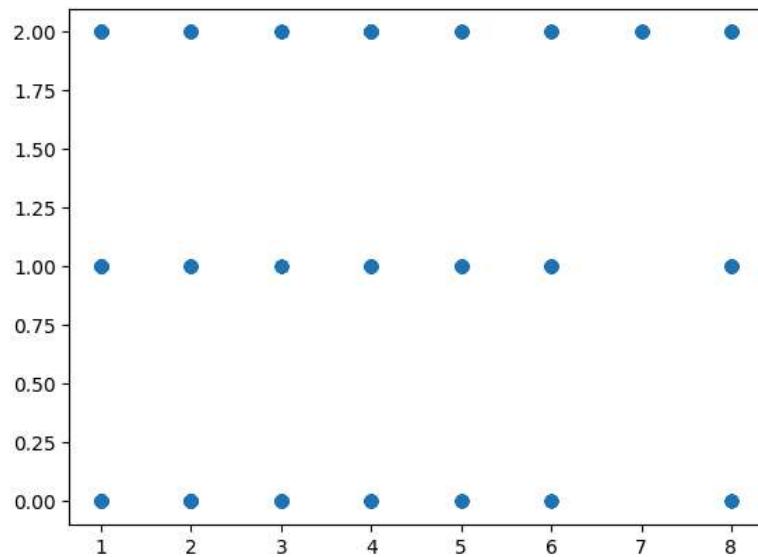
```
from matplotlib import pyplot as pt
x=d['Swallowing Difficulty']
y=d['Level']
pt.plot(x,y)
```

```
⤵ [<matplotlib.lines.Line2D at 0x798c07fb8d30>]
```



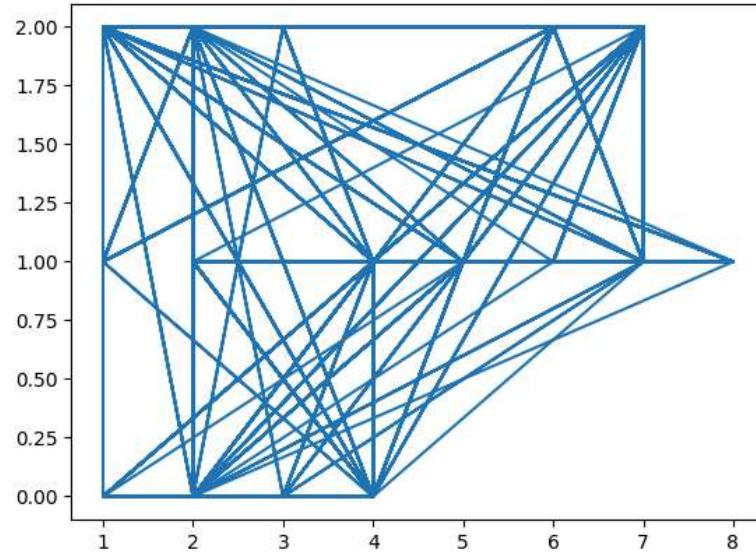
```
pt.scatter(x,y)
```

```
[-] <matplotlib.collections.PathCollection at 0x798c07e2dc30>
```



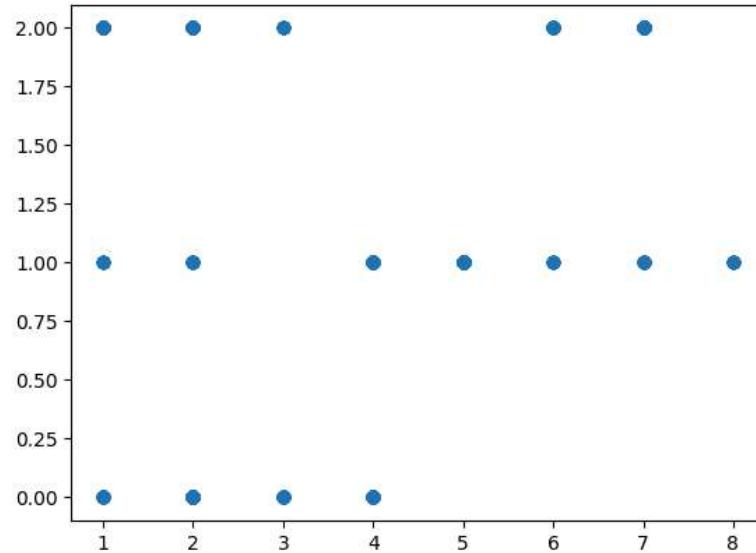
```
from matplotlib import pyplot as pt
x=d['Wheezing']
y=d['Level']
pt.plot(x,y)
```

```
↳ [ <matplotlib.lines.Line2D at 0x798c07eaea40> ]
```



```
pt.scatter(x,y)
```

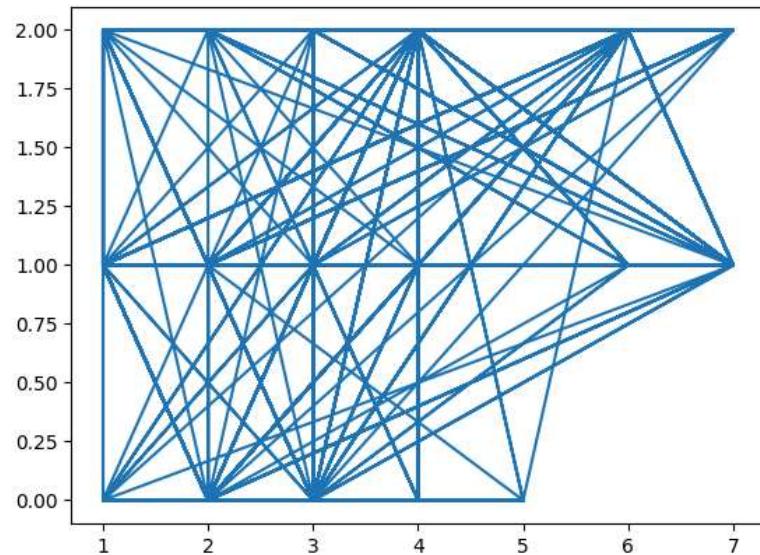
```
↳ <matplotlib.collections.PathCollection at 0x798c07d2fa00>
```



```
from matplotlib import pyplot as pt  
x=d['Frequent Cold']
```

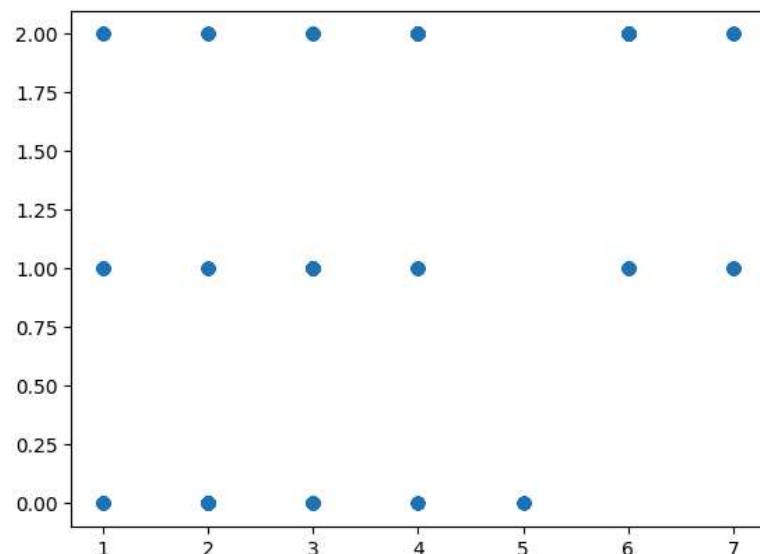
```
y=d['Level']  
pt.plot(x,y)
```

↳ <matplotlib.lines.Line2D at 0x798c07dd0df0>



```
pt.scatter(x,y)
```

↳ <matplotlib.collections.PathCollection at 0x798c07c3cca0>



```
print(d.isnull())
```

	index	Patient Id	Age	Gender	Air Pollution	Alcohol use	\
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	
..	
995	False	False	False	False	False	False	
996	False	False	False	False	False	False	
997	False	False	False	False	False	False	
998	False	False	False	False	False	False	
999	False	False	False	False	False	False	
	Dust Allergy	OccuPational Hazards	Genetic Risk	chronic Lung Disease	Lung Disease	\	
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	
..	
995	False	False	False	False	False	False	
996	False	False	False	False	False	False	
997	False	False	False	False	False	False	
998	False	False	False	False	False	False	
999	False	False	False	False	False	False	
	... Fatigue	Weight Loss	Shortness of Breath	Wheezing	Wheezing	\	
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	
..	
995	... False	False	False	False	False	False	
996	... False	False	False	False	False	False	
997	... False	False	False	False	False	False	
998	... False	False	False	False	False	False	
999	... False	False	False	False	False	False	
	Swal0ing Difficulty	Clubbing of Finger Nails	Frequent Cold	Dry Cough	Dry Cough	\	
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	
..	
995	False	False	False	False	False	False	
996	False	False	False	False	False	False	
997	False	False	False	False	False	False	
998	False	False	False	False	False	False	
999	False	False	False	False	False	False	
	Snoring Level						
0	False	False					
1	False	False					
2	False	False					
3	False	False					
4	False	False					

```
print(d.describe())
print(d)
```

```
index      Age   Gender Air Pollution Alcohol use \
count 1000.000000 1000.000000 1000.000000 1000.000000 1000.000000
mean 499.500000 37.174000 1.402000 3.8400 4.563000
std 288.819436 12.005493 0.490547 2.0304 2.620477
min 0.000000 14.000000 1.000000 1.0000 1.000000
25% 249.750000 27.750000 1.000000 2.0000 2.000000
50% 499.500000 36.000000 1.000000 3.0000 5.000000
75% 749.250000 45.000000 2.000000 6.0000 7.000000
max 999.000000 73.000000 2.000000 8.0000 8.000000
```

```
Dust Allergy Occupational Hazards Genetic Risk chronic Lung Disease \
count 1000.000000 1000.000000 1000.000000 1000.000000
mean 5.165000 4.840000 4.580000 4.380000
std 1.980833 2.107805 2.126999 1.848518
min 1.000000 1.000000 1.000000 1.000000
25% 4.000000 3.000000 2.000000 3.000000
50% 6.000000 5.000000 5.000000 4.000000
75% 7.000000 7.000000 7.000000 6.000000
max 8.000000 8.000000 7.000000 7.000000
```

```
Balanced Diet ... Fatigue Weight Loss Shortness of Breath \
count 1000.000000 ... 1000.000000 1000.000000 1000.000000
mean 4.491000 ... 3.856000 3.855000 4.240000
std 2.135528 ... 2.244616 2.206546 2.285087
min 1.000000 ... 1.000000 1.000000 1.000000
25% 2.000000 ... 2.000000 2.000000 2.000000
50% 4.000000 ... 3.000000 3.000000 4.000000
75% 7.000000 ... 5.000000 6.000000 6.000000
max 7.000000 ... 9.000000 8.000000 9.000000
```

```
Wheezing Swallowing Difficulty Clubbing of Finger Nails \
count 1000.000000 1000.000000 1000.000000
mean 3.777000 3.746000 3.923000
std 2.041921 2.270383 2.388048
min 1.000000 1.000000 1.000000
25% 2.000000 2.000000 2.000000
50% 4.000000 4.000000 4.000000
75% 5.000000 5.000000 5.000000
max 8.000000 8.000000 9.000000
```

```
Frequent Cold Dry Cough Snoring Level
count 1000.000000 1000.000000 1000.000000 1000.000000
mean 3.536000 3.853000 2.926000 1.062000
std 1.832502 2.039007 1.474686 0.815365
min 1.000000 1.000000 1.000000 0.000000
25% 2.000000 2.000000 2.000000 0.000000
50% 3.000000 4.000000 3.000000 1.000000
75% 5.000000 6.000000 4.000000 2.000000
max 7.000000 7.000000 7.000000 2.000000
```

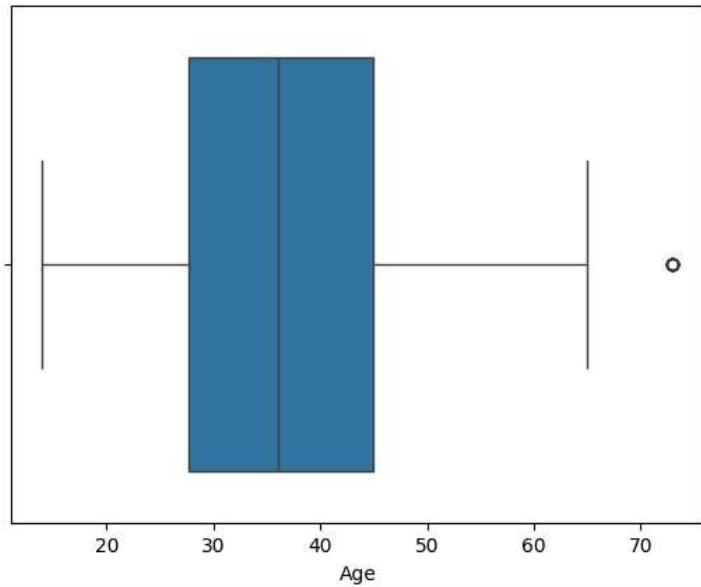
```
[8 rows x 25 columns]
index Patient Id Age Gender Air Pollution Alcohol use Dust Allergy \
0 0 P1 33 1 2 4 5
1 1 P10 17 1 3 1 5
2 2 P100 35 1 4 5 6
3 3 P1000 37 1 7 7 7
4 4 P101 46 1 6 8 7
```

```
d.columns
```

```
Index(['index', 'Patient Id', 'Age', 'Gender', 'Air Pollution', 'Alcohol use',  
       'Dust Allergy', 'Occupational Hazards', 'Genetic Risk',  
       'chronic Lung Disease', 'Balanced Diet', 'Obesity', 'Smoking',  
       'Passive Smoker', 'Chest Pain', 'Coughing of Blood', 'Fatigue',  
       'Weight Loss', 'Shortness of Breath', 'Wheezing', 'Swallowing Difficulty',  
       'Clubbing of Finger Nails', 'Frequent Cold', 'Dry Cough', 'Snoring',  
       'Level'],  
      dtype='object')
```

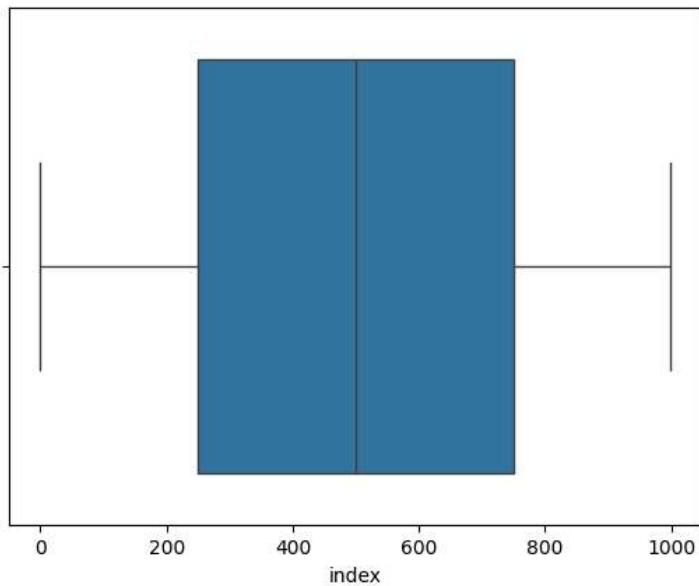
```
import seaborn as sns  
sns.boxplot(data=d,x='Age')
```

```
<Axes: xlabel='Age'>
```



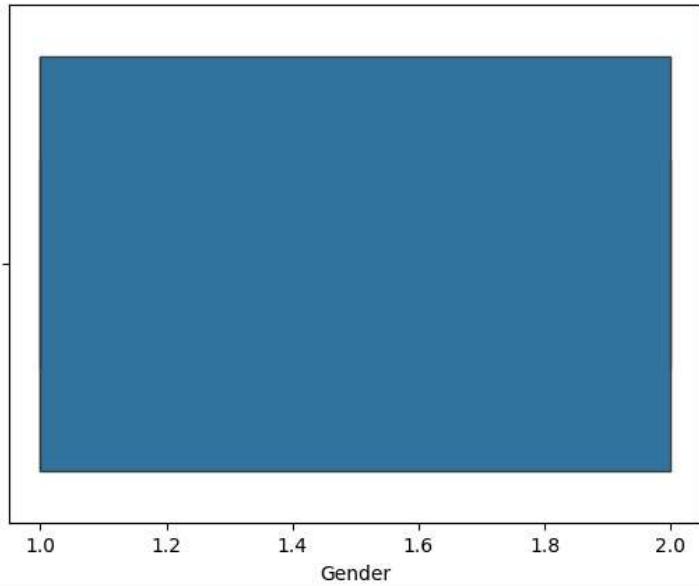
```
import seaborn as sns  
sns.boxplot(data=d,x='index')
```

```
⤵ <Axes: xlabel='index'>
```



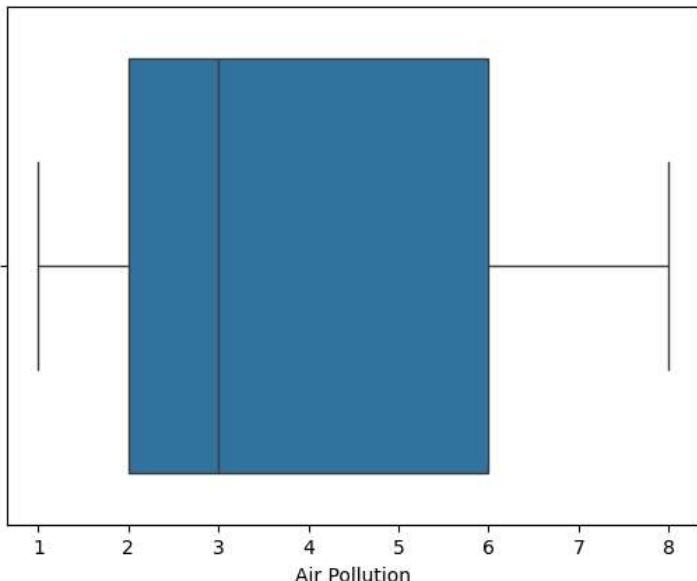
```
import seaborn as sns  
sns.boxplot(data=d,x='Gender')
```

```
⤵ <Axes: xlabel='Gender'>
```



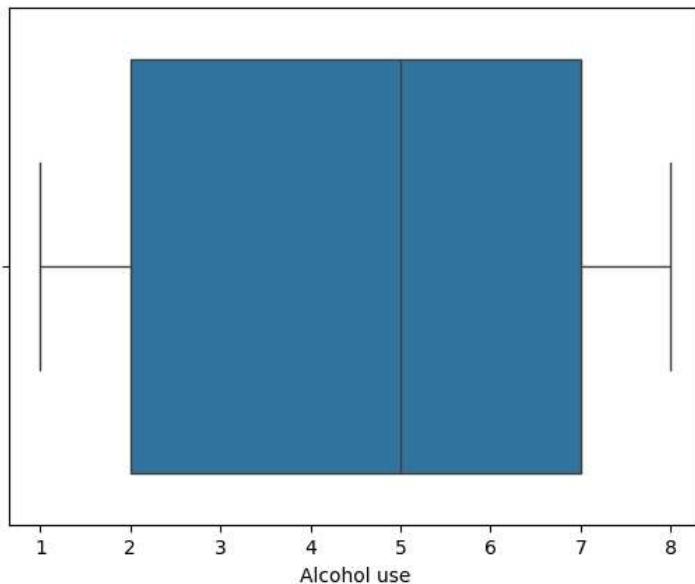
```
import seaborn as sns  
sns.boxplot(data=d,x='Air Pollution')
```

⤵ <Axes: xlabel='Air Pollution'>



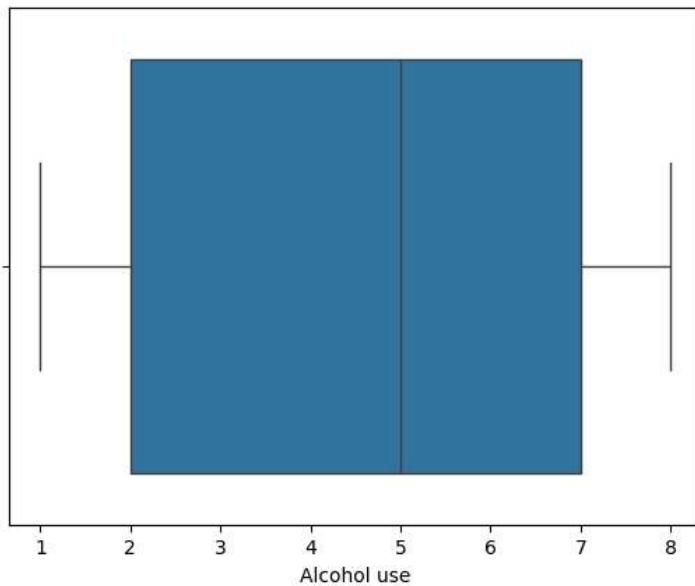
```
import seaborn as sns  
sns.boxplot(data=d,x='Alcohol use')
```

```
⤵ <Axes: xlabel='Alcohol use'>
```



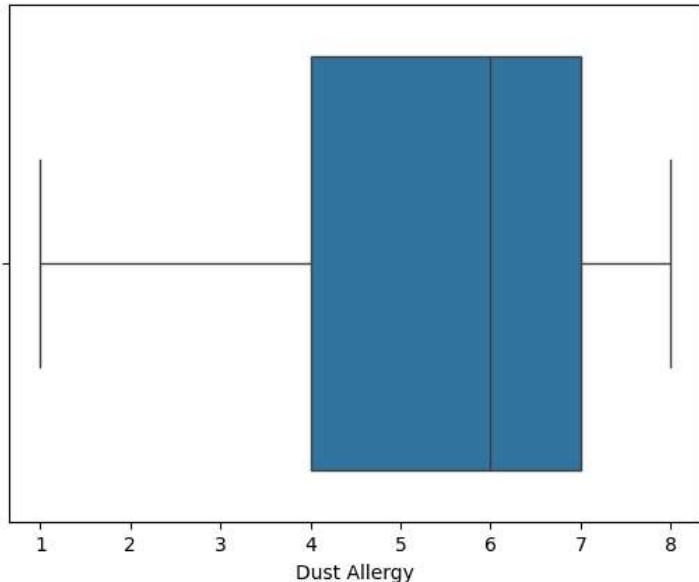
```
import seaborn as sns  
sns.boxplot(data=d,x='Alcohol use')
```

```
⤵ <Axes: xlabel='Alcohol use'>
```



```
import seaborn as sns  
sns.boxplot(data=d,x='Dust Allergy')
```

⤵ <Axes: xlabel='Dust Allergy'>



```
import seaborn as sns  
sns.boxplot(data=d,x='Occupational Hazards')
```

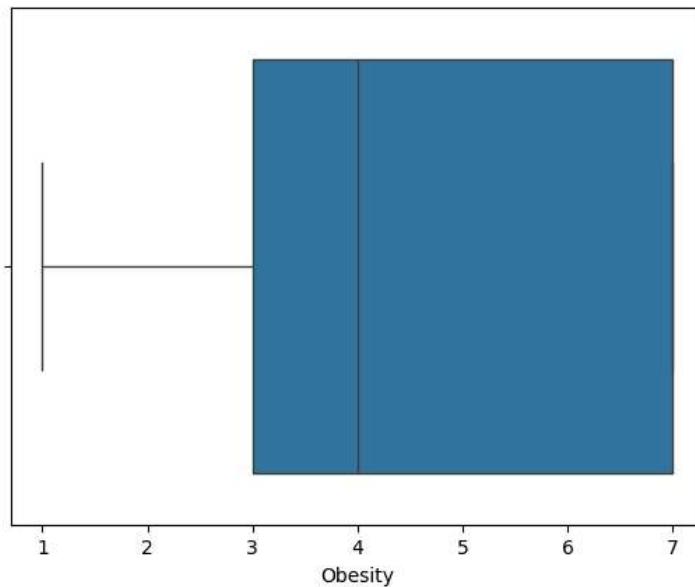
```
import seaborn as sns  
sns.boxplot(data=d,x='Genetic Risk')
```

```
import seaborn as sns  
sns.boxplot(data=d,x='Chronic Lung Disease')
```

```
import seaborn as sns  
sns.boxplot(data=d,x='Balanced Diet')
```

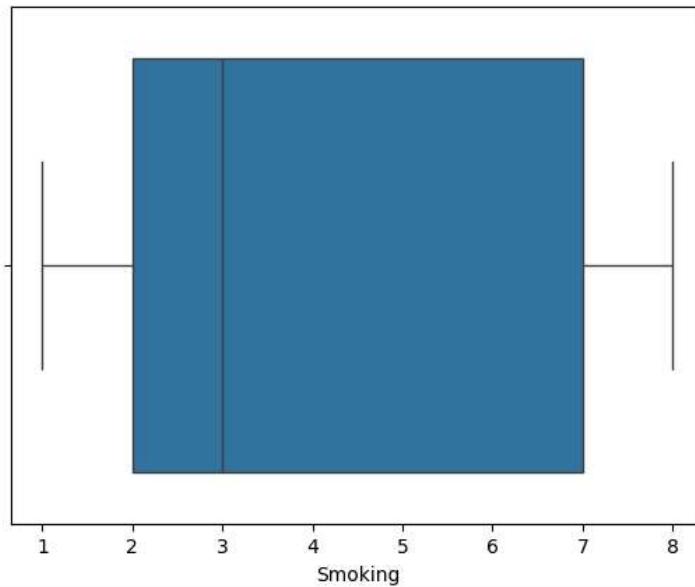
```
import seaborn as sns  
sns.boxplot(data=d,x='Obesity')
```

```
⤵ <Axes: xlabel='Obesity'>
```



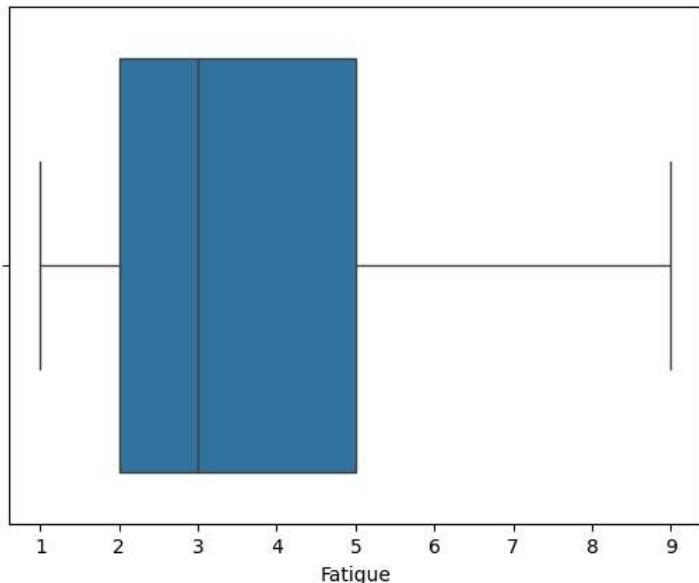
```
import seaborn as sns  
sns.boxplot(data=d,x='Smoking')
```

```
⤵ <Axes: xlabel='Smoking'>
```



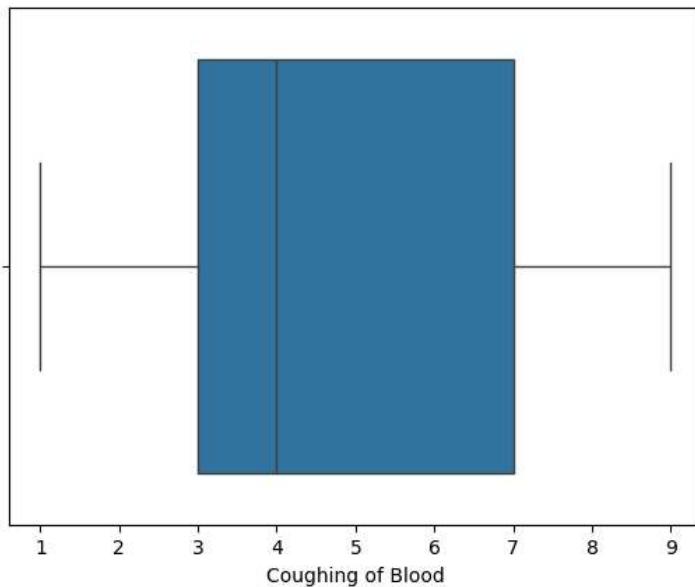
```
import seaborn as sns  
sns.boxplot(data=d,x='Fatigue')
```

⤵ <Axes: xlabel='Fatigue'>



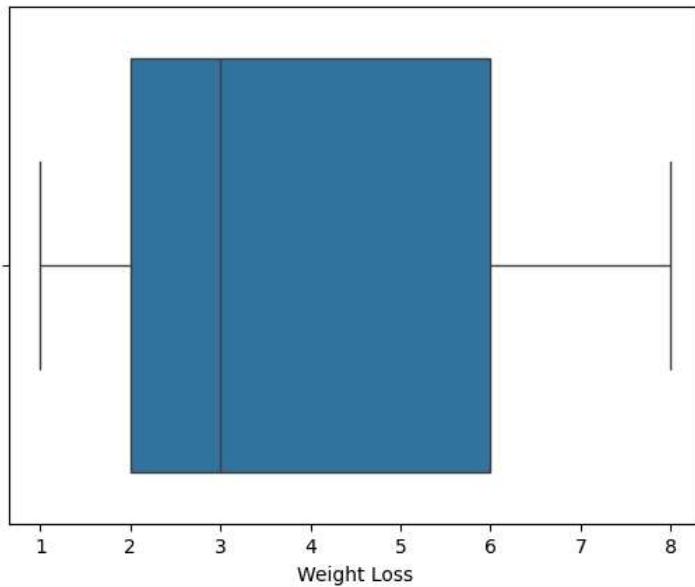
```
import seaborn as sns  
sns.boxplot(data=d,x='Coughing of Blood')
```

```
⤵ <Axes: xlabel='Coughing of Blood'>
```



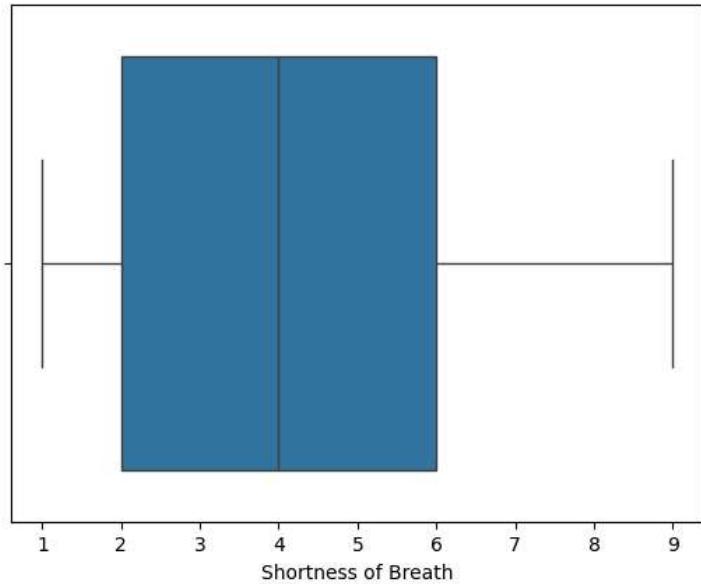
```
import seaborn as sns  
sns.boxplot(data=d,x='Weight Loss')
```

```
⤵ <Axes: xlabel='Weight Loss'>
```



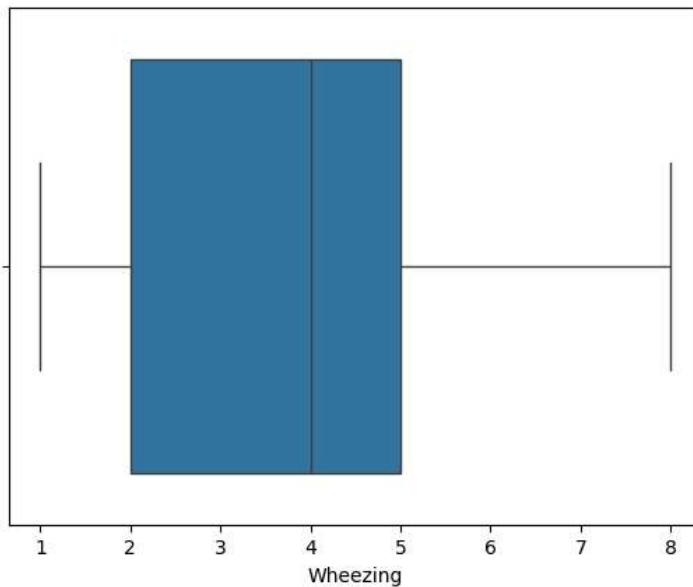
```
import seaborn as sns  
sns.boxplot(data=d,x='Shortness of Breath')
```

⤵ <Axes: xlabel='Shortness of Breath'>



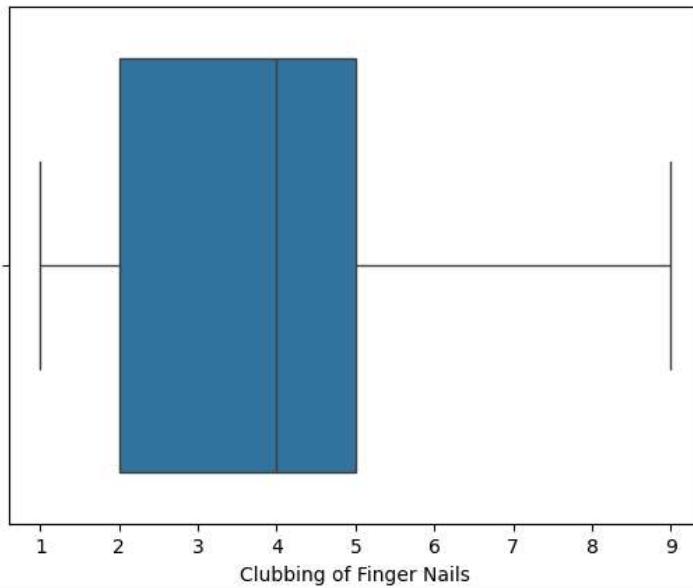
```
import seaborn as sns  
sns.boxplot(data=d,x='Wheezing')
```

```
⤵ <Axes: xlabel='Wheezing'>
```



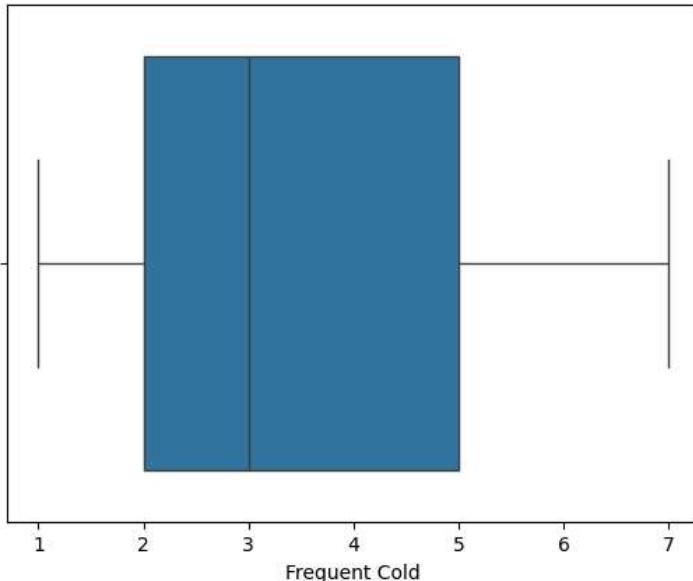
```
import seaborn as sns  
sns.boxplot(data=d,x='Clubbing of Finger Nails')
```

```
⤵ <Axes: xlabel='Clubbing of Finger Nails'>
```



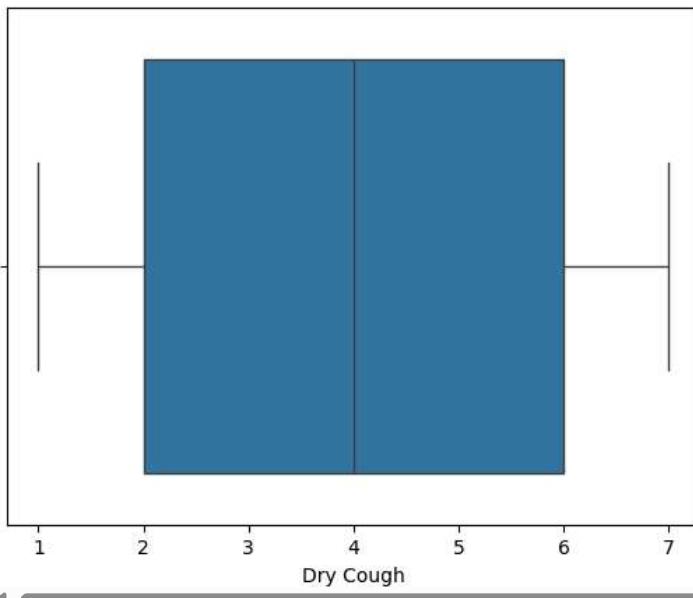
```
import seaborn as sns  
sns.boxplot(data=d,x='Frequent Cold')
```

```
⤵ <Axes: xlabel='Frequent Cold'>
```



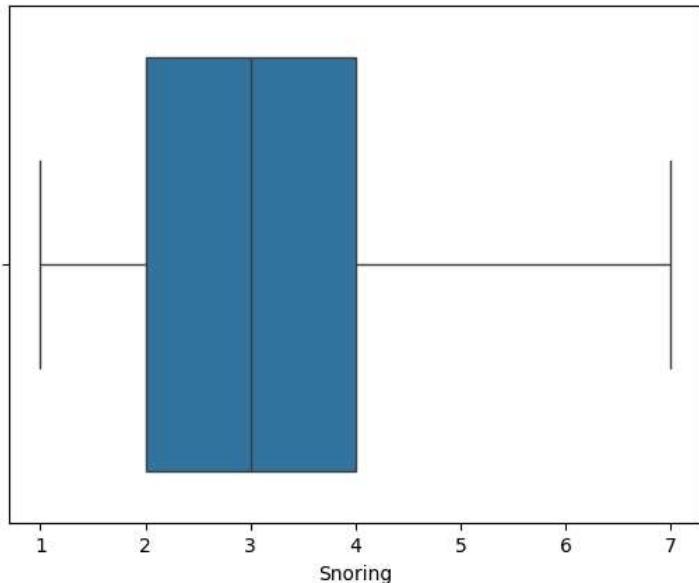
```
import seaborn as sns  
sns.boxplot(data=d,x='Dry Cough')
```

```
⤵ <Axes: xlabel='Dry Cough'>
```



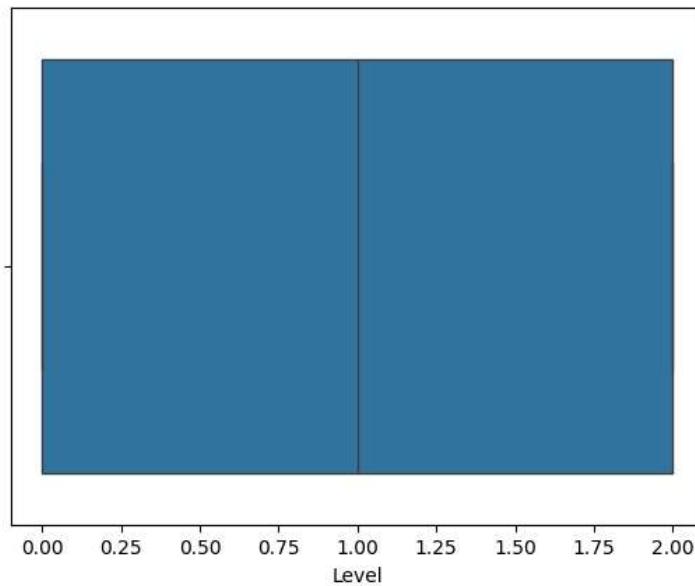
```
import seaborn as sns  
sns.boxplot(data=d,x='Snoring')
```

⤵ <Axes: xlabel='Snoring'>



```
import seaborn as sns  
sns.boxplot(data=d,x='Level')
```

```
⤵ <Axes: xlabel='Level'>
```



```
print(d.dtypes)
```

```
⤵ index          int64
Patient Id      object
Age             int64
Gender           int64
Air Pollution   int64
Alcohol use     int64
Dust Allergy    int64
OccuPational Hazards int64
Genetic Risk    int64
chronic Lung Disease int64
Balanced Diet   int64
Obesity         int64
Smoking          int64
Passive Smoker   int64
Chest Pain       int64
Coughing of Blood int64
Fatigue          int64
Weight Loss      int64
Shortness of Breath int64
Wheezing          int64
Swal0ing Difficulty int64
Clubbing of Finger Nails int64
```

```
Frequent Cold      int64  
Dry Cough        int64  
Snoring          int64  
Level            int64  
dtype: object
```

```
import numpy as np
```

```
d_numeric = d.select_dtypes(include=[np.number])  
corr_matrix = d_numeric.corr()  
print(corr_matrix)
```

```
index      index    Age   Gender Air Pollution \
index  1.000000  0.002674 -0.025739  0.053307
Age     0.002674  1.000000 -0.202086  0.099494
Gender   -0.025739 -0.202086  1.000000 -0.246912
Air Pollution  0.053307  0.099494 -0.246912  1.000000
Alcohol use    0.041374  0.151742 -0.227636  0.747293
Dust Allergy   0.037960  0.035202 -0.204312  0.637503
OccuPational Hazards  0.032355  0.062177 -0.192343  0.608924
Genetic Risk    0.030725  0.073151 -0.222727  0.705276
chronic Lung Disease  0.025177  0.128952 -0.205061  0.626701
Balanced Diet   0.030743  0.004863 -0.099741  0.524873
Obesity        0.050584  0.034337 -0.123813  0.601468
Smoking         0.018407  0.075333 -0.206924  0.481902
Passive Smoker   0.019517  0.004908 -0.184826  0.606764
Chest Pain       0.022210  0.012864 -0.218426  0.585734
Coughing of Blood  0.049401  0.053006 -0.146505  0.607829
Fatigue          0.042346  0.095059 -0.116467  0.211724
Weight Loss       0.026393  0.106946 -0.057993  0.258016
Shortness of Breath  0.027950  0.035329 -0.045972  0.269558
Wheezing          0.015078 -0.095354 -0.076304  0.055368
Swallowing Difficulty  0.005573 -0.105833 -0.058324 -0.080918
Clubbing of Finger Nails  0.015706  0.039258 -0.034219  0.241065
Frequent Cold    0.045687 -0.012706 -0.000526  0.174539
Dry Cough         0.003793  0.012128 -0.123001  0.261489
Snoring          -0.002957 -0.004700 -0.181618 -0.021343
Level            0.057758  0.060048 -0.164985  0.636038

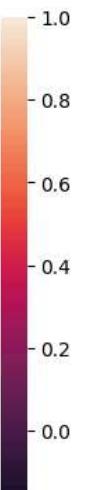
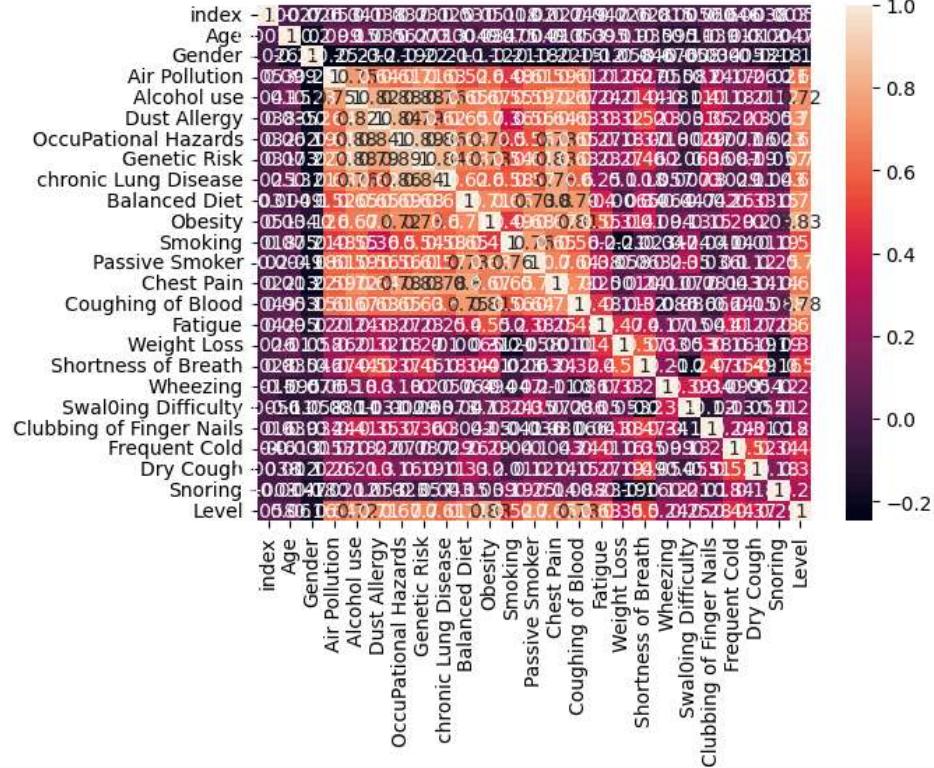
index      Alcohol use  Dust Allergy OccuPational Hazards \
index  0.041374  0.037960  0.032355
Age     0.151742  0.035202  0.062177
Gender   -0.227636 -0.204312 -0.192343
Air Pollution  0.747293  0.637503  0.608924
Alcohol use    1.000000  0.818644  0.878786
Dust Allergy   0.818644  1.000000  0.835860
OccuPational Hazards  0.878786  0.835860  1.000000
Genetic Risk    0.877210  0.787904  0.893049
chronic Lung Disease  0.763576  0.619556  0.858284
Balanced Diet   0.653352  0.647197  0.691509
Obesity          0.669312  0.700676  0.722191
Smoking          0.547035  0.358691  0.497693
Passive Smoker   0.592576  0.560002  0.555311
Chest Pain        0.717242  0.639983  0.775619
Coughing of Blood  0.667612  0.625291  0.645947
Fatigue           0.237245  0.332472  0.267844
Weight Loss        0.207851  0.321756  0.176226
Shortness of Breath  0.435785  0.518682  0.366482
```

Wheezing	0.180817	0.304850	0.178925
Swallowing Difficulty	-0.114073	0.031141	-0.002853
Clubbing of Finger Nails	0.414992	0.345714	0.366447
Frequent Cold	0.180778	0.219389	0.077166
Dry Cough	0.211277	0.300195	0.159887
Snoring	0.122694	0.052844	0.022916
Level	0.718710	0.713839	0.673255

	Genetic Risk	chronic Lung Disease	Balanced Diet	\
index	0.030725	0.025177	0.030743	
Age	0.073151	0.128952	0.004863	
Gender	0.222727	0.220561	0.200741	

```
import seaborn as sn
import matplotlib.pyplot as plt
sn.heatmap(corr_matrix, annot=True)
```

→ <Axes: >



```
import numpy as np
import pandas as pd

# Assuming 'd' is your DataFrame
d_numeric = d.select_dtypes(include=[np.number])
```

```
cov_matrix = d_numeric.cov()
print(cov_matrix)
```

	index	Age	Gender	Air Pollution	\
index	83416.666667	9.273273	-3.646647	31.260260	
Age	9.273273	144.131856	-1.190138	2.425265	
Gender	-3.646647	-1.190138	0.240637	-0.245926	
Air Pollution	31.260260	2.425265	-0.245926	4.122523	
Alcohol use	31.313814	4.773812	-0.292619	3.976056	
Dust Allergy	21.717217	0.837127	-0.198529	2.563964	
OccuPational Hazards	19.696697	1.573413	-0.198879	2.606006	
Genetic Risk	18.874875	1.867948	-0.232392	3.045846	
chronic Lung Disease	13.441441	2.861742	-0.185946	2.352152	
Balanced Diet	18.961461	0.124691	-0.104486	2.275836	
Obesity	31.044545	0.875966	-0.129059	2.594995	
Smoking	13.269269	2.257305	-0.253349	2.442122	
Passive Smoker	13.031532	0.136206	-0.209600	2.848048	
Chest Pain	14.626627	0.352140	-0.244320	2.711792	
Coughing of Blood	34.642142	1.545079	-0.174492	2.996436	
Fatigue	27.452452	2.561618	-0.128240	0.964925	
Weight Loss	16.820320	2.833063	-0.062773	1.155956	
Shortness of Breath	18.446446	0.969209	-0.051532	1.250651	
Wheezing	8.892392	-2.337536	-0.076430	0.229550	
Swallowing Difficulty	3.654655	-2.884689	-0.064957	-0.373013	
Clubbing of Finger Nails	10.832332	1.125524	-0.040086	1.168849	
Frequent Cold	24.180180	-0.279544	-0.000472	0.649409	
Dry Cough	2.233734	0.296875	-0.123029	1.082563	
Snoring	-1.259259	-0.083207	-0.131383	-0.063904	
Level	13.601602	0.587800	-0.065990	1.052973	

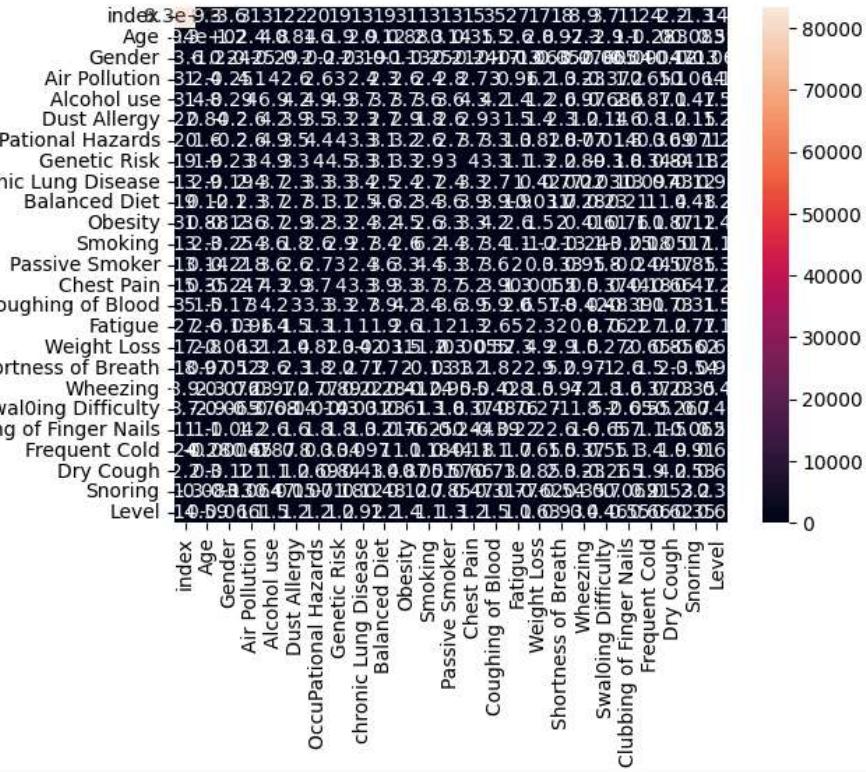
	Alcohol use	Dust Allergy	OccuPational Hazards	\
index	31.313814	21.717217	19.696697	
Age	4.773812	0.837127	1.573413	
Gender	-0.292619	-0.198529	-0.198879	
Air Pollution	3.976056	2.563964	2.606006	
Alcohol use	6.866898	4.249354	4.853934	
Dust Allergy	4.249354	3.923699	3.489890	
OccuPational Hazards	4.853934	3.489890	4.442843	
Genetic Risk	4.889349	3.319620	4.003804	
chronic Lung Disease	3.698759	2.268569	3.344144	
Balanced Diet	3.656223	2.737723	3.112673	
Obesity	3.726932	2.949224	3.234635	
Smoking	3.577854	1.773353	2.618298	
Passive Smoker	3.589805	2.564389	2.705906	
Chest Pain	4.285692	2.890621	3.727808	
Coughing of Blood	4.247631	3.007272	3.305746	
Fatigue	1.395467	1.478238	1.267227	
Weight Loss	1.201837	1.406331	0.819620	
Shortness of Breath	2.609489	2.347748	1.765165	
Wheezing	0.967517	1.233028	0.770090	
Swallowing Difficulty	-0.678677	0.140050	-0.013654	
Clubbing of Finger Nails	2.596948	1.635340	1.844525	
Frequent Cold	0.868100	0.796356	0.298058	
Dry Cough	1.128890	1.212467	0.687167	
Snoring	0.474136	0.154364	0.071231	

Level	1.535630	1.152923	1.157077
index	Genetic Risk	chronic Lung Disease	Balanced Diet
Age	18.874875	13.441441	18.961461

```
import seaborn as sn
```

```
import matplotlib.pyplot as plt  
sn.heatmap(cov_matrix, annot=True)
```

 <Axes: >



```
x=d.iloc[:,3:8]  
y=d.iloc[:,8:9]  
print(x)  
print(y)
```

	Gender	Air Pollution	Alcohol use	Dust Allergy	Occupational Hazards
0	1	2	4	5	4
1	1	3	1	5	3
2	1	4	5	6	5
3	1	7	7	7	7
4	1	6	8	7	7
..
995	1	6	7	7	7
996	2	6	8	7	7

```
997      2        4        5        6        5
998      2        6        8        7        7
999      1        6        5        6        5
```

```
[1000 rows x 5 columns]
```

```
  Genetic Risk
```

```
0        3
1        4
2        5
3        6
4        7
..
995     ..
996     7
997     5
998     7
999     5
```

```
[1000 rows x 1 columns]
```

```
from sklearn.model_selection import train_test_split
```

```
from imblearn.over_sampling import SMOTE
sm=SMOTE(random_state=42)
x,y=sm.fit_resample(x,y)
print(x.shape)
x_train,x_test,y_train,y_test = train_test_split(x,y,random_state=True,test_size=0.20)
```

```
→ (2289, 5)
```

```
from sklearn.linear_model import LogisticRegression
```

```
lr=LogisticRegression()
```

```
mm=lr.fit(x_train,y_train)
```

```
→ /usr/local/lib/python3.10/dist-packages/scikit-learn/utils/validation.py:1183: DataConversionWarning: A column-vector y was passed when a 1d array was expected. Please change the
      y = column_or_1d(y, warn=True)
/usr/local/lib/python3.10/dist-packages/scikit-learn/linear_model/_logistic.py:460: ConvergenceWarning: lbfgs failed to converge (status=1):
STOP: TOTAL NO. OF ITERATIONS REACHED LIMIT.
```

Increase the number of iterations (`max_iter`) or scale the data as shown in:

<https://scikit-learn.org/stable/modules/preprocessing.html>

Please also refer to the documentation for alternative solver options:

https://scikit-learn.org/stable/modules/linear_model.html#logistic-regression

`n_iter_i = _check_optimize_result(`

```
yp=mm.predict(x_test)
```

```
print(yp)

[2 1 1 4 7 2 2 3 5 5 7 2 7 2 4 2 7 1 7 4 5 3 1 4 1 3 5 7 3 7 5 1 1 7 5 2 7
5 6 1 2 5 5 7 6 6 7 7 5 2 4 1 1 5 5 1 1 5 2 5 4 2 2 2 1 7 3 5 2 4 4 7 4 1
4 1 3 7 3 1 2 2 1 5 2 6 1 7 5 4 4 4 2 3 7 4 1 5 5 6 1 7 1 7 6 5 7 4 3 5 1
1 5 2 5 3 5 5 4 4 7 1 2 3 4 2 5 3 2 1 1 4 3 7 7 1 2 7 5 2 4 1 5 1 1 1 5 2
1 2 4 7 5 5 5 2 2 1 2 1 7 1 3 7 2 1 2 2 7 1 1 1 3 5 7 7 5 1 2 3 7 1 1 1 3
2 5 1 2 5 1 5 1 2 1 4 5 2 1 7 6 4 7 5 7 5 1 7 5 2 4 5 4 7 7 7 5 1 4 2 1 7
1 2 3 5 7 4 2 4 7 5 1 2 7 6 1 2 5 4 1 7 2 7 1 5 4 1 5 6 1 5 5 4 7 1 2 5 1
7 7 1 1 3 7 5 7 7 1 5 2 7 4 7 5 7 5 5 1 2 1 5 4 5 7 2 1 7 7 2 5 6 1 4 1
7 2 1 7 1 7 5 5 1 1 6 5 5 3 5 7 7 7 7 2 1 4 5 7 3 7 4 4 7 1 4 1 2 2 2 7
5 7 1 4 4 4 1 7 1 6 2 7 4 5 1 4 1 5 5 7 1 5 1 3 2 2 1 2 3 2 1 7 1 1 7 2
1 5 2 1 1 7 7 3 1 2 4 2 7 1 1 1 7 2 4 1 4 2 4 1 5 7 7 7 1 1 2 3 1 7 5 2 2
2 1 4 5 4 2 2 1 1 5 5 1 2 2 2 5 4 2 4 4 5 7 1 5 2 1 4 1 1 2 5 7 7 5 1 5 7
5 7 5 1 1 3 7 2 2 6 6 1 1 4]
```

```
from sklearn.metrics import accuracy_score
print(accuracy_score(yp,y_test))
```

```
0.6615720524017468
```

```
from sklearn.tree import DecisionTreeClassifier
```

```
classifier=DecisionTreeClassifier(criterion='entropy',random_state=0)
mm=classifier.fit(x_train,y_train)
```

```
yp=mm.predict(x_test)
```

```
print(yp)
```

```
[4 1 3 4 7 2 3 1 5 5 7 2 7 4 2 3 7 1 6 4 5 3 1 2 1 1 5 7 3 7 5 1 1 7 6 3 7
5 6 3 3 5 5 6 6 6 7 7 5 2 2 1 1 5 5 1 3 5 2 5 2 4 4 4 1 7 1 5 2 2 4 7 2 1
4 1 3 7 3 3 3 3 1 5 2 6 3 6 5 4 4 2 2 3 6 4 1 5 5 6 1 7 1 7 6 5 7 2 3 5 1
1 5 2 5 3 5 5 4 4 7 1 4 3 4 2 5 3 2 1 1 2 3 7 7 1 4 7 5 4 2 1 5 1 1 3 5 2
1 2 4 7 5 5 5 3 4 1 2 1 7 1 3 7 4 1 2 4 6 3 1 1 3 6 7 7 6 1 2 3 7 1 1 3 3
2 5 1 2 5 1 5 3 2 1 4 6 2 1 7 3 4 7 5 7 5 3 7 5 3 2 5 4 7 7 7 5 1 4 3 1 6
1 2 3 5 7 6 4 4 7 5 1 3 7 6 1 3 5 2 1 7 4 7 1 5 2 3 5 6 1 5 5 2 7 2 4 5 1
7 7 1 1 3 6 5 7 7 3 5 2 7 2 7 5 7 5 5 3 2 1 5 4 5 7 4 1 7 7 4 5 6 1 2 1
7 3 1 7 1 7 5 5 1 1 6 5 5 3 5 7 7 7 6 7 4 1 4 5 7 3 7 2 2 7 3 2 1 3 2 4 7
5 7 1 4 4 4 1 7 1 3 2 6 2 5 1 2 2 5 5 7 1 5 1 3 3 2 1 4 3 6 1 7 1 3 1 7 2
1 5 4 1 1 7 7 2 2 4 2 2 7 2 1 7 2 2 1 2 4 4 2 5 7 6 7 1 2 4 2 1 7 5 5 2
```