

Analysis Of Diseases and Symptoms

Project Objective : To analyse the Symptoms of a Disease and its Diagnostics.

The dataset provided to us helps us to analyse the symptoms of a disease and helps us to diagnose it and tells us how fatal the disease is on a scale of 0 to 3(0 being the least fatal and 3 being most)

Description of Data Set columns :

Three Databases are used in this project :-

Dataset1 (symptom) :

- **syd:>Symptom ID:**
Gives us the unique Identification number of a specific symptom
Datatype:int64
Non-null values:272
- **symptom:>Tells us the possible symptoms of all diseases**
Datatype:object
Non-null values:246

```
import pandas as pd
sym=pd.read_csv("symptom.csv")
dia=pd.read_csv("diagnose.csv")
data=pd.read_csv("datamatch.csv")
#print(dis)
print(sym.info())
#print(dia.info())
#print(data.info())
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 272 entries, 0 to 271
Data columns (total 2 columns):
syd          272 non-null int64
symptom      246 non-null object
dtypes: int64(1), object(1)
memory usage: 4.3+ KB
None
```

Dataset2 (diagnose) :

- **did:>Disease ID .Gives us the unique Identification number of a specific disease**

Datatype:int64

Non-null values:1166

- diagnosis:>Tells us the possible diseases after diagnosing the symptoms.

Datatype:object

Non-null values:1166

```
import pandas as pd
sym=pd.read_csv("symptom.csv")
dia=pd.read_csv("diagnose.csv")
data=pd.read_csv("datamatch.csv")
#print(dis)
#print(sym.info())
print(dia.info())
#print(data.info())
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 1166 entries, 0 to 1165
Data columns (total 2 columns):
did          1166 non-null int64
diagnose      1166 non-null object
dtypes: int64(1), object(1)
memory usage: 18.3+ KB
None
```

Dataset3 (datamatch) :

- syd:>Symptom ID .Gives us the unique Identification number of a specific symptom
Datatype:int64
Non-null values:5568
- did:>Disease ID .Gives us the unique Identification number of a specific disease
Datatype:int64
Non-null values:5568
- wei:>how possible the fatal a disease can be depending upon the lives it claims annually
Datatype:float64
Non-null values:5371

```
import pandas as pd
sym=pd.read_csv("symptom.csv")
dia=pd.read_csv("diagnose.csv")
data=pd.read_csv("datamatch.csv")
#print(dis)
#print(sym.info())
#print(dia.info())
print(data.info())
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 5568 entries, 0 to 5567
Data columns (total 3 columns):
syd      5568 non-null int64
did      5568 non-null int64
wei      5371 non-null float64
dtypes: float64(1), int64(2)
memory usage: 130.6 KB
None
```

Data Interpretation :-

Dataset1 (symptom) :

```
import pandas as pd
sym=pd.read_csv("symptom.csv")
#dia=pd.read_csv("diagnose.csv")
#data=pd.read_csv("datamatch.csv")
print(sym)
```

	syd	symptom
0	1	Upper abdominal pain
1	2	Lower abdominal pain
2	3	Abscess (Collection of pus)
3	4	Alcohol abuse
4	5	Anxiety (Nervousness)
5	6	Arm ache or pain
6	7	Back ache or pain
7	8	Bleeding tendency
8	9	Blood in vomit
9	10	Bloody diarrhea
10	11	Pain or soreness of breast
11	12	Calf pain
12	13	Chest pressure
13	14	Chills
14	15	Change in behavior
15	16	Constipation
16	17	Cough
17	18	Dark stools
18	19	Depressed
19	20	Diarrhea
20	21	Dizziness
21	22	Double vision (Diplopia)
22	23	Ear pressure
23	24	Pain in the ear
24	25	Elbow ache or pain
25	26	Eye pain (Irritation)
26	27	Facial pain
27	28	Fainting
28	29	Fever
29	30	Fever in the returning traveler
..
242	257	Snoring
243	258	Dry skin
244	259	Itchy eyes
245	261	Elbow swelling

Dataset2 (*diagnose*) :

```
import pandas as pd
#sym=pd.read_csv("symptom.csv")
dia=pd.read_csv("diagnose.csv")
#data=pd.read_csv("datamatch.csv")
print(dia)
```

	did	diagnose
0	1	Abdominal aortic aneurysm□(enlarged major bloo...
1	2	Abdominal swelling
2	3	Abdominal trauma
3	4	Abrasions□ (scrapes)
4	5	ACE inhibitor induced cough□blood pressure med...
5	6	acetaminophen overdose□Adverse reaction to ace...
6	7	Tylenol □acetaminophen poisoning
7	8	Achilles tendonitis□ (heel tendon inflammation)
8	9	Achilles tendon rupture□(heel tendon tear)
9	10	Acid □LSD abuse
10	11	Acidosis□ (excessive acid in the body)
11	12	Acoustic neuroma□(ear nerve tumor)
12	13	AC joint separation□acromioclavicular joint se...
13	14	Acute angle closure glaucoma□increased inner e...
14	15	Acute fatty liver of pregnancy
15	16	Adenoiditis□(a type of lymph node inflammation)
16	17	Adenovirus infection□ (virus infection)
17	18	Frozen shoulder□ (adhesive capsulitis of shoul...
18	19	Adjustment disorder□ (poor adjustment to life ...
19	20	Alcohol □ethanol intoxication
20	21	Alcohol □ethanol abuse
21	22	Alcohol □ethanol poisoning□ (overdose)
22	23	Alcohol withdrawal syndrome□ (mild)
23	25	Alcoholism
24	26	Allergic reaction
25	27	Allergic rhinitis□ (allergic reaction in the n...
26	28	Allergy
27	29	Confusion□ (altered mental status)
28	30	Altered mental status□confusion
29	31	Altitude illness□Illnesses due to high altitud...
...
1136	1467	Cerebellar Ataxia
1137	1469	Complex partial seizures□psychomotor epilepsy
1138	1471	Meralgia Paresthetica
1139	1473	Retinopathv

Dataset3 (*datamatch*) :

```
import pandas as pd
#sym=pd.read_csv("symptom.csv")
#dia=pd.read_csv("diagnose.csv")
data=pd.read_csv("datamatch.csv")
print(data)
```

	syd	did	wei
0	1	163	2.0
1	1	164	2.0
2	1	165	1.0
3	1	187	2.0
4	1	306	2.0
5	1	307	1.0
6	1	308	2.0
7	1	309	2.0
8	1	354	1.0
9	1	401	1.0
10	1	411	1.0
11	1	513	1.0
12	1	546	2.0
13	1	722	1.0
14	2	56	3.0
15	2	179	2.0
16	2	236	1.0
17	2	388	2.0
18	2	539	1.0
19	2	540	1.0
20	2	557	1.0
21	2	600	1.0
22	2	793	2.0
23	2	795	1.0
24	3	44	1.0
25	3	106	1.0
26	3	108	0.0
27	3	707	2.0
28	3	209	2.0
29	3	250	1.0
...
5538	277	650	0.0
5539	277	1034	2.0
5540	277	227	1.0
5541	277	1080	0.0

Analysis performed on the basis of various parameters :

1. Perform analysis on diseases which are most untraceable
2. Perform analysis on the drug related diseases and their fatality
3. Perform analysis to find the easiest Drug Related Disease
4. Perform analysis to find the most curable cancer
5. Perform analyses to find the most fatal cancer
6. Perform analysis to find the top 10 Severe cancer type.
7. Perform analysis to find breast related diseases with their specific symptoms
8. Perform analysis to find the number of diseases whose fatality is 1 or 2
9. Perform analysis on the fact to find possible diseases if the symptoms are delusions and hallucinations.
10. Perform analysis on the fact that diseases which have symptoms of both headache and fever
11. Perform analysis on the fact that Showing, the diseases based on fatality rate
12. Perform analysis on the fact that the number of Bacterial and cervical diseases and their fatality rate
13. Perform analysis on the fact that the symptoms related to eye (irritation) and has a fatality rate equal to 3
14. Perform analysis on the fact that what is the chance of dying from cold symptoms(Graph)
15. Perform analysis on the fact that Diseases with more than twenty five symptoms
16. Perform analysis on the fact that If delusions and hallucinations occur in a patient then it leads to mental illness

- 17.** Perform analysis on the fact that most common symptoms found in the diseases of highest fatality
- 18.** Perform analysis on the fact that fatality rate of diseases which have symptoms of vomiting
- 19.** Perform analysis on the fact that the symptoms related to different types of organ failures and has a fatality rate equal to 3
- 20.** Perform analysis to find the occurrence of symptoms of highest fatality
- 21.** Perform analysis on the fact that diseases with symptoms of upper abdominal pain whose fatality is greater than 2
- 22.** Perform analysis to calculate the Death percentage in top 3 Diseases.

Correlation data on the basis of various parameters(pre-data cleaning) :

1. Perform analysis on diseases which are most untraceable
 - ❖ Diseases vs symptom id: 0.6
2. Perform analysis on the drug related diseases and their fatality
 - ❖ Diseases vs fatality: -0.11
3. Perform analysis to find the easiest Drug Related Diseases
 - ❖ Diseases vs symptom id: 0.6
4. Perform analysis to find the most curable cancer
 - ❖ Diseases vs fatality: -0.11
5. Perform analyses to find the most fatal cancer
 - ❖ Diseases vs fatality: -0.11
6. Perform analysis to find the top 10 Severe cancer type.
 - ❖ Diseases vs fatality: -0.11
7. Perform analysis to find breast related diseases with their specific symptoms
 - ❖ Diseases vs symptom id: 0.6
8. Perform analysis to find the number of diseases whose fatality is 1 or 2
 - ❖ Disease ID vs fatality : -0.19
9. Perform analysis on the fact to find possible diseases if the symptoms are delusions and hallucinations.
 - ❖ Diseases vs symptoms: -0.03
10. Perform analysis on the fact that diseases which have symptoms of both headache and fever
 - ❖ Diseases vs symptoms: -0.03
11. Perform analysis on the fact that Showing, the diseases based on fatality rate
 - ❖ Diseases vs fatality: -0.11
12. Perform analysis on the fact that the number of Bacterial and cervical diseases and their fatality rate

❖ Diseases vs fatality: -0.11

13. Perform analysis on the fact that the symptoms related to eye (irritation) and has a fatality rate equal to 3

❖ Diseases vs fatality: -0.11

14. Perform analysis on the fact that what is the chance of dying from cold symptoms(Graph)

❖ Symptom vs fatality: -0.11

15. Perform analysis on the fact that Diseases with more than twenty five symptoms

❖ Diseases vs symptoms: -0.03

16. Perform analysis on the fact that If delusions and hallucinations occur in a patient then it leads to mental illness

❖ Diseases vs symptoms: -0.03

17. Perform analysis on the fact that most common symptoms found in the diseases of highest fatality

❖ Symptom vs fatality: -0.11

18. Perform analysis on the fact that fatality rate of diseases which have symptoms of vomiting

❖ Symptom vs fatality: -0.11

19. Perform analysis on the fact that the symptoms related to different types of organ failures and has a fatality rate equal to 3

❖ Disease ID vs fatality : -0.19

20. Perform analysis to find the occurrence of symptoms of highest fatality

❖ Symptom vs fatality: -0.11

21. Perform analysis on the fact that diseases with symptoms of upper abdominal pain whose fatality is greater than 2

❖ Symptom vs fatality: -0.11

22. Perform analysis to calculate the Death percentage in top 3 Diseases.

❖ Diseases vs fatality: -0.11

Correlation data on the basis of various parameters(post-data cleaning) :

1. Perform analysis on diseases which are most untraceable
 - ❖ Diseases vs symptom id: 0.42
2. Perform analysis on the drug related diseases and their fatality
 - ❖ Diseases vs fatality: -0.10
3. Perform analysis to find the easiest Drug Related Diseases
 - ❖ Diseases vs symptom id: 0.42
4. Perform analysis to find the most curable cancer
 - ❖ Diseases vs fatality: -0.10
5. Perform analyses to find the most fatal cancer
 - ❖ Diseases vs fatality: -0.10
6. Perform analysis to find the top 10 Severe cancer type.
 - ❖ Diseases vs fatality: -0.10
7. Perform analysis to find breast related diseases with their specific symptoms
 - ❖ Diseases vs symptom id: 0.42
8. Perform analysis to find the number of diseases whose fatality is 1 or 2
 - ❖ Disease ID vs fatality : -0.19
9. Perform analysis on the fact to find possible diseases if the symptoms are delusions and hallucinations.
 - ❖ Diseases vs symptoms: -0.06
10. Perform analysis on the fact that diseases which have symptoms of both headache and fever
 - ❖ Diseases vs symptoms: -0.06

11. Perform analysis on the fact that Showing, the diseases based on fatality rate

❖ Diseases vs fatality: -0.10

12 .Perform analysis on the fact that the number of Bacterial and cervical diseases and their fatality rate

❖ Diseases vs fatality: -0.10

13. Perform analysis on the fact that the symptoms related to eye (irritation) and has a fatality rate equal to 3

❖ Diseases vs fatality: -0.10

14. Perform analysis on the fact that what is the chance of dying from cold symptoms(Graph)

❖ Symptom vs fatality: -0.12

15. Perform analysis on the fact that Diseases with more than twenty five symptoms

❖ Diseases vs symptoms: -0.06

16. Perform analysis on the fact that If delusions and hallucinations occur in a patient then it leads to mental illness

❖ Diseases vs symptoms: -0.06

17. Perform analysis on the fact that most common symptoms found in the diseases of highest fatality

❖ Symptom vs fatality: -0.12

18. Perform analysis on the fact that fatality rate of diseases which have symptoms of vomiting

❖ Symptom vs fatality: -0.12

19. Perform analysis on the fact that the symptoms related to different types of organ failures and has a fatality rate equal to 3

❖ Disease ID vs fatality : -0.19

20. Perform analysis to find the occurrence of symptoms of highest fatality

❖ Symptom vs fatality: -0.12

21. Perform analysis on the fact that diseases with symptoms of upper abdominal pain whose fatality is greater than 2

❖ Symptom vs fatality: -0.12

22. Perform analysis to calculate the Death percentage in top 3 Diseases.

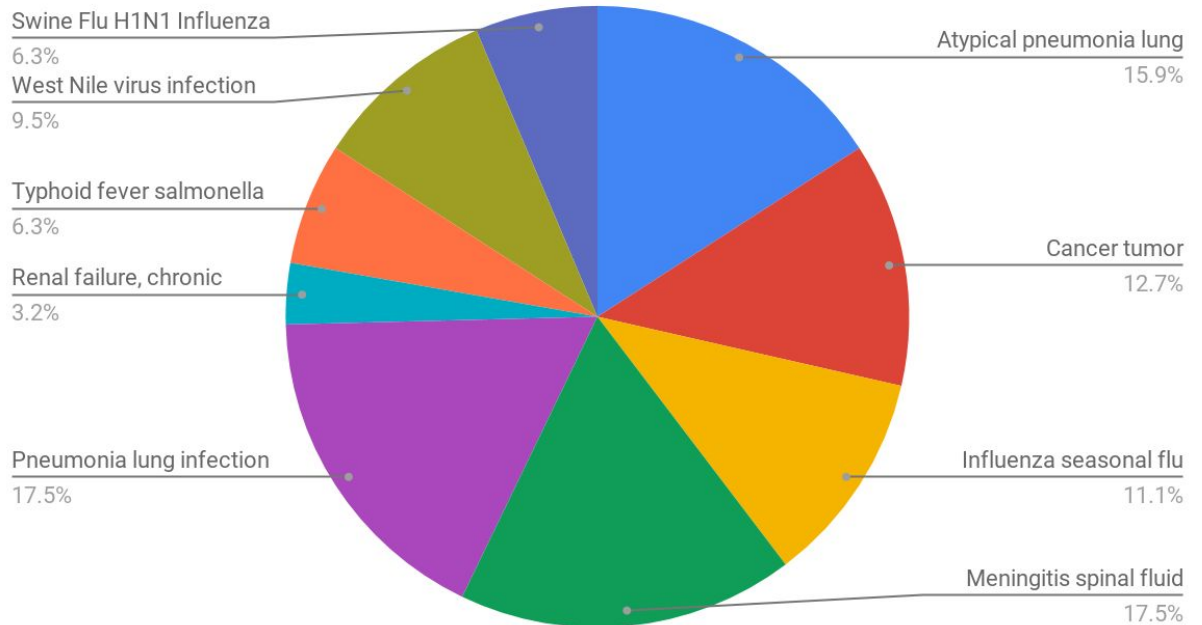
❖ Diseases vs fatality: -0.10

Analysis performed on the basis of various parameters with code, charts and inferences :

1. Perform analysis on diseases which are most untraceable

```
f=0
severeDiagSheet=pd.DataFrame()
unidentifiedDf.sort_values('Diagnosis Id', inplace=True, ascending=True)
group=unidentifiedDf.groupby('Diagnosis Id')
for diagId,diagDf in group:
    if diagDf.shape[0]>3:
        if f==0:
            severeDiagSheet=diagDf
            f=1
            continue
        severeDiagSheet=pd.concat([severeDiagSheet,diagDf])
print(severeDiagSheet)
severeDiagSheet.to_excel(r'E:\Python project files\notsevereDisease.xlsx')
```

Most Severe Disease



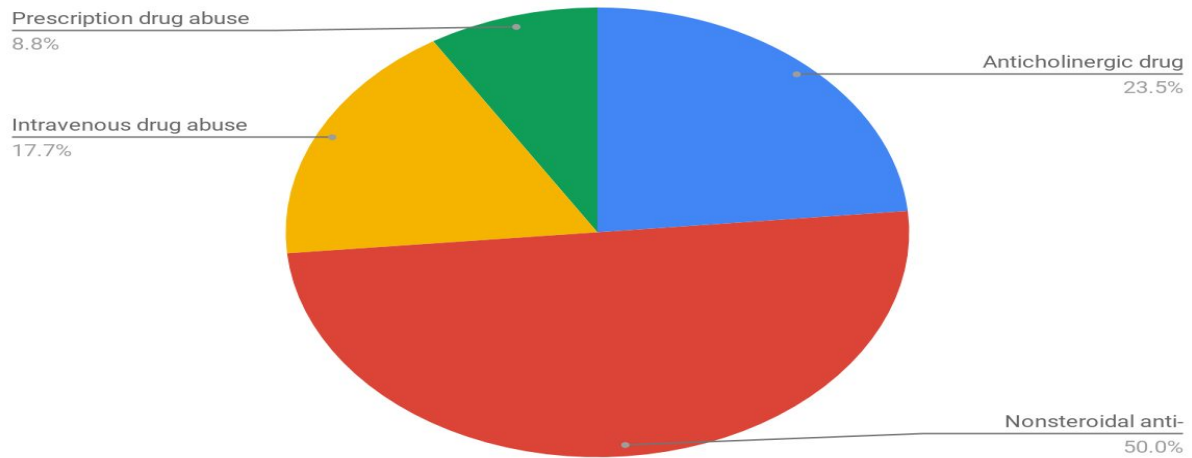
Inference: These are the diseases with most of the symptoms are unknown.

2. Perform analysis on the drug related diseases and their fatality

```
drugDisease=pd.read_excel(r'E:\Python project files\drugDisease.xlsx')
drugDisease.set_index("Disease ID",inplace=True)
print(drugDisease)
emptyindex=[]
for index ,rows in drugDisease.iterrows():
    emptyindex.append(index)
emptyIndexArray=np.asarray(emptyindex)
print(emptyIndexArray)
fatalitysheet.sort_values('Diagnosis Id', inplace=True, ascending=True)
fatalitysheet
fatalitysheetDiagnosisgroup=fatalitysheet.groupby("Diagnosis Id")
f=0
DrugSymptomSheet=pd.DataFrame()
for i in emptyIndexArray:
    if f==0:
        DrugSymptomSheet=fatalitysheetDiagnosisgroup.get_group(i)
        f=1
        continue
    r=fatalitysheetDiagnosisgroup.get_group(i)
    DrugSymptomSheet=pd.concat([DrugSymptomSheet,r])
#print(DrugSymptomSheet)
#DrugSymptomSheet.to_excel(r'E:\Python project files\DrugSymptomSheet.xlsx')
```

Disease ID	diagnose
48	Anticholinergic drug overdose
379	Nonsteroidal anti-inflammatory drug overdose M...
398	Intravenous drug abuse IVDA
601	Prescription drug abuse

Most fatal Drug diagnosis



Inference : The most fatal drug diagnosis is **Nonsteroidal anti-inflammatory drug overdose Motrin, Advil** with fatality rate of **2.833**

3. Perform analysis to find the easiest Drug Related Disease

```
#diseaseSheet=pd.read_excel(r'E:\Python project files\Diagnosis name.xlsx')
#diseaseSheet.set_index('Disease ID',inplace=True)
#drugDisease=diseaseSheet[diseaseSheet['diagnose'].str.contains('drug')]
drugDisease=pd.read_excel(r'E:\Python project files\drugDisease.xlsx')
drugDisease.set_index("Disease ID",inplace=True)
print(drugDisease)
emptyindex=[]
for index ,rows in drugDisease.iterrows():
    emptyindex.append(index)
emptyIndexArray=np.asarray(emptyindex)
print(emptyIndexArray)
fatalitysheet.sort_values('Diagnosis Id', inplace=True, ascending=True)
fatalitysheet
fatalitysheetDiagnosisgroup=fatalitysheet.groupby("Diagnosis Id")
f=0
DrugSymptomSheet=pd.DataFrame()
for i in emptyIndexArray:
    if f==0:
        DrugSymptomSheet=fatalitysheetDiagnosisgroup.get_group(i)
        f=1
        continue
    r=fatalitysheetDiagnosisgroup.get_group(i)
    DrugSymptomSheet=pd.concat([DrugSymptomSheet,r])
#print(DrugSymptomSheet)
#DrugSymptomSheet.to_excel(r'E:\Python project files\DrugSymptomSheet.xlsx')
```

	Symptom Id	Diagnosis Id	Fatality Rate
1803	77	48	3
1712	148	48	1
1785	153	48	2
2292	192	48	1
2731	223	48	1
2811	226	48	1
2810	227	48	1
2882	228	48	1
2883	229	48	1
1852	9	379	3
788	71	379	3
840	77	379	3
1794	153	379	3
1842	156	379	2
1845	157	379	3
869	79	398	NaN
3628	4	601	NaN
3640	5	601	NaN
3800	15	601	NaN
3842	19	601	NaN
3887	21	601	NaN
4078	38	601	NaN
4264	66	601	NaN
1046	96	601	1
4450	113	601	NaN
4492	121	601	NaN
4677	275	601	NaN

Inference : **Prescription Drug abuse** is the easiest to diagnose with 11 visual symptoms.

4. Perform analysis to find the most curable cancer.

```
CancerSheet=pd.read_excel(r'E:\Python project files\CancerDisease.xlsx')
CancerSheet.set_index('Disease ID',inplace=True)
emptyindex=[]
for index ,rows in CancerDisease.iterrows():
    emptyindex.append(index)
emptyIndexArray=np.asarray(emptyindex)
print(emptyIndexArray)
fatalitySheet=pd.read_excel(r'E:\Python project files\fatality sheet.xlsx')
fatalitySheet
fatalityGroupSheet=fatalitySheet.groupby('Diagnosis Id')
f=0
for i in emptyIndexArray:
    if f==0:
        CancerDf=fatalityGroupSheet.get_group(i)
        f=1
        continue
    r=fatalityGroupSheet.get_group(i)
    CancerDf=pd.concat([CancerDf,r])
print(CancerDf)
g=0
CancerGroupdf=CancerDf.groupby('Diagnosis Id')
for i,idf in CancerGroupdf:
    if g==0:
        largestCancerRow=idf
        g=1
        continue
    if idf.shape[0] > largestCancerRow.shape[0]:
        largestCancerRow=idf
print(largestCancerRow)
```

The symptom table of the most curable disease:

-	Symptom Id	Diagnosis Id	Fatality Rate
3368	12	1119	0.0
3370	25	1119	0.0
3372	35	1119	0.0
3374	196	1119	0.0
3376	42	1119	0.0
3380	45	1119	0.0
3382	57	1119	0.0
3383	58	1119	0.0
3385	251	1119	0.0
3387	89	1119	0.0
3389	233	1119	0.0
3391	118	1119	0.0
3393	217	1119	0.0
3396	177	1119	0.0
3397	245	1119	0.0
3398	6	1119	0.0
3399	213	1119	0.0
3401	234	1119	0.0
4689	275	1119	0.0

Inference : **Sarcoma soft tissue cancer** is the most curable cancer with 19 visual symptoms and with fatality value 0.0

5. Perform analysis to find the most fatal cancer.

```
CancerSheet=pd.read_excel(r'E:\Python project files\CancerDisease.xlsx')
CancerSheet.set_index('Disease ID',inplace=True)
emptyindex=[]
for index ,rows in CancerDisease.iterrows():
    emptyindex.append(index)
emptyIndexArray=np.asarray(emptyindex)
print(emptyIndexArray)
fatalitySheet=pd.read_excel(r'E:\Python project files\fatality sheet.xlsx')
fatalitySheet
fatalityGroupSheet=fatalitySheet.groupby('Diagnosis Id')
f=0
for i in emptyIndexArray:
    if f==0:
        CancerDf=fatalityGroupSheet.get_group(i)
        f=1
        continue
    r=fatalityGroupSheet.get_group(i)
    CancerDf=pd.concat([CancerDf,r])
print(CancerDf)
g=0
CancerGrouppdf=CancerDf.groupby('Fatality Rate')
for i,idf in CancerGrouppdf:
    if g==0:
        largestCancerRow=idf
        g=1
        continue
    if idf.shape[0] < largestCancerRow.shape[0]:
        largestCancerRow=idf
print(largestCancerRow)
```

	Symptom Id	Diagnosis Id	Fatality Rate
1023	94	107	3.0
1129	104	107	3.0
1590	137	107	3.0
1924	164	107	3.0
1950	165	107	3.0
3302	246	107	3.0
3434	255	107	3.0
4817	294	107	3.0

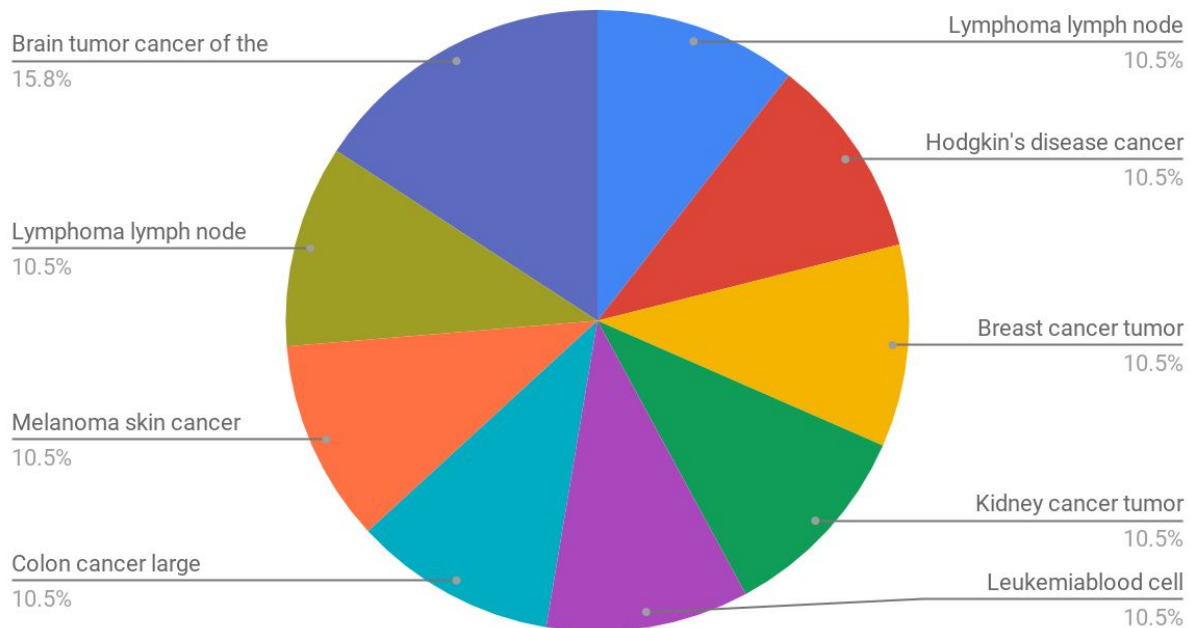
Inference : The most fatal type of cancer is brain tumor or commonly known as cancer of the brain

6. Perform analysis to find the top 10 Severe cancer type.

```
CancerSheet=pd.read_excel(r'E:\Python project files\CancerDisease.xlsx')
CancerSheet.set_index('Disease ID',inplace=True)
emptyindex=[]
for index ,rows in CancerDisease.iterrows():
    emptyindex.append(index)
emptyIndexArray=np.asarray(emptyindex)
print(emptyIndexArray)
fatalitySheet=pd.read_excel(r'E:\Python project files\fatality sheet.xlsx')
fatalitySheet
fatalityGroupSheet=fatalitySheet.groupby('Diagnosis Id')
f=0
for i in emptyIndexArray:
    if f==0:
        CancerDf=fatalityGroupSheet.get_group(i)
        f=1
        continue
    r=fatalityGroupSheet.get_group(i)
    CancerDf=pd.concat([CancerDf,r])
print(CancerDf)
g=0
CancerGroupdf=CancerDf.groupby('Fatality Rate')
for i,idf in CancerGroupdf:
    if g==0:
        largestCancerRow=idf
        g=1
        continue
    if idf.shape[0] < largestCancerRow.shape[0]:
        largestCancerRow=idf
print(largestCancerRow)
```

	Symptom Id	Diagnosis Id	Fatality Rate
1438	131	Lymphoma lymph node cancer	2
2124	180	Hodgkin's disease cancer of the lymph system	2
113	11	Breast cancer tumor	2
1690	147	Kidney cancer tumor	2
2127	180	Leukemiablood cell cancer	2
2644	218	Colon cancer large intestine tumor	2
2416	201	Colon cancer large intestine tumor	2
182	16	Colon cancer large intestine tumor	2
2036	174	Melanoma skin cancer	2
2492	206	Melanoma skin cancer	2
4826	295	Breast cancer tumor	2
3562	287	Breast cancer tumor	2
2125	180	Lymphoma lymph node cancer	2
1924	164	Brain tumor cancer of the brain	3
4817	294	Brain tumor cancer of the brain	3
1129	104	Brain tumor cancer of the brain	3
1023	94	Brain tumor cancer of the brain	3
1950	165	Brain tumor cancer of the brain	3
3302	246	Brain tumor cancer of the brain	3
3434	255	Brain tumor cancer of the brain	3
1590	137	Brain tumor cancer of the brain	3

Top Severe Cancer Disease



Inference : Most of the cancer whose fatality is 2 are quite untraceable since they have very few symptoms

7. Perform analysis to find breast related diseases with their specific symptoms

```
tempdf=dia.loc[dia['Disease'].str.match('Breast')]
tempdf2=data.loc[data['Disease ID'].isin(tempdf['Disease ID'])]
tempdf3=sym.loc[sym['Symptom ID'].isin(tempdf2['Symptom ID'])]
c=0
d=0
g=0
f=0
for i in tempdf2["Fatality"]:
    if i==1:
        c=c+1
    if i==2:
        d=d+1
    if i==3:
        f=f+1
    if i==0:
        g=g+1
print(c,f,g,d)

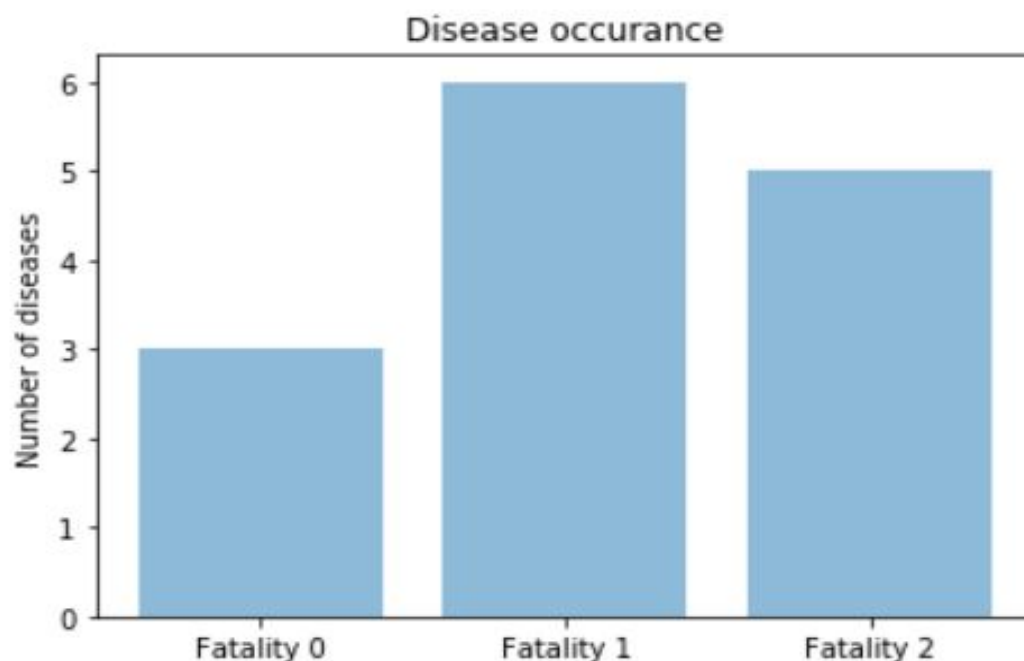
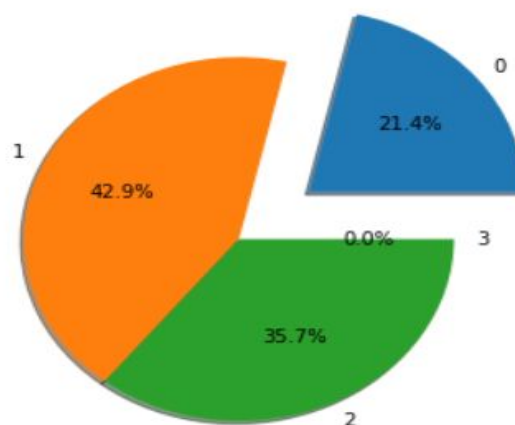
expval=[g,c,d,f]
exphead=[0,1,2,3]
plt.axis("equal")
plt.pie(expval,labels=exphead,radius=1.2,autopct='%0.1f%%',shadow=True,explode=[0.5,0,0,0])
plt.show()

l4=[]
grp=tempdf2.groupby('Fatality')
for d,group in grp:
    i=(group.shape[0])
    l3=[i]
    l4.extend(l3)
arr1=np.asarray(l4)
objects1=('Fatality 0','Fatality 1','Fatality 2')
y_pos=np.arange(len(objects1))

plt.bar(y_pos,arr1,align='center',alpha=0.5)
plt.xticks(y_pos,objects1)
plt.ylabel('Number of diseases')
plt.title('Disease occurance')
plt.show()
```

	Disease ID		Disease
101	108	Breast abscess□collection of pus in the breast	
102	109	Breast cancer□tumor	
103	110	Breast fat necrosis□dead breast fat	
104	111	Breast fibroadenoma□benign breast lumps	
1158	1523	Breast cyst	
Symptom ID	Disease ID	Fatality	
26	3	108	0.0
112	11	108	1.0
113	11	109	2.0
114	11	111	1.0
3560	287	111	1.0
3562	287	109	2.0
3566	287	108	0.0
3867	21	109	1.0
4577	262	109	1.0
4818	295	108	2.0
4819	295	110	0.0
4821	295	111	2.0
4826	295	109	2.0
5160	287	1523	1.0

6 0 3 5



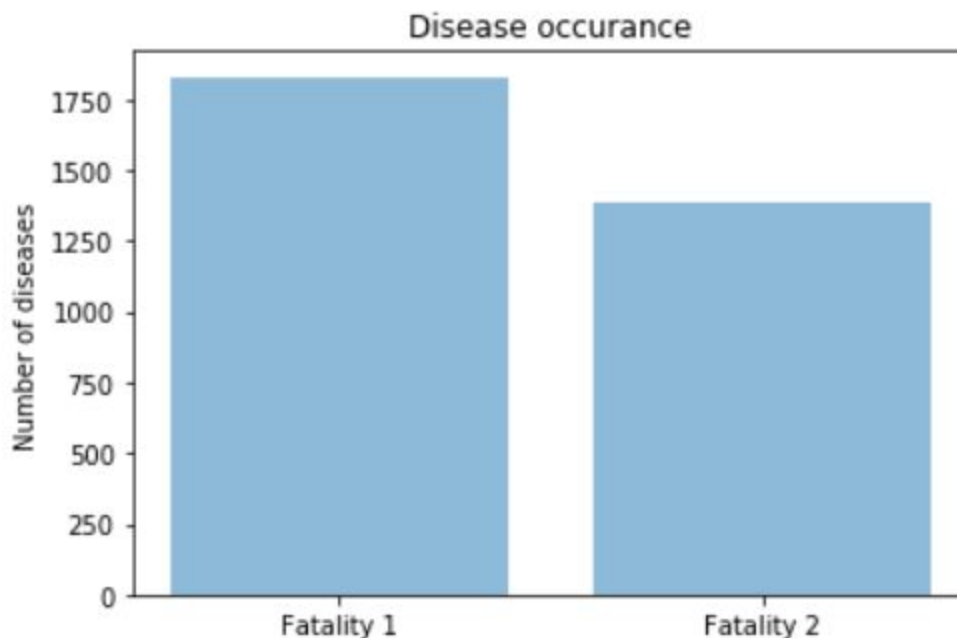
Inference : Most of the breasts related diseases have minimal fatality

8. Perform analysis to find the number of diseases whose fatality is 1 or 2

```
arr=[1,2]
tempdf6=data.loc[data["Fatality"].isin(arr)]
tempdf7=dia.loc[dia["Disease ID"].isin(tempdf6["Disease ID"])]
print(tempdf6)

l2=[]
grp=tempdf6.groupby('Fatality')
for d,group in grp:
    i=(group.shape[0])
    l1=[i]
    l2.extend(l1)
arr=np.asarray(l2)
objects=('Fatality 1','Fatality 2')
y_pos=np.arange(len(objects))

plt.bar(y_pos,arr,align='center',alpha=0.5)
plt.xticks(y_pos,objects)
plt.ylabel('Number of diseases')
plt.title('Disease occurance')
plt.show()
```



Inference : Most of the diseases have minimal fatality or fatality equal to one

9. Perform analysis on the fact to find possible diseases if the symptoms are delusions and hallucinations.

```
arr2=['Delusions','hallucinations']
empdf=sym.loc[sym['Symptom'].str.contains('|'.join(arr2))]
empdf1=data.loc[data['Symptom ID'].isin(empdf['Symptom ID'])]
empdf2=dia.loc[dia['Disease ID'].isin(empdf1['Disease ID'])]
print(empdf1['Fatality'])

l4=[]
grp1=empdf1.groupby('Fatality')
for d1,group1 in grp1:
    i1=(group1.shape[0])
    l3=[i1]
    l4.extend(l3)
arr1=np.asarray(l4)
objects=('Fatality 0','Fatality 1','Fatality 2','Fatality 3')
y_pos=np.arange(len(objects))

plt.bar(y_pos,arr1,align='center',alpha=0.5)
plt.xticks(y_pos,objects)
plt.ylabel('Number of diseases')
plt.title('Disease occurrence')
plt.show()

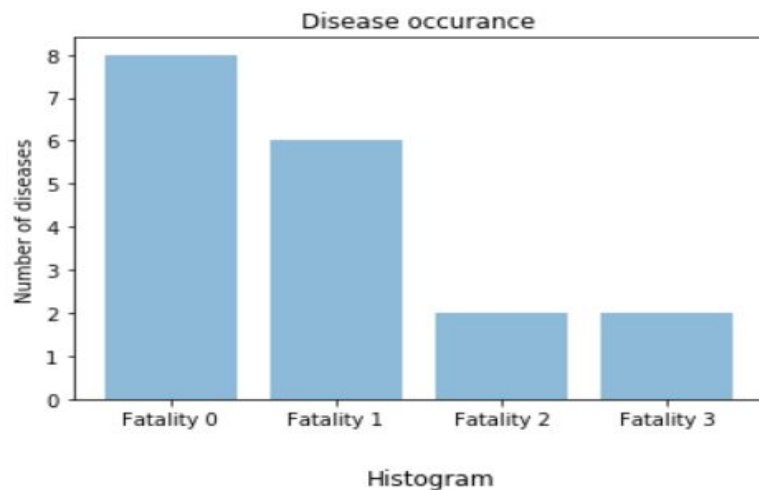
#Histogram
population_age = [2,2,0,1,3,3,0,0,0,1,1,1,0,1,1]
bins = [0,2,4]
plt.hist(population_age, bins, histtype='bar',rwidth=0.2)
plt.xlabel('age groups')
plt.ylabel('Number of people')
plt.title('Histogram')
plt.show()

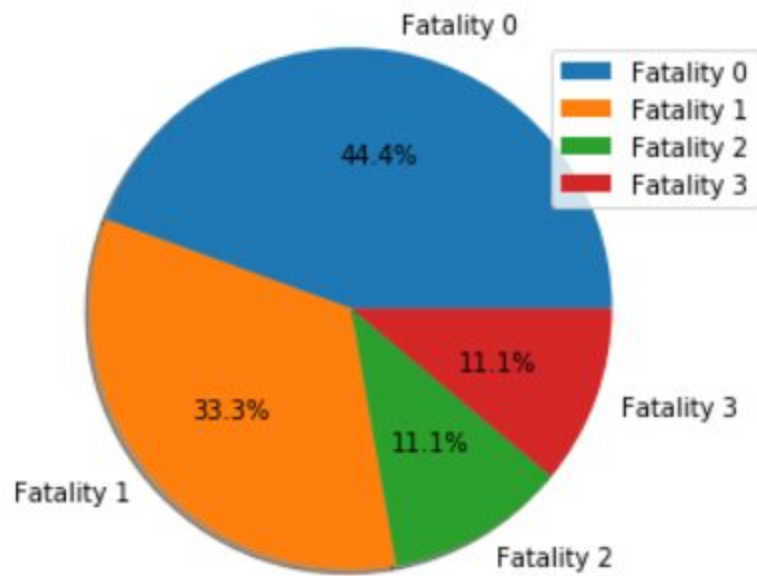
expval=[5,6,2,2]
exphead=['Fatality 0','Fatality 1','Fatality 2','Fatality 3']
plt.axis("equal")
plt.pie(expval,labels=exphead,radius=1.2,autopct='%0.1f%%',shadow=True,explode=[0,0,0,0])
plt.legend()
plt.show()
```

443	2.0
444	2.0
445	0.0
446	1.0
447	3.0
448	3.0
4070	0.0
4071	0.0
4072	0.0
4073	1.0
4074	1.0
4075	0.0
4076	1.0
4077	0.0
4078	0.0
4079	0.0
4080	1.0
5141	1.0

Name: Fatality, dtype: float64

	Disease ID	Disease
18	19	Adjustment disorder (poor adjustment to life ...
22	23	Alcohol withdrawal syndrome (mild)
23	25	Alcoholism
31	33	Amphetamine abuse
48	51	Anxiety disordergeneralized anxiety disorderGAD
88	93	Bipolar disordermanic depressive disorder
165	175	Cocaine abuse
202	212	Depressionexcessive sadness
217	227	Drug reaction
414	444	Magic mushroom ingestionpsilocybin
415	445	Major depressive disordersevere depression
553	595	Post-traumatic stress disorderPTSD
558	601	Prescription drug abuse
631	676	Schizoaffective disorderfeatures of schizophr...
632	677	Schizophreniachronic impaired reality perception
700	749	Temporal lobe epilepsynon-convulsive seizure
1004	1112	Hepatic encephalopathyconfusion from liver fa...
1112	1403	Delusional disorder





Inference : Most Diseases related delusions and hallucinations are much less fatal. As understood from the above bar graph.

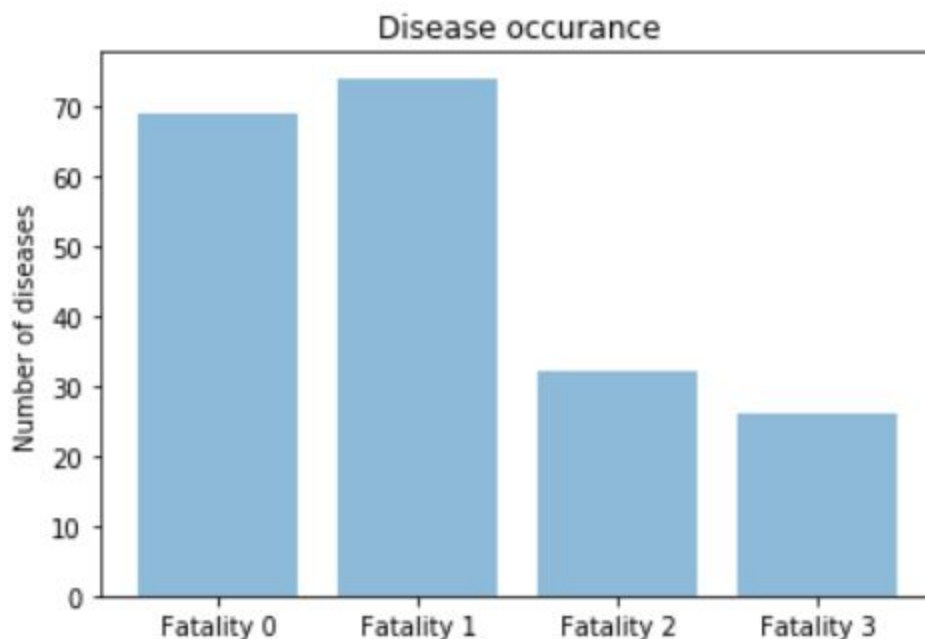
10.Perform analysis on the fact that diseases which have symptoms of both headache and fever

```
sym.fillna('no',inplace=True)
arr2=['Headache','Fever']
empdf=sym.loc[sym['Symptom'].str.contains('|'.join(arr2))]
empdf1=data.loc[data['Symptom ID'].isin(empdf['Symptom ID'])]
empdf2=dia.loc[dia['Disease ID'].isin(empdf1['Disease ID'])]
print(empdf2)

l4=[]
grp1=empdf1.groupby('Fatality')
for d1,group1 in grp1:
    i1=(group1.shape[0])
    l3=[i1]
    l4.extend(l3)
arr1=np.asarray(l4)
objects=('Fatality 0','Fatality 1','Fatality 2','Fatality 3')
y_pos=np.arange(len(objects))

plt.bar(y_pos,arr1,align='center',alpha=0.5)
plt.xticks(y_pos,objects)
plt.ylabel('Number of diseases')
plt.title('Disease occurrence')
plt.show()
```

Disease ID	Disease
3	4 Abrasions (scrapes)
23	25 Alcoholism
25	27 Allergic rhinitis (allergic reaction in the n...
30	32 Amebiasis (ameba infection)
31	33 Amphetamine abuse
48	51 Anxiety disorder (generalized anxiety disorder) GAD
65	69 Atypical pneumonia (lung infection)
66	70 Autoimmune conditions
71	75 Bacterial infection
79	83 Basilar skull fracture (broken skull)
83	87 Bell's palsy (facial muscle weakness)
88	93 Bipolar disorder (manic depressive disorder)
100	107 Brain tumor (cancer of the brain)
108	115 Bronchitis (bronchial tube infection)
118	125 Campylobacter infection (intestinal bacterial i...
119	126 Cancer (tumor)
121	128 Carbon monoxide poisoning (odorless, poisonous gas)
124	131 Carotid artery dissection (neck artery tear)
128	135 Cavernous sinus aneurysm (head vein dilation)
129	136 Cavernous sinus thrombosis (head vein blood clot)
130	137 Cavernous sinus tumor (head vein cancer)
131	138 Cavity (tooth caries)
133	140 Cellulitis (skin infection)
134	142 Cerebellar hemorrhage (bleeding in back of brain)
136	144 Cerebral contusion (bruise of brain)
138	146 Cerebral vascular accident (stroke)
139	147 Cerebrospinal fluid rhinorrhea (leakage of brai...
145	153 Cervical spondylosis (neck arthritis)
147	155 Chagas disease (trypanosomiasis)
153	163 Cholecystitis (inflammation of the gallbladder)
...	...
1029	1155 Tension headache (stress headache)
1049	1261 Mycoplasma infection (bacteria)
1053	1269 Essential thrombocythemia (excessive blood plat...
1055	1273 Herpangina (mouth blisters)
1056	1275 Ewing's sarcoma (cancer)
1057	1277 Extragonadal germ cell tumors
1058	1279 Fallopian tube cancer
1060	1283 Farsightedness (hyperopia/hypermétropia)
1063	1289 Foot ulcer
1066	1295 Gallbladder and bile duct cancer (Gallbladder c...
1069	1301 Head injury in children (Non-accidental traumat...
1070	1303 Hemorrhagic stroke (CVA, cerebrovascular accident)
1071	1305 Hepatitis A



Inference : Most of the diseases with the symptoms of headache and fever have minimal fatality

11. Perform analysis on the fact that Showing, the diseases based on fatality rate

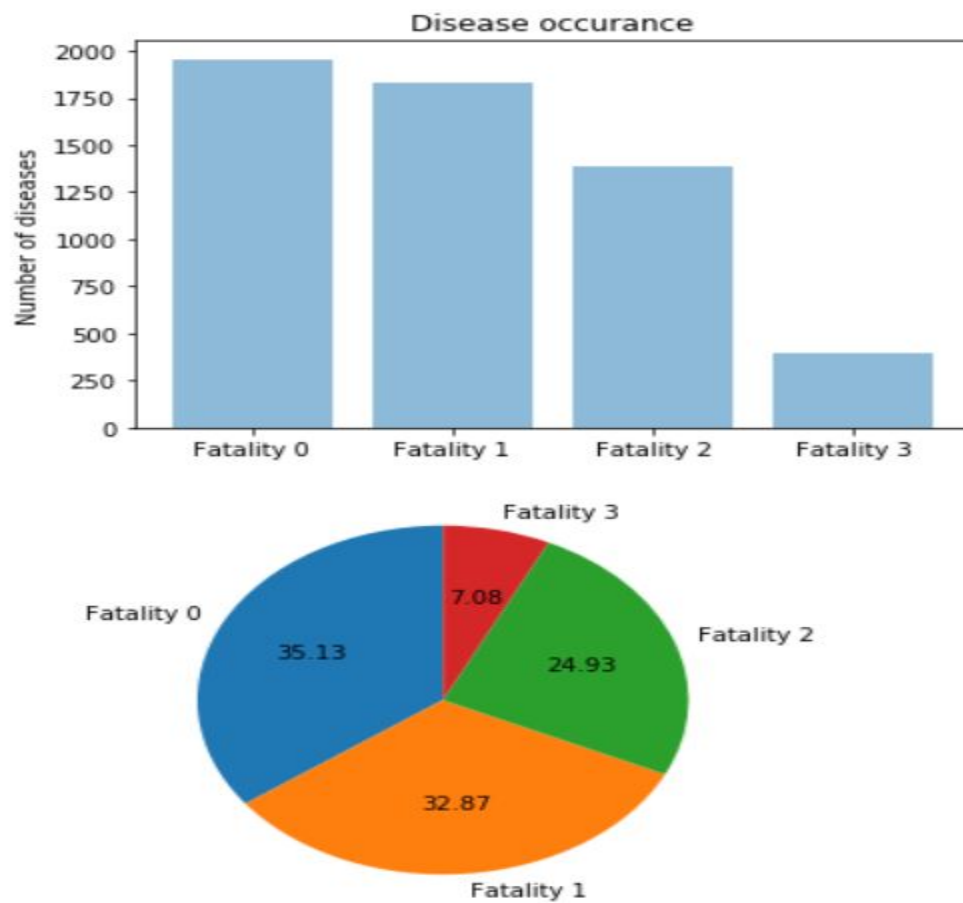
```
grp1=data.groupby('Fatality')
l4=[]
for d,group in grp1:
    i1=(group.shape[0])
    l3=[i1]
    l4.extend(l3)
arr1=np.asarray(l4)
#print(arr1)
objects=('Fatality 0','Fatality 1','Fatality 2','Fatality 3')
y_pos=np.arange(len(objects))

plt.bar(y_pos,arr1,align='center',alpha=0.5)
plt.xticks(y_pos,objects)
plt.ylabel('Number of diseases')
plt.title('Disease occurance')
plt.show()

figureObjects, axesObject = plt.subplots()

axesObject.pie(arr1,
               labels=objects,
               autopct='%1.2f',
               startangle=90)

axesObject.axis('equal')
plt.show()
```

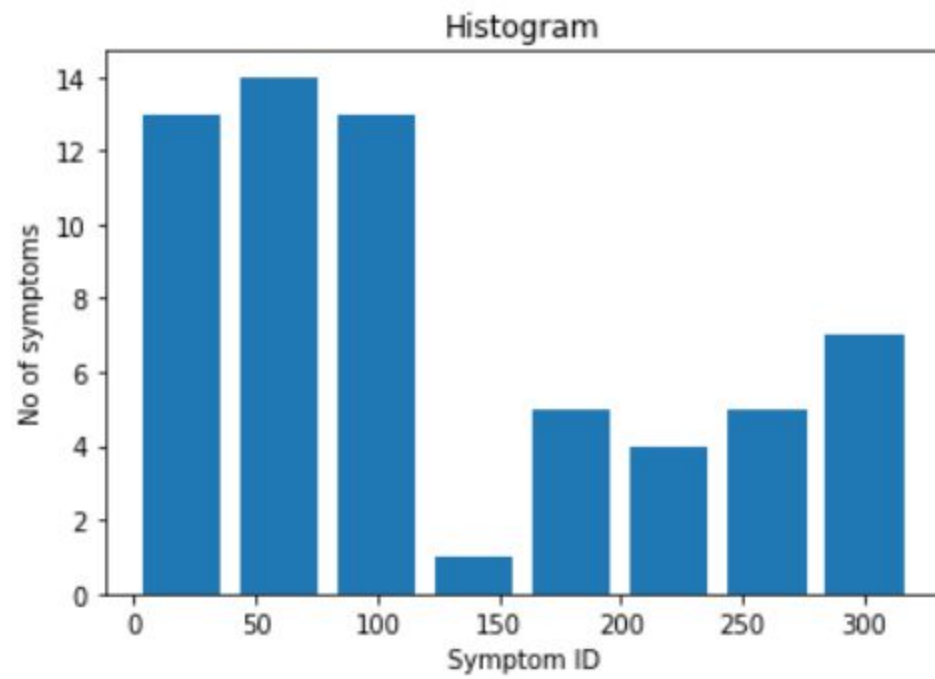
Inference : Most of the diseases are curable and does not cause any sort of major health issues

12. Perform analysis on the fact that the number of Bacterial and cervical diseases and their fatality rate using histogram

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt

#data=pd.read_csv("datamatch.csv")
#dia=pd.read_csv("diagnose.csv")
#sym=pd.read_csv("symptom.csv")
dia.rename(columns = {"did": "Disease ID", "diagnose": "Disease"}, inplace = True)
sym.rename(columns = {"syd": "Symptom ID", "symptom": "Symptom"}, inplace = True)
data.rename(columns = {"did": "Disease ID", "syd": "Symptom ID", "wei": "Fatality"}, inplace = True)
search_values=['Bacterial', 'Cervical']
tempdf4=dia.loc[dia['Disease'].str.contains('|'.join(search_values))]
tempdf5=data.loc[data["Disease ID"].isin(tempdf4["Disease ID"])]
#print(tempdf5)
print(tempdf4)
l4=[]
for i in tempdf5['Symptom ID']:
    l3=[i]
    l4.extend(l3)
#print(l4)
population_age = l4
bins = [0,40,80,120,160,200,240,280,320]
plt.hist(population_age, bins, histtype='bar',rwidth=0.8)
plt.xlabel('age groups')
plt.ylabel('Number of people')
plt.title('Histogram')
plt.show()
```

	Disease ID	Disease
70	74	Bacterial dysentery□bacterial infection of the...
71	75	Bacterial infection
72	76	Bacterial overgrowth of small intestine□Bacter...
73	77	Bacterial vaginosis□vaginal infection
141	149	Cervical cancer□tumor
142	150	Cervical lymphadenopathy□enlarged neck lymph n...
143	151	Cervical radiculopathy□pinched nerve in neck
144	152	Cervical spine fracture□broken neck bone
145	153	Cervical spondylosis□neck arthritis
288	305	Bacterial vaginosis□BV, garnerella vaginalis
799	862	Cervical spine stenosis□narrowing of spinal canal
800	863	Cervical myelopathy□spinal cord compression
1143	1487	Bacterial tracheitis



Inference : Neck stiffness or tightness is the most common symptom in bacterial and cervical diseases.

13. Perform analysis on the fact that the symptoms related to eye (irritation) and has a fatality rate equal to 3

```
dies=pd.read_excel("Disease.xlsx")
fatal=pd.read_excel("Disease_fatality.xlsx")
tempdf=sym.loc[sym['symptom'].str.contains('|'.join(arr))]
#print(tempdf)
u=3
l2=[]
tempdf.set_index('syd',inplace=True)
print(tempdf)
for i,group in tempdf.iterrows():
    l1=[i]
    l2.extend(l1)
print(l2)
l3=[]
tempdf1=fatal.groupby('syd')
for i in l2:
    for j,group1 in tempdf1:
        if i==j:
            if(i==160 and i==198):
                df1=tempdf1.get_group(i)
                df2=df1.groupby('wei')
                df3=df2.get_group(u)
                p=df3.shape[0]
                l4=[p]
                l3.extend(l4)

l6=[]
l3.insert(1,0)
l3.insert(3,0)
print(l3)
sym.set_index('syd',inplace=True)
for j in l2:
    for i,gro in sym.iterrows():
        if i==j:
```

```
            if i==j:
                l5=[gro['symptom']]
                l6.extend(l5)
rr1=np.asarray(l6)
bjects =arr1
_pos = np.arange(len(objects))
lt.figure(figsize=(12, 7))
lt.bar(y_pos,l3, align='center', alpha=1)
lt.xticks(y_pos, objects)
lt.ylabel('Number of Diseases with a Particular fatality frequency')
lt.xlabel('Symptoms')
lt.title('')
lt.show()
rr2=arr1
figureObject, axesObject = plt.subplots()
xesObject.pie(l3,

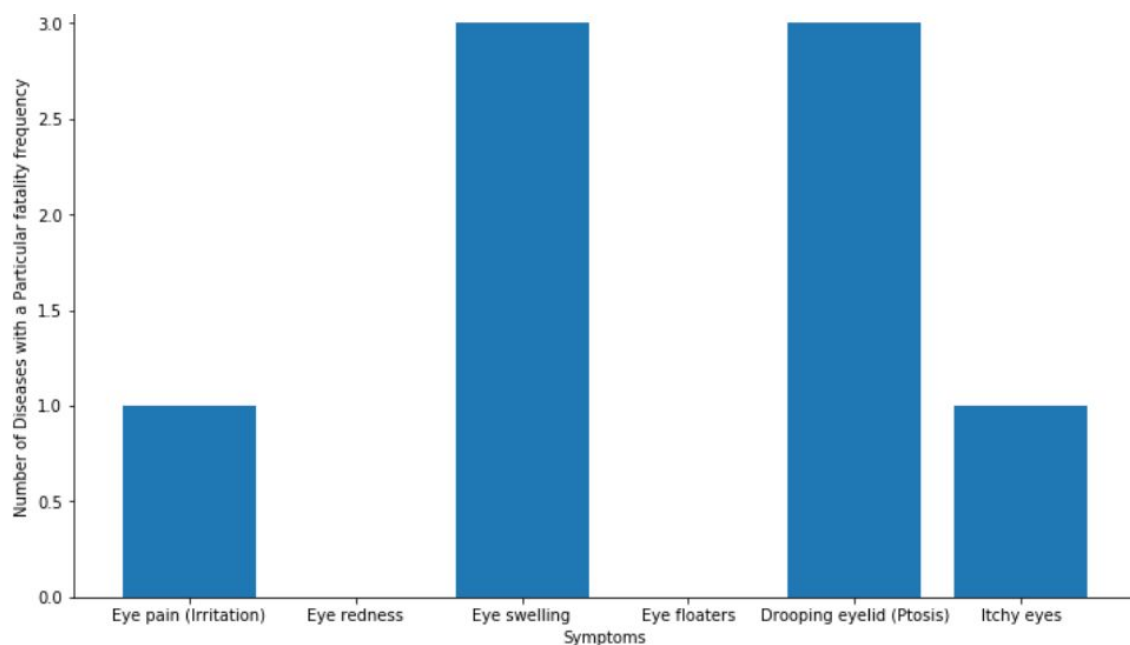
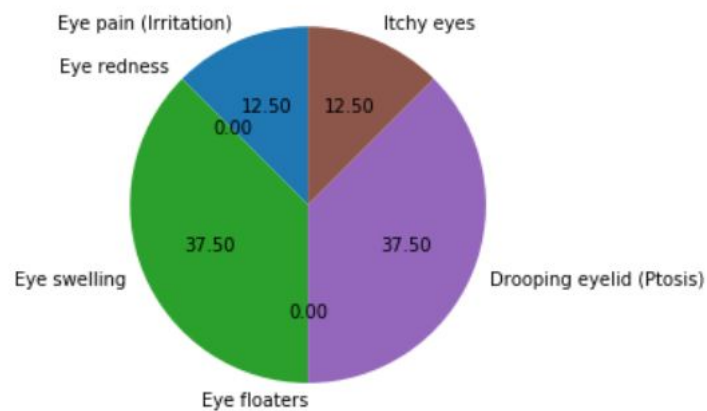
                labels=arr2,

                autopct='%1.2f',

                startangle=90)

xesObject.axis('equal')
lt.show()
```

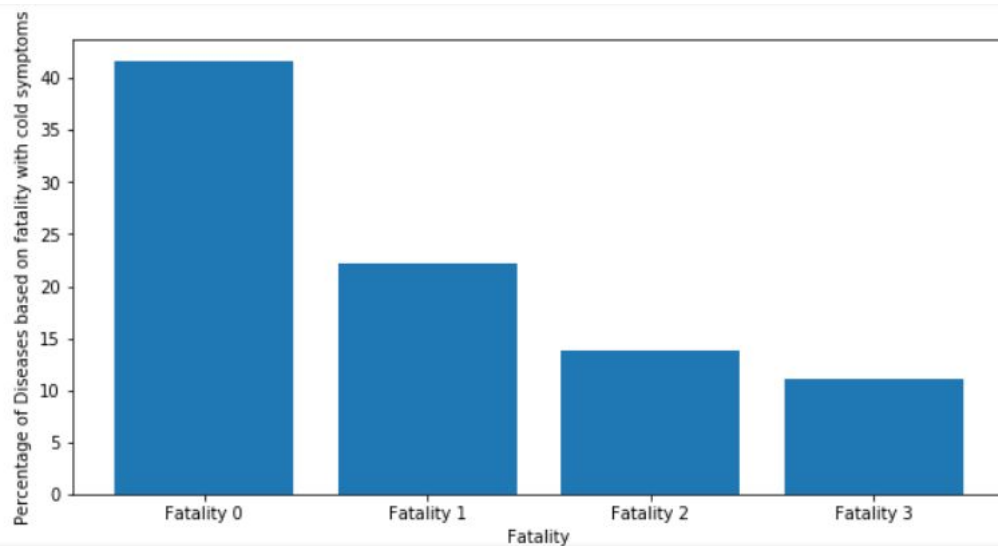
syd	symptom
26	Eye pain (Irritation)
160	Eye redness
195	Eye swelling
198	Eye floaters
255	Drooping eyelid (Ptosis)
259	Itchy eyes

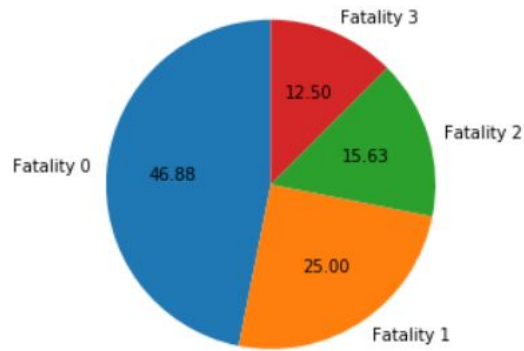


Inference : The most common eye related symptoms are “Eye swelling”, “Drooping eyelids” as they occur in 3 diseases.

14. Perform analysis on the fact that what is the chance of dying from cold symptoms(Graph)

```
tempdf=sym.loc[sym['symptom'].str.match('Cough')]
print(tempdf)
tempdf.set_index('syd',inplace=True)
for i,name in tempdf.iterrows():
    symid=i
    print(symid)
    dise=fatal.groupby('syd')
    coughdis=dise.get_group(symid)
    print(coughdis)
    coughdis1=coughdis.groupby('wei')
    for i,group in coughdis1:
        print(group)
        c=group.shape[0]
        print(c)
    fat0=(15/36)*100
    print(fat0)
    fat1=(8/36)*100
    print(fat1)
    fat2=(5/36)*100
    print(fat2)
    fat3=(4/36)*100
    print(fat3)
    l1=[fat0,fat1,fat2,fat3]
    arr1=np.asarray(l1)
    print(arr1)
    objects = ('Fatality 0','Fatality 1','Fatality 2','Fatality 3')
    y_pos = np.arange(len(objects))
    plt.figure(figsize=(10, 5))
    plt.bar(y_pos,arr1, align='center', alpha=1)
    plt.xticks(y_pos, objects)
    plt.ylabel('Percentage of Diseases based on fatality with cold symptoms')
    plt.xlabel('Fatality')
    plt.title('')
    plt.show()
    arr2=['Fatality 0','Fatality 1','Fatality 2','Fatality 3']
    figureObject, axesObject = plt.subplots()
    axesObject.pie(arr1,labels=arr2,autopct='%1.2f',startangle=90)
    axesObject.axis('equal')
    plt.show()
```





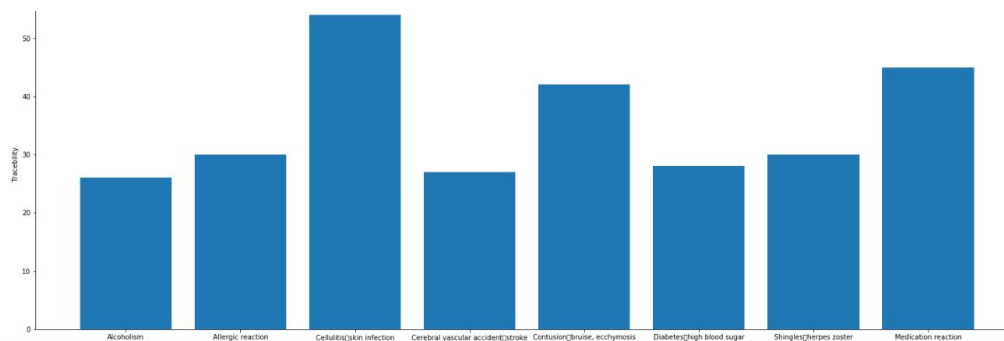
Inference : Cold symptoms do not pose much threat as most of the diseases related to cold symptoms have fatality rate 0.

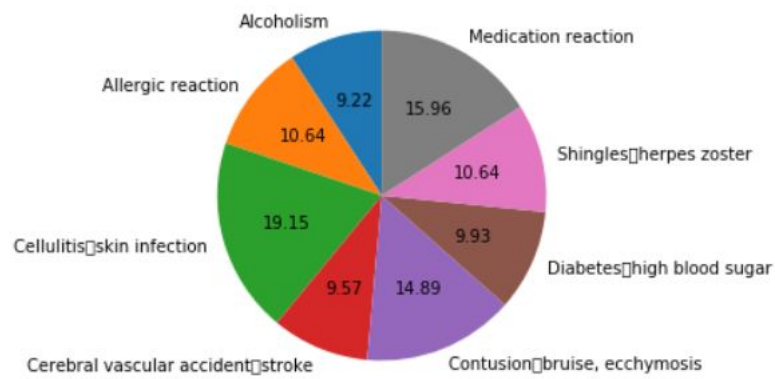
15. Perform analysis on the fact that Diseases with more than twenty five symptoms

```

disease1=fatal.groupby('did')
l2=[]
l4=[]
symp1=[]
arr2=[]
diseaisd=[25,26,140,146,191,213,698,1034]
for index,group in disease1:
    symno=group.shape[0]
    if symno>25:
        symp=[symno]
        symp1.extend(symp)
        l1=[group['did']]
        l2.extend(l1)
arr=np.asarray(l2)
arr2=np.asarray(symp1)
print(arr2)
print(arr)
dies.set_index('did',inplace=True)
j=0
for j in diseaisd:
    for i,gro in dies.iterrows():
        if i==j:
            l3=[gro['diagnose']]
            l4.extend(l3)
arr1=np.asarray(l4)
objects = arr1
y_pos = np.arange(len(objects))
plt.figure(figsize=(26, 9))
plt.bar(y_pos,arr2, align='center', alpha=1)
plt.xticks(y_pos, objects)
plt.ylabel('Traceability')
plt.xlabel('Diseases above symptoms 25')
plt.title('')
plt.show()
figureObject, axesObject = plt.subplots()
axesObject.pie(arr2,labels=arr1,autopct='%1.2f',startangle=90)
axesObject.axis('equal')
plt.show()

```





Inference : Diseases with Symptoms Above 25. Among them The easily Traceable Disease is Cellulitis Skin Infection

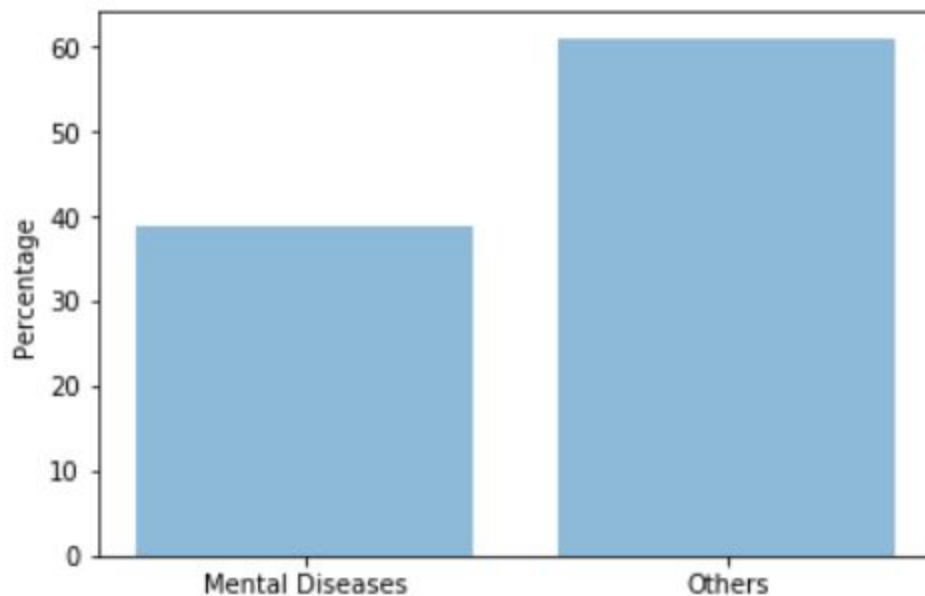
occur in a patient then it leads to mental illness.

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
data=pd.read_csv("datamatch.csv")
dia=pd.read_csv("diagnose.csv")
sym=pd.read_csv("symptom.csv")
sym.fillna('no',inplace=True)
arr2=["Delusion","hallucinations"]
empdf=sym.loc[sym['symptom'].str.contains('|'.join(arr2))]
print(empdf)
empdf1=data.loc[data['syd'].isin(empdf['syd'])]
print(empdf1)
empdf2=dia.loc[dia['did'].isin(empdf1['did'])]
print(empdf2)
dis=[19,51,93,212,445,595,1403]
c=0
for i in dis:
    c=c+1
print(c)
d=empdf2.count()
print(d)
avg1=(c/d*100)
avg2=(100-c/d*100)
arr1=np.array(avg1,avg2)
#print(arr1)
object=('Mental Diseases','Others')
y_pos=np.arange(len(object))

plt.bar(y_pos,[38.8888888889,61.1111111111],align='center',alpha=0.5)
plt.xticks(y_pos,object)
plt.ylabel('Percentage')
plt.show()
```

	syd	symptom	
	37	38	Delusions or hallucinations
		syd	did wei
443	38	23	2.0
444	38	33	2.0
445	38	93	0.0
446	38	444	1.0
447	38	677	3.0
448	38	749	3.0
4070	38	19	0.0
4071	38	25	0.0
4072	38	51	0.0
4073	38	175	1.0
4074	38	212	1.0
4075	38	227	NaN
4076	38	445	1.0
4077	38	595	NaN
4078	38	601	NaN
4079	38	676	0.0
4080	38	1112	1.0
5141	38	1403	1.0

	did	diagnose
18	19	Adjustment disorder□ (poor adjustment to life ...
22	23	Alcohol withdrawal syndrome□ (mild)
23	25	Alcoholism
31	33	Amphetamine abuse
48	51	Anxiety disorder□generalized anxiety disorder□GAD
88	93	Bipolar disorder□manic depressive disorder
165	175	Cocaine abuse
202	212	Depression□excessive sadness
217	227	Drug reaction
414	444	Magic mushroom ingestion□psilocybin
415	445	Major depressive disorder□severe depression
553	595	Post-traumatic stress disorder□PTSD
558	601	Prescription drug abuse
631	676	Schizoaffective disorder□features of schizophr...
632	677	Schizophrenia□chronic impaired reality perception
700	749	Temporal lobe epilepsy□non-convulsive seizure
1004	1112	Hepatic encephalopathy□confusion from liver fa...
1112	1403	Delusional disorder



Inference : Delusions and hallucinations are not the main symptoms for mental illness.

17. Perform analysis on the fact that most common symptoms found in the diseases of highest fatality

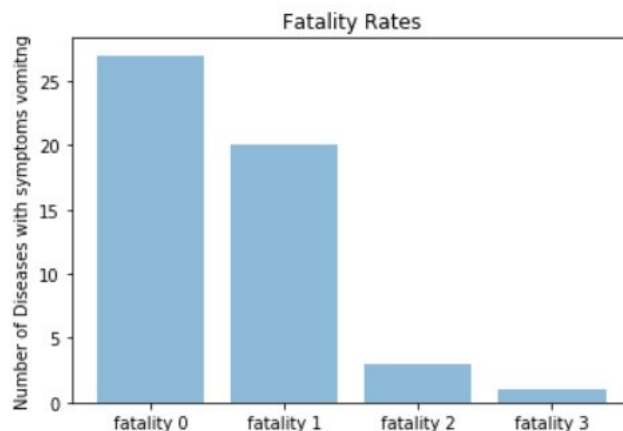
```
import pandas as pd
import matplotlib.pyplot as plt
import numpy as np
sym=pd.read_excel("Symptoms.xlsx")
fatal=pd.read_excel("Disease_fatality.xlsx")
disease1=fatal.groupby('wei')
diseasevomit=disease1.get_group(3)
disease12=diseasevomit.groupby('syd')
#diseasevomit.set_index('syd', inplace=True)
print(diseasevomit)
p=0
min1=[]
max2=[]
for grp,fta in disease12:
    #print(fta)
    if p==0:
        max=fta.shape[0]
        p=1
    r=fta.shape[0]
    if(r>max):
        max=r
        comsym=fta['syd']
print(comsym)
for t,grp1 in disease12:
    min=[grp1.shape[0]]
    min1.extend(min)
    max1=[grp1['syd']]
    max2.extend(max1)
#print(max2)
print(min1)
arr1=np.array(min1)
arr2=np.array(max2)
figureObject, axesObject = plt.subplots()
axesObject.pie(arr1,autopct='%1.2f',startangle=90,axesObject.axis('equal'))
plt.show()
```

```
[394 rows x 3 columns]
4726    292
4730    292
4733    292
4737    292
4742    292
4746    292
4747    292
4752    292
4757    292
4761    292
Name: syd, dtype: int64
```

Inference : The most common symptom found among diseases which are most fatal is “Confusion”

18. Perform analysis on the fact that fatality rate of diseases which have symptoms of vomiting

```
sym=pd.read_excel("Symptoms.xlsx")
fatal=pd.read_excel("Disease_fatality.xlsx")
index1=0
sym.set_index('syd',inplace=True)
print(sym)
for index,sympt in sym.iterrows():
    print(index)
    print(sympt)
    if sympt["symptom"]=="Vomiting":
        index1=index
        break
print(index1)
disease1=fatal.groupby('syd')
diseasevomit=disease1.get_group(index1)
print(diseasevomit)
fatalrate=[]
i=0
vomitfatal=diseasevomit.groupby('wei')
#vomitfatal.set_index('syd',inplace=True)
for fat,grp in vomitfatal:
    print(fat)
    print(grp)
    ftal1=(grp.shape[0])
    fatalrat=[ftal1]
    fatalrate.extend(fatalrat)
print(fatalrate)
fatalrate1=pd.DataFrame(fatalrate,index=['fatality 0','fatality 1','fatality 2','fatality 3'])
print(fatalrate1)
fatal2=np.asarray(fatalrate)
print(fatal2)
objects = ('fatality 0','fatality 1','fatality 2','fatality 3')
y_pos = np.arange(len(objects))
plt.bar(y_pos,fatal2, align='center', alpha=0.5)
plt.xticks(y_pos, objects)
plt.ylabel('Number of Diseases with symptoms vomitng')
plt.title('Fatality Rates')
plt.show()
```



Inference : Vomiting causes the least fatal diseases.

19. Perform analysis on the fact that the symptoms related to different types of organ failures and has a fatality rate equal to 3

```
arr=['failure','Failure']
sym=pd.read_excel("Symptoms.xlsx")
sym['symptom'].fillna('No',inplace=True)
dies=pd.read_excel("Disease.xlsx")
fatal=pd.read_excel("Disease_fatality.xlsx")
tempdf=sym.loc[sym['symptom'].str.contains('|'.join(arr))]
#print(tempdf)
u=3
l2=[]
tempdf.set_index('syd',inplace=True)
print(tempdf)
for i,group in tempdf.iterrows():
    l1=[i]
    l2.extend(l1)
print(l2)
l3=[]
tempdf1=fatal.groupby('syd')
for i in l2:
    for j,group1 in tempdf1:
        if i==j:
            df1=tempdf1.get_group(i)
            df2=df1.groupby('wei')
            df3=df2.get_group(u)
            p=df3.shape[0]
            l4=[p]
            l3.extend(l4)
l6=[]
print(l3)
sym.set_index('syd',inplace=True)
for j in l2:
    for i,gro in sym.iterrows():
        if i==j:
            l5=[gro['symptom']]
            l6.extend(l5)
arr1=np.asarray(l6)
```

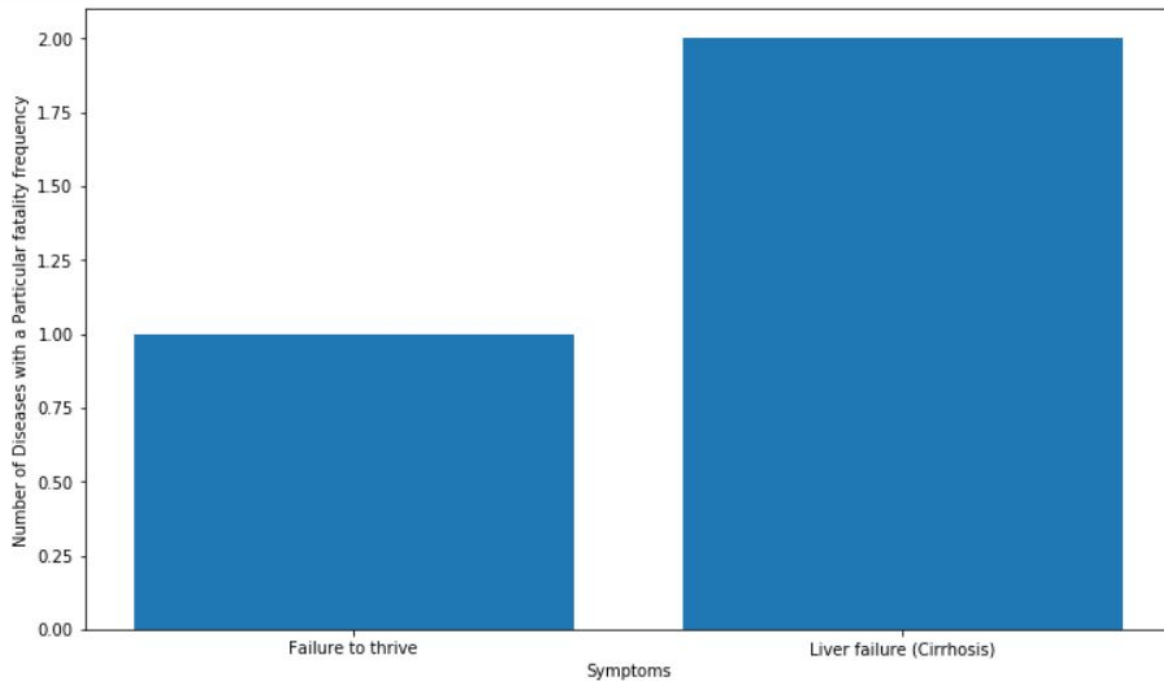


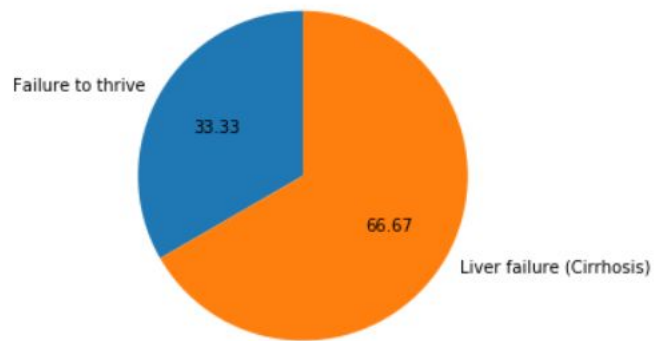
```

print(l3)
sym.set_index('syd',inplace=True)
for j in l2:
    for i,gro in sym.iterrows():
        if i==j:
            l5=[gro['symptom']]
            l6.extend(l5)
arr1=np.asarray(l6)
objects =arr1
y_pos = np.arange(len(objects))
plt.figure(figsize=(12, 7))
plt.bar(y_pos,l3, align='center', alpha=1)
plt.xticks(y_pos, objects)
plt.ylabel('Number of Diseases with a Particular fatality frequency')
plt.xlabel('Symptoms')
plt.title('')
plt.show()
arr2=arr1
figureObject, axesObject = plt.subplots()
axesObject.pie(l3,labels=arr2,autopct='%1.2f',startangle=90,axesObject.axis('equal'))
plt.show()

```

	syd	symptom
132		Failure to thrive
149		Liver failure (Cirrhosis)





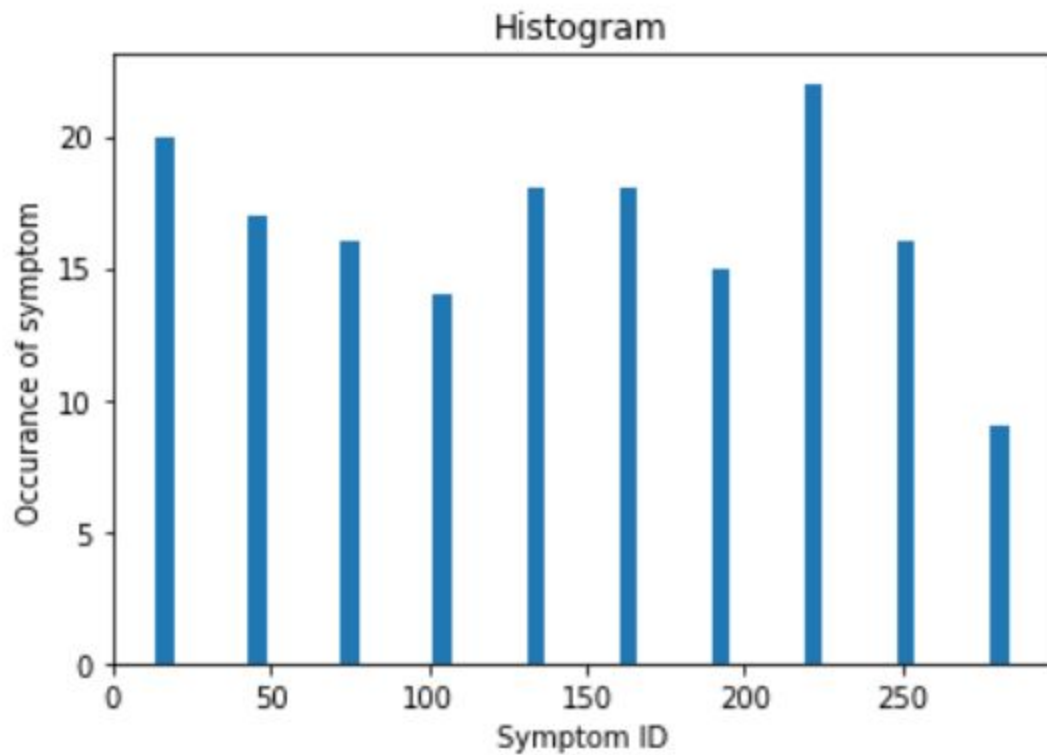
Inference : The most common organ failures are “Liver failure”, “Failure to thrive”. Liver failure being the most common amongst the two.

20. Perform analysis to find the occurrence of symptoms of highest fatality

```
import pandas as pd
import numpy as np
symid1=[]
data=pd.read_csv("datamatch.csv")
dia=pd.read_csv("diagnose.csv")
sym=pd.read_csv("symptom.csv")
for i,group in sym.iterrows():
    symid=group['syd']
    symid1.extend(symid)
#print(symid1)
dia.rename(columns = {"did": "Disease ID", "diagnose":"Disease"}, inplace = True)
sym.rename(columns = {"syd": "Symptom ID", "symptom":"Symptom"}, inplace = True)
data.rename(columns = {"did": "Disease ID", "syd":"Symptom ID","wei":"Fatality"}, inplace = True)
empdf4=data.loc[data['Fatality']==3]
#print(empdf4)
empdf5=sym.loc[sym['Symptom ID'].isin(empdf4['Symptom ID'])]
#print(empdf5)
l4=[]
for i in empdf4['Symptom ID']:
    #print(i)
    l3=[i]
    l4.extend(l3)
#print(l4)

final_list = []
for num in l4:
    if num not in final_list:
        final_list.append(num)

print(final_list)
#Histogram
population_age = final_list
bins = 10
plt.hist(population_age, bins, histtype='bar',rwidth=0.2)
plt.xlabel('Symptom ID')
plt.ylabel('Occurance of symptom')
plt.title('Histogram')
plt.show()
```



Inference : The most common Symptoms found of fatality 3 are “Confusion”, “Change in behaviour”, “Fainting”, “Fever in the returning traveller”, “Ingestion” and “Headache after trauma”.

21. Perform analysis on the fact that diseases with symptoms of upper abdominal pain whose fatality is greater than or equal to 2

```
str1="Upper"
sym.set_index('syd',inplace=True)
index1=0
diarr2=[]
for index,symptoms in sym.iterrows():
    for j in symptoms:
        smp=j.split(" ")
        for o in smp:
            for t in o:
                print(t)
                if t==str1:
                    index1=index
                    break
            break
        break
    break
print(index1)

fatal.set_index('did',inplace=True)
#print(fatal)
for did,fatals in fatal.iterrows():
    #print(did)
    #print(fatals)
    if fatals['syd']==index1:
        if(fatals['wei']==2 or fatals['wei']>2):
            diarr=[did]
            diarr2.extend(diarr)
print(diarr2)

dies.set_index('did',inplace=True)

for dyd,dname in dies.iterrows():
    for i in diarr2:
        if(i==dyd):
            print(dname['diagnose'])
```

Output:

```
[163, 164, 187, 306, 308, 309, 546, 988, 1115]
Cholecystitis□inflammation of the gallbladder
Choledocholithiasis□stone in bile duct
Constipation
Gastric ulcer□stomach ulcer
Gastroenteritis□intestinal infection
Gastroesophageal reflux□GERD, heartburn
Pancreatitis□pancreas inflammation
Lactose intolerance
Ventral hernia□bulging of the abdominal wall
```

Inference: Most of the diseases related to upper abdominal pain are gastric problem

22. Perform analysis to calculate the Death percentage in top 3 Diseases.

```
fatal=pd.read_excel("Disease_fatality.xlsx")
tempdf=dies.loc[dies['diagnose'].str.match('Subdural')]
p=tempdf['did']
print(tempdf)
tempdf1=dies.loc[dies['diagnose'].str.match('Anaphylaxis')]
p1=tempdf1['did']
tempdf3=dies.loc[dies['diagnose'].str.match('Poison ivy')]
p2=tempdf3['did']
size3=[]
df1=fatal.groupby('did')
df11=df1.get_group(736)
df112=df11.groupby('wei')
for i,group in df112:
    if i==3:
        size1=group.shape[0]

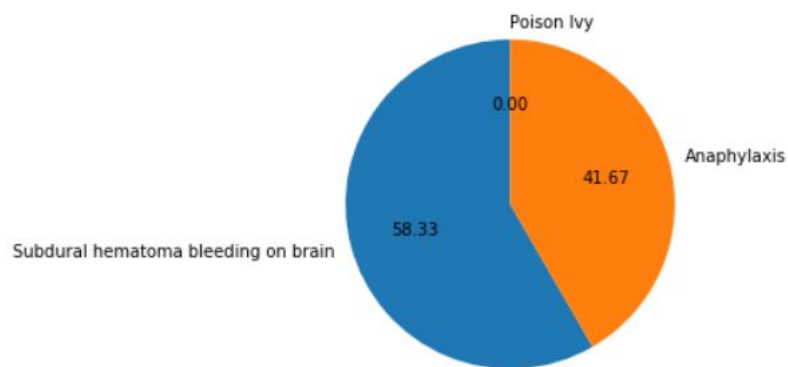
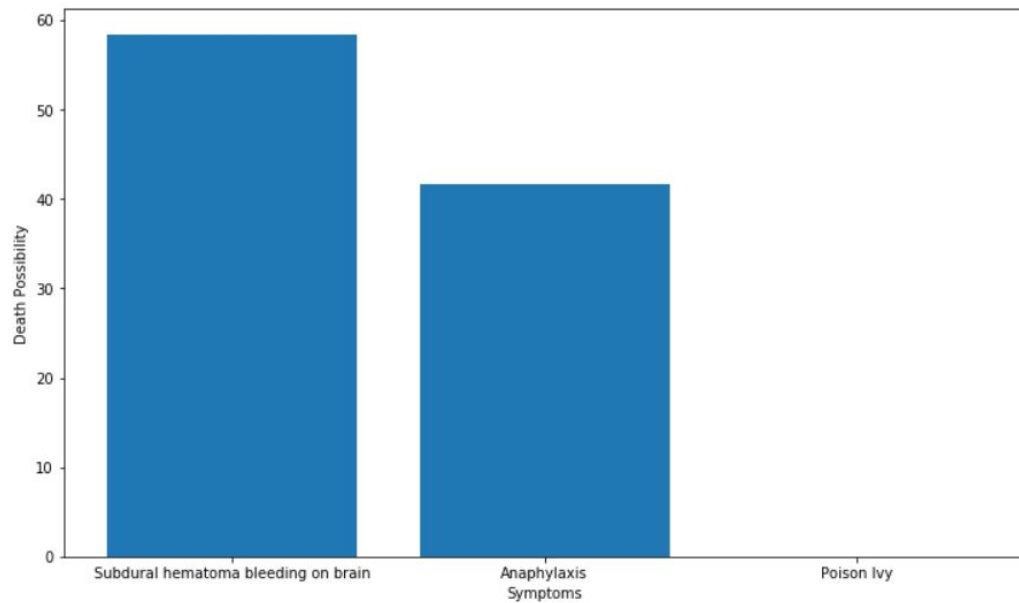
df11=df1.get_group(38)
df113=df11.groupby('wei')
for i,group1 in df113:
    if i==3:
        size2=group1.shape[0]

df11=df1.get_group(189)
df114=df11.groupby('wei')
for i,group2 in df114:
    if i==3:
        size4=group2.shape[0]
fat0=(7/12)*100
print(fat0)
fat1=(5/12)*100
print(fat1)
fat2=(0/12)*100
print(fat2)
l1=[fat0,fat1,fat2]
arr1=np.asarray(l1)
```

```
objects =('Subdural hematoma bleeding on brain','Anaphylaxis','Poison Ivy')
y_pos = np.arange(len(objects))
plt.figure(figsize=(12, 7))
plt.bar(y_pos,arr1, align='center', alpha=1)
plt.xticks(y_pos, objects)
plt.ylabel('Death Possibility')
plt.xlabel('Symptoms')
plt.title('')
plt.show()
arr2=arr1
figureObject, axesObject = plt.subplots()
axesObject.pie(arr1,labels=('Subdural hematoma bleeding on brain','Anaphylaxis','Poison Ivy'),autopct='%1.2f',startangle=90)
axesObject.axis('equal')
plt.show()
```

Death Percentage:

```
58.333333333333336
41.66666666666667
0.0
```



Inference: The Death percentage of Subdural hematoma bleeding on brain is 58.33

The Death percentage of Anaphylaxis is 41.33

The Death percentage of Poison Ivy is 0.0