

Disease Prediction based on Symptoms

```
In [1]: from mpl_toolkits.mplot3d import Axes3D
        from sklearn.preprocessing import StandardScaler
        import matplotlib.pyplot as plt
        from tkinter import *
        import numpy as np
        import pandas as pd
        import os
```

```
In [2]: l1=['back_pain','constipation','abdominal_pain','diarrhoea','mild_fever',
            'yellow_urine',
            'yellowing_of_eyes','acute_liver_failure','fluid_overload','swelling_of_stomach',
            'swollen_lymph_nodes','malaise','blurred_and_distorted_vision','phlegm',
            'throat_irritation',
            'redness_of_eyes','sinus_pressure','runny_nose','congestion','chest_pain',
            'weakness_in_limbs',
            'fast_heart_rate','pain_during_bowel_movements','pain_in_anal_region',
            'bloody_stool',
            'irritation_in_anus','neck_pain','dizziness','cramps','bruising','obesity',
            'swollen_legs',
            'swollen_blood_vessels','puffy_face_and_eyes','enlarged_thyroid','brittle_nails',
            'swollen_extremities','excessive_hunger','extra_marital_contacts',
            'drying_and_tingling_lips',
            'slurred_speech','knee_pain','hip_joint_pain','muscle_weakness','stiff_neck',
            'swelling_joints',
            'movement_stiffness','spinning_movements','loss_of_balance','unsteadiness',
            'weakness_of_one_body_side','loss_of_smell','bladder_discomfort','foul_smell_of_urine',
            'continuous_feel_of_urine','passage_of_gases','internal_itching','toxic_look_(typhos)',
```



```
0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0]
```

```
In [5]: df=pd.read_csv("C:/Users/Damodhar Sai Pesay/Desktop/training.csv")
DF= pd.read_csv('C:/Users/Damodhar Sai Pesay/Desktop/training.csv', ind
ex_col='prognosis')
df.replace({'prognosis':{'Fungal infection':0,'Allergy':1,'GERD':2,'Chr
onic cholestasis':3,'Drug Reaction':4,
'Peptic ulcer diseae':5,'AIDS':6,'Diabetes ':7,'Gastroenteritis':8,
'Bronchial Asthma':9,'Hypertension ':10,
'Migraine':11,'Cervical spondylosis':12,
'Paralysis (brain hemorrhage)':13,'Jaundice':14,'Malaria':15,'Chick
en pox':16,'Dengue':17,'Typhoid':18,'hepatitis A':19,
'Hepatitis B':20,'Hepatitis C':21,'Hepatitis D':22,'Hepatitis E':23
,'Alcoholic hepatitis':24,'Tuberculosis':25,
'Common Cold':26,'Pneumonia':27,'Dimorphic hemmorhoids(piles)':28,
'Heart attack':29,'Varicose veins':30,'Hypothyroidism':31,
'Hyperthyroidism':32,'Hypoglycemia':33,'Osteoarthritis':34,'Arthri
tis':35,
'(vertigo) Paroymsal Positional Vertigo':36,'Acne':37,'Urinary tra
ct infection':38,'Psoriasis':39,
'Impetigo':40}},inplace=True)
#df.head()
DF.head()
```

Out[5]:

	itching	skin_rash	nodal_skin_eruptions	continuous_sneezing	shivering	chills	joint
prognosis							
Fungal infection	1	1	1	0	0	0	
Fungal infection	0	1	1	0	0	0	
Fungal infection	1	0	1	0	0	0	
Fungal infection	1	1	0	0	0	0	

Fungal infection	1	1	1	0	0	0
------------------	---	---	---	---	---	---

5 rows × 132 columns

```
In [6]: def plotPerColumnDistribution(df1, nGraphShown, nGraphPerRow):
    unique = df1.nunique()
    df1 = df1[[col for col in df1 if unique[col] > 1 and unique[col] < 50]]
    nRow, nCol = df1.shape
    columnNames = list(df1)
    nGraphRow = (nCol + nGraphPerRow - 1) // nGraphPerRow
    plt.figure(num = None, figsize = (6 * nGraphPerRow, 8 * nGraphRow),
    dpi = 80, facecolor = 'w', edgecolor = 'k')
    for i in range(min(nCol, nGraphShown)):
        plt.subplot(nGraphRow, nGraphPerRow, i + 1)
        columnDf = df1.iloc[:, i]
        if (not np.issubdtype(type(columnDf.iloc[0]), np.number)):
            valueCounts = columnDf.value_counts()
            valueCounts.plot.bar()
        else:
            columnDf.hist()
            plt.ylabel('counts')
            plt.xticks(rotation = 90)
            plt.title(f'{columnNames[i]} (column {i})')
    plt.tight_layout(pad = 1.0, w_pad = 1.0, h_pad = 1.0)
    plt.show()
```

```
In [7]: def plotScatterMatrix(df1, plotSize, textSize):
    df1 = df1.select_dtypes(include=[np.number])

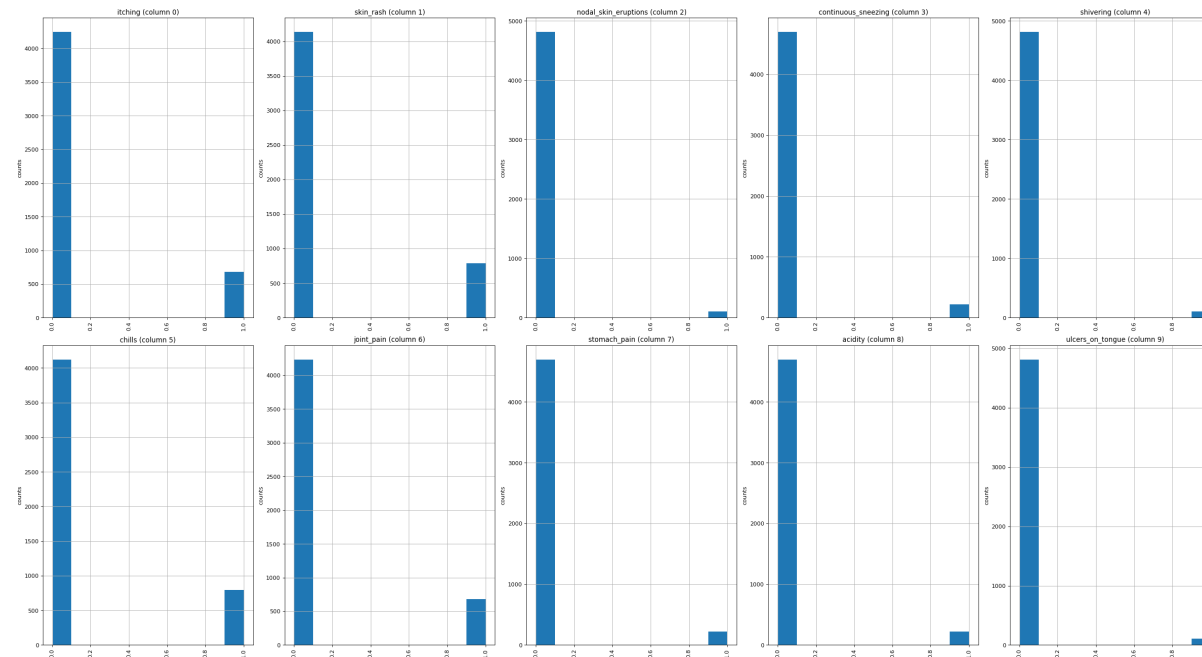
    df1 = df1.dropna('columns')
    df1 = df1[[col for col in df1 if df1[col].nunique() > 1]]
    columnNames = list(df1)
    if len(columnNames) > 10:
        columnNames = columnNames[:10]
    df1 = df1[columnNames]
    ax = pd.plotting.scatter_matrix(df1, alpha=0.75, figsize=[plotSize,
```

```

plotSize], diagonal='kde')
corrs = df1.corr().values
for i, j in zip(*plt.np.triu_indices_from(ax, k = 1)):
    ax[i, j].annotate('Corr. coef = %.3f' % corrs[i, j], (0.8, 0.2
), xycoords='axes fraction', ha='center', va='center', size=textSize)
plt.suptitle('Scatter and Density Plot')
plt.show()

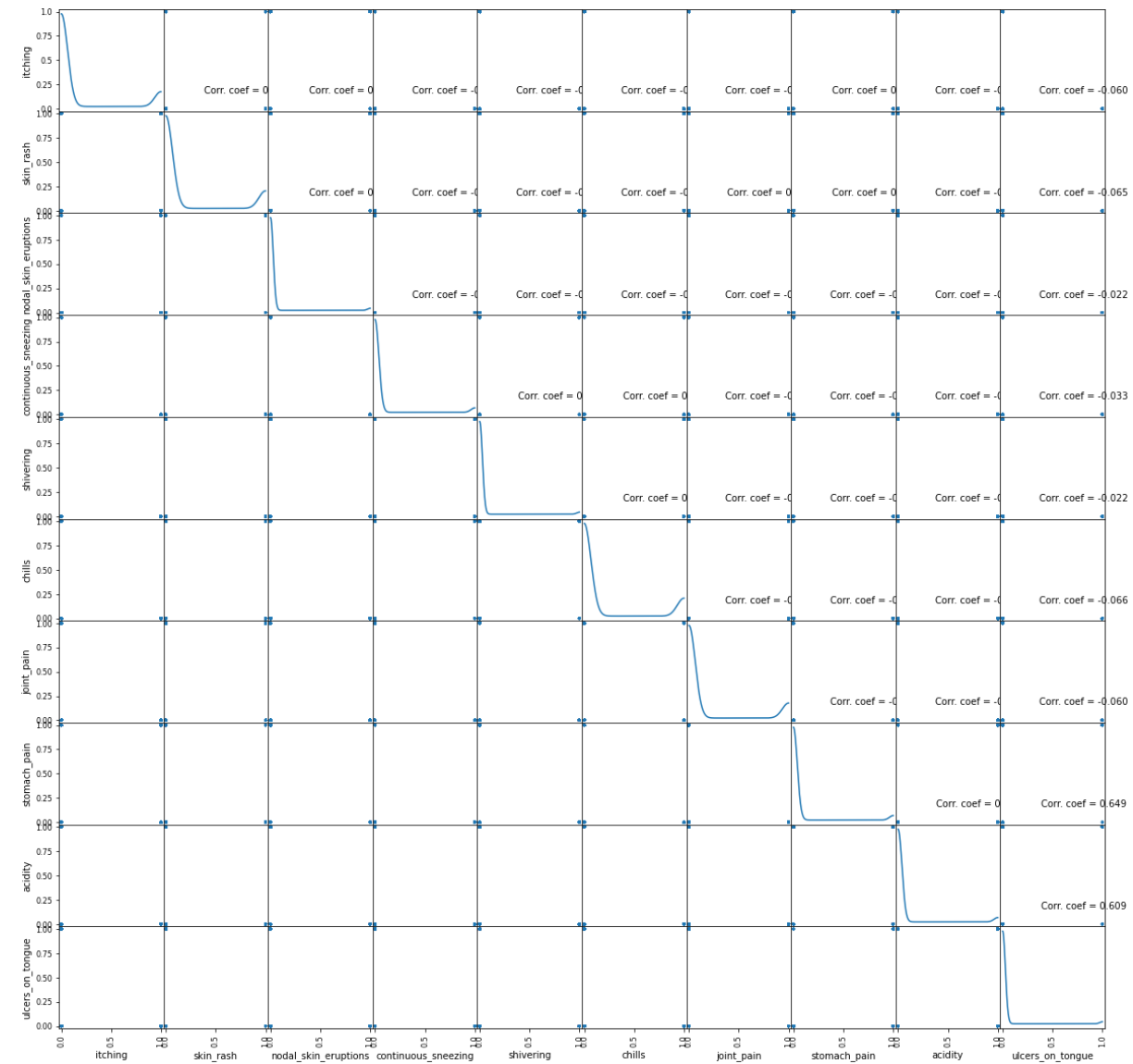
```

In [8]: `plotPerColumnDistribution(df, 10, 5)`



In [9]: `plotScatterMatrix(df, 20, 10)`

Scatter and Density Plot



```
In [10]: X= df[l1]
y = df[["prognosis"]]
np.ravel(y)
print(X)
```

	back_pain	constipation	abdominal_pain	diarrhoea	mild_fever	\
0	0	0	0	0	0	
1	0	0	0	0	0	
2	0	0	0	0	0	
3	0	0	0	0	0	
4	0	0	0	0	0	
...	
4915	0	0	0	0	0	
4916	0	0	0	0	0	
4917	0	0	0	0	0	
4918	0	0	0	0	0	
4919	0	0	0	0	0	

	yellow_urine	yellowing_of_eyes	acute_liver_failure	fluid_overl	oad	\
0	0	0	0	0	0	
1	0	0	0	0	0	
2	0	0	0	0	0	
3	0	0	0	0	0	
4	0	0	0	0	0	
...	
4915	0	0	0	0	0	
4916	0	0	0	0	0	
4917	0	0	0	0	0	
4918	0	0	0	0	0	
...	

4919	0	0	0
0			
	swelling_of_stomach	...	pus_filled_pimples
g \			blackheads
0	0	...	0
0			0
1	0	...	0
0			0
2	0	...	0
0			0
3	0	...	0
0			0
4	0	...	0
0			0
...
...			
4915	0	...	0
0			0
4916	0	...	1
1			1
4917	0	...	0
0			0
4918	0	...	0
0			0
4919	0	...	0
0			0
	skin_peeling	silver_like_dusting	small_dents_in_nails \
0	0	0	0
1	0	0	0
2	0	0	0
3	0	0	0
4	0	0	0
...
4915	0	0	0
4916	0	0	0
4917	0	0	0
4918	1	1	1


```

4919          0          0          0
      inflammatory_nails blister red_sore_around_nose yellow_crust_o
oze
0          0          0          0
  0
1          0          0          0
  0
2          0          0          0
  0
3          0          0          0
  0
4          0          0          0
  0
...          ...          ...          ...
...
4915          0          0          0
  0
4916          0          0          0
  0
4917          0          0          0
  0
4918          1          0          0
  0
4919          0          1          1
  1

[4920 rows x 95 columns]

```

In [11]: `print(y)`

```

      prognosis
0          0
1          0
2          0
3          0
4          0
...          ...
4915        36

```

```
4916      37
4917      38
4918      39
4919      40
```

```
[4920 rows x 1 columns]
```

```
In [12]: tr=pd.read_csv("C:/Users/Damodhar Sai Pesay/Desktop/testing.csv")

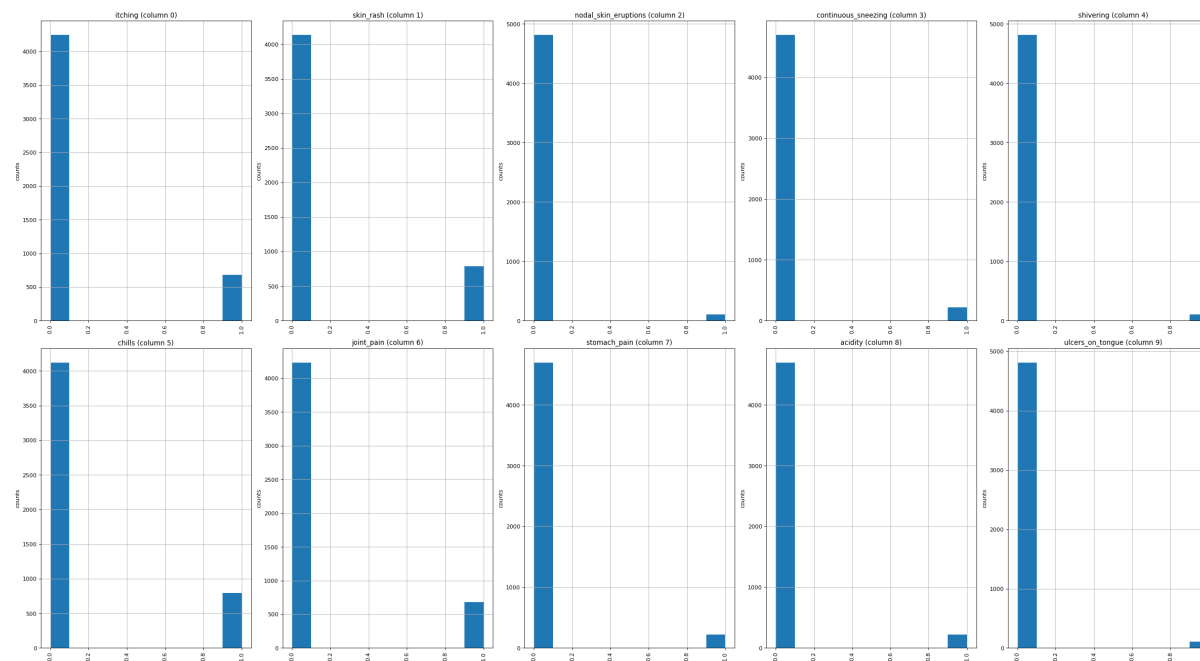
tr.replace({'prognosis':{'Fungal infection':0,'Allergy':1,'GERD':2,'Chronic cholestasis':3,'Drug Reaction':4,
    'Peptic ulcer disease':5,'AIDS':6,'Diabetes ':7,'Gastroenteritis':8,
    'Bronchial Asthma':9,'Hypertension ':10,
    'Migraine':11,'Cervical spondylosis':12,
    'Paralysis (brain hemorrhage)':13,'Jaundice':14,'Malaria':15,'Chicken pox':16,'Dengue':17,'Typhoid':18,'hepatitis A':19,
    'Hepatitis B':20,'Hepatitis C':21,'Hepatitis D':22,'Hepatitis E':23,
    'Alcoholic hepatitis':24,'Tuberculosis':25,
    'Common Cold':26,'Pneumonia':27,'Dimorphic hemmorhoids(piles)':28,
    'Heart attack':29,'Varicose veins':30,'Hypothyroidism':31,
    'Hyperthyroidism':32,'Hypoglycemia':33,'Osteoarthritis':34,'Arthritis':35,
    '(vertigo) Paroymsal Positional Vertigo':36,'Acne':37,'Urinary tract infection':38,'Psoriasis':39,
    'Impetigo':40}}),inplace=True)
tr.head()
```

```
Out[12]:
```

	itching	skin_rash	nodal_skin_eruptions	continuous_sneezing	shivering	chills	joint_pain	st
0	1	1	1	0	0	0	0	
1	0	0	0	1	1	1	0	
2	0	0	0	0	0	0	0	
3	1	0	0	0	0	0	0	
4	1	1	0	0	0	0	0	

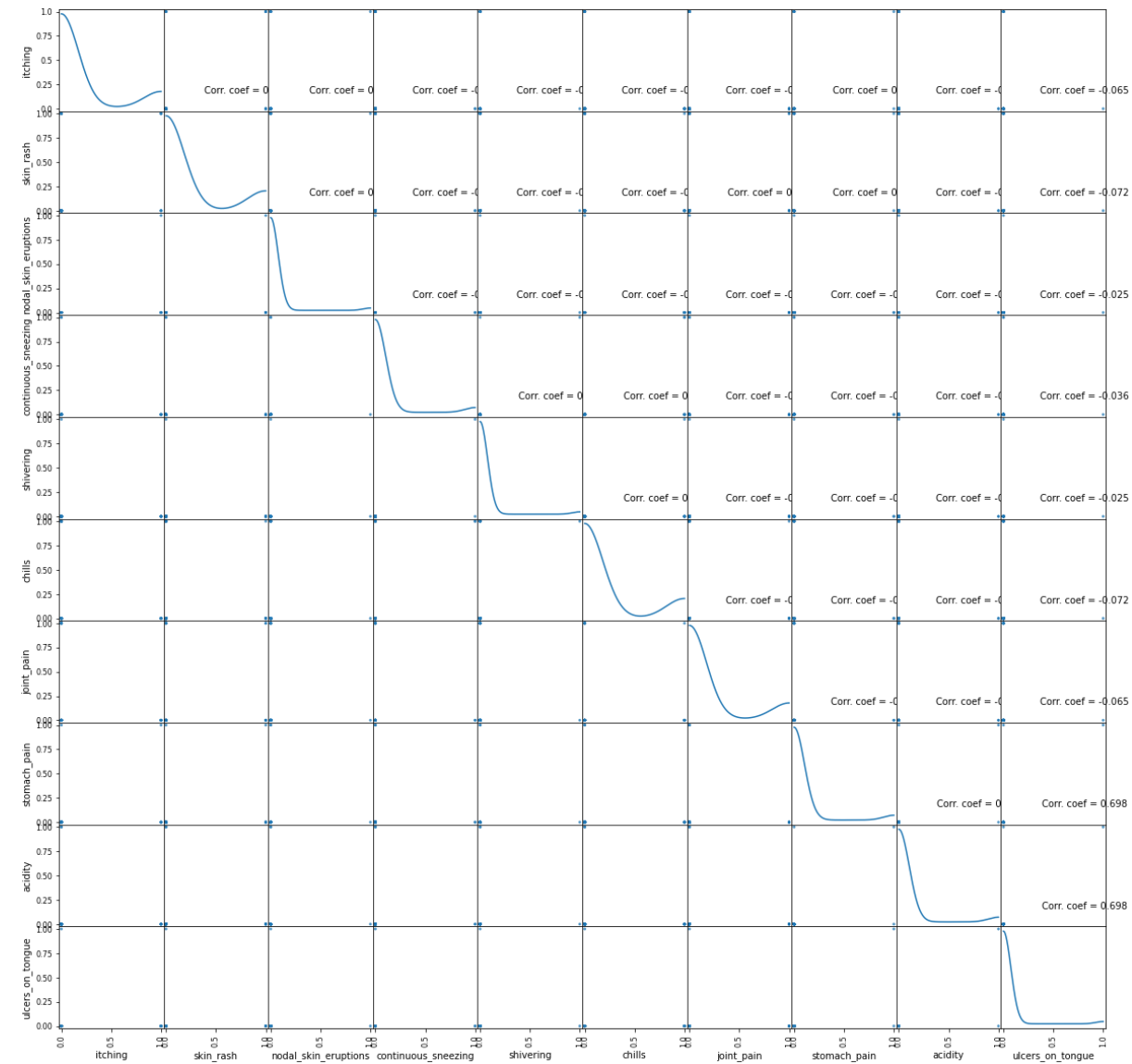
5 rows × 133 columns

```
In [13]: plotPerColumnDistribution(tr, 10, 5)
```



```
In [14]: plotScatterMatrix(tr, 20, 10)
```

Scatter and Density Plot



```
In [15]: X_test= tr[l1]
y_test = tr[["prognosis"]]
np.ravel(y_test)
print(X_test)
```

	back_pain	constipation	abdominal_pain	diarrhoea	mild_fever	\
0	0	0	0	0	0	
1	0	0	0	0	0	
2	0	0	0	0	0	
3	0	0	1	0	0	
4	0	0	0	0	0	
5	0	0	1	0	0	
6	0	0	0	0	0	
7	0	0	0	0	0	
8	0	0	0	1	0	
9	0	0	0	0	0	
10	0	0	0	0	0	
11	0	0	0	0	0	
12	1	0	0	0	0	
13	0	0	0	0	0	
14	0	0	1	0	0	
15	0	0	0	1	0	
16	0	0	0	0	1	
17	1	0	0	0	0	
18	0	1	1	1	0	
19	0	0	1	1	1	
20	0	0	1	0	0	
21	0	0	0	0	0	
22	0	0	1	0	0	
23	0	0	1	0	0	
24	0	0	1	0	0	
25	0	0	0	0	1	
26	0	0	0	0	0	
27	0	0	0	0	0	
28	0	1	0	0	0	
29	0	0	0	0	0	
30	0	0	0	0	0	
31	0	0	0	0	0	
32	0	0	0	1	0	
33	0	0	0	0	0	

34	0	0	0	0	0
35	0	0	0	0	0
36	0	0	0	0	0
37	0	0	0	0	0
38	0	0	0	0	0
39	0	0	0	0	0
40	0	0	0	0	0

	yellow_urine	yellowing_of_eyes	acute_liver_failure	fluid_overloa
d \				
0	0	0	0	
0				
1	0	0	0	
0				
2	0	0	0	
0				
3	0	1	0	
0				
4	0	0	0	
0				
5	0	0	0	
0				
6	0	0	0	
0				
7	0	0	0	
0				
8	0	0	0	
0				
9	0	0	0	
0				
10	0	0	0	
0				
11	0	0	0	
0				
12	0	0	0	
0				
13	0	0	0	
0				
14	0	0	0	

0			
15	0	0	0
0			
16	0	0	0
0			
17	0	0	0
0			
18	0	0	0
0			
19	0	1	0
0			
20	1	1	0
0			
21	0	1	0
0			
22	0	1	0
0			
23	0	1	1
0			
24	0	0	0
0			
25	0	1	0
0			
26	0	0	0
0			
27	0	0	0
0			
28	0	0	0
0			
29	0	0	0
0			
30	0	0	0
0			
31	0	0	0
0			
32	0	0	0
0			
33	0	0	0
0			

34	0	0	0
0			
35	0	0	0
0			
36	0	0	0
0			
37	0	0	0
0			
38	0	0	0
0			
39	0	0	0
0			
40	0	0	0
0			

	swelling_of_stomach	...	pus_filled_pimples	blackheads	scurring
\					
0	0	...	0	0	0
1	0	...	0	0	0
2	0	...	0	0	0
3	0	...	0	0	0
4	0	...	0	0	0
5	0	...	0	0	0
6	0	...	0	0	0
7	0	...	0	0	0
8	0	...	0	0	0
9	0	...	0	0	0
10	0	...	0	0	0

11	0 ...	0	0	0
12	0 ...	0	0	0
13	0 ...	0	0	0
14	0 ...	0	0	0
15	0 ...	0	0	0
16	0 ...	0	0	0
17	0 ...	0	0	0
18	0 ...	0	0	0
19	0 ...	0	0	0
20	0 ...	0	0	0
21	0 ...	0	0	0
22	0 ...	0	0	0
23	0 ...	0	0	0
24	1 ...	0	0	0
25	0 ...	0	0	0
26	0 ...	0	0	0
27	0 ...	0	0	0
28	0 ...	0	0	0
29	0 ...	0	0	0
30	0 ...	0	0	0

31	0	...	0	0	0
32	0	...	0	0	0
33	0	...	0	0	0
34	0	...	0	0	0
35	0	...	0	0	0
36	0	...	0	0	0
37	0	...	1	1	1
38	0	...	0	0	0
39	0	...	0	0	0
40	0	...	0	0	0

	skin_peeling	silver_like_dusting	small_dents_in_nails	\
0	0	0	0	
1	0	0	0	
2	0	0	0	
3	0	0	0	
4	0	0	0	
5	0	0	0	
6	0	0	0	
7	0	0	0	
8	0	0	0	
9	0	0	0	
10	0	0	0	
11	0	0	0	
12	0	0	0	
13	0	0	0	
14	0	0	0	
15	0	0	0	

16	0	0	0
17	0	0	0
18	0	0	0
19	0	0	0
20	0	0	0
21	0	0	0
22	0	0	0
23	0	0	0
24	0	0	0
25	0	0	0
26	0	0	0
27	0	0	0
28	0	0	0
29	0	0	0
30	0	0	0
31	0	0	0
32	0	0	0
33	0	0	0
34	0	0	0
35	0	0	0
36	0	0	0
37	0	0	0
38	0	0	0
39	1	1	1
40	0	0	0

	inflammatory_nails	blister	red_sore_around_nose	yellow_crust_ooz
e				
0	0	0	0	
0				
1	0	0	0	
0				
2	0	0	0	
0				
3	0	0	0	
0				
4	0	0	0	
0				
5	0	0	0	

0			
6	0	0	0
0			
7	0	0	0
0			
8	0	0	0
0			
9	0	0	0
0			
10	0	0	0
0			
11	0	0	0
0			
12	0	0	0
0			
13	0	0	0
0			
14	0	0	0
0			
15	0	0	0
0			
16	0	0	0
0			
17	0	0	0
0			
18	0	0	0
0			
19	0	0	0
0			
20	0	0	0
0			
21	0	0	0
0			
22	0	0	0
0			
23	0	0	0
0			
24	0	0	0
0			

25	0	0	0
0			
26	0	0	0
0			
27	0	0	0
0			
28	0	0	0
0			
29	0	0	0
0			
30	0	0	0
0			
31	0	0	0
0			
32	0	0	0
0			
33	0	0	0
0			
34	0	0	0
0			
35	0	0	0
0			
36	0	0	0
0			
37	0	0	0
0			
38	0	0	0
0			
39	1	0	0
0			
40	0	1	1
1			

[41 rows x 95 columns]

In [16]: `print(y_test)`

```

prognosis
0          0

```

1	1
2	2
3	3
4	4
5	5
6	6
7	7
8	8
9	9
10	10
11	11
12	12
13	13
14	14
15	15
16	16
17	17
18	18
19	19
20	20
21	21
22	22
23	23
24	24
25	25
26	26
27	27
28	28
29	29
30	30
31	31
32	32
33	33
34	34
35	35
36	36
37	37
38	38

```
39         39
40         40
```

To build the precision of the model we have used four algorithms.

1) Decision Tree algorithm 2) Random Forest algorithm 3) KNearestNeighbour algorithm 4) Naive Bayes algorithm

```
In [17]: def scatterplt(disea):
          x = ((DF.loc[disea]).sum())
          x.drop(x[x==0].index,inplace=True)
          print(x.values)
          y = x.keys()
          print(len(x))
          print(len(y))
          plt.title(disea)
          plt.scatter(y,x.values)
          plt.show()

          def scatterinp(sym1,sym2,sym3,sym4,sym5):
              x = [sym1,sym2,sym3,sym4,sym5]
              y = [0,0,0,0,0]
              if(sym1!='Select Here'):
                  y[0]=1
              if(sym2!='Select Here'):
                  y[1]=1
              if(sym3!='Select Here'):
                  y[2]=1
              if(sym4!='Select Here'):
                  y[3]=1
              if(sym5!='Select Here'):
                  y[4]=1
              print(x)
              print(y)
              plt.scatter(x,y)
              plt.show()
```

Decision Tree Algorithm

```
In [18]: root = Tk()
pred1=StringVar()
def DecisionTree():
    if len(NameEn.get()) == 0:
        pred1.set(" ")
        comp=messagebox.askokcancel("System","Kindly fill the Name")
        if comp:
            root.mainloop()
    elif((Symptom1.get()=="Select Here") or (Symptom2.get()=="Select Here")):
        pred1.set(" ")
        sym=messagebox.askokcancel("System","Kindly fill atleast first two Symptoms")
        if sym:
            root.mainloop()
    else:
        from sklearn import tree

        clf3 = tree.DecisionTreeClassifier()
        clf3 = clf3.fit(X,y)

        from sklearn.metrics import classification_report,confusion_matrix,accuracy_score
        y_pred=clf3.predict(X_test)
        print("Decision Tree")
        print("Accuracy")
        print(accuracy_score(y_test, y_pred))
        print(accuracy_score(y_test, y_pred,normalize=False))
        print("Confusion matrix")
        conf_matrix=confusion_matrix(y_test,y_pred)
        print(conf_matrix)

        psymptoms = [Symptom1.get(),Symptom2.get(),Symptom3.get(),Symptom4.get(),Symptom5.get()]

        for k in range(0,len(l1)):
```



```

        for z in psymptoms:
            if(z==l1[k]):
                l2[k]=1

inputtest = [l2]
predict = clf3.predict(inputtest)
predicted=predict[0]

h='no'
for a in range(0,len(disease)):
    if(predicted == a):
        h='yes'
        break

if (h=='yes'):
    pred1.set(" ")
    pred1.set(disease[a])
else:
    pred1.set(" ")
    pred1.set("Not Found")

import sqlite3
conn = sqlite3.connect('database.db')
c = conn.cursor()
c.execute("CREATE TABLE IF NOT EXISTS DecisionTree(Name StringVar,Symtom1 StringVar,Symtom2 StringVar,Symtom3 StringVar,Symtom4 TEXT,Symtom5 TEXT,Disease StringVar)")
c.execute("INSERT INTO DecisionTree(Name,Symtom1,Symtom2,Symtom3,Symtom4,Symtom5,Disease) VALUES(?,?,?,?,?,?,?)", (NameEn.get(),Symptom1.get(),Symptom2.get(),Symptom3.get(),Symptom4.get(),Symptom5.get(),pred1.get()))
conn.commit()
c.close()
conn.close()

scatterinp(Symptom1.get(),Symptom2.get(),Symptom3.get(),Symptom

```

```
4.get(),Symptom5.get())
scatterplt(pred1.get())
```

Random Forest Algorithm

```
In [19]: pred2=StringVar()
def randomforest():
    if len(NameEn.get()) == 0:
        pred1.set(" ")
        comp=messagebox.askokcancel("System","Kindly Fill the Name")
        if comp:
            root.mainloop()
    elif((Symptom1.get()=="Select Here") or (Symptom2.get()=="Select Here")):
        pred1.set(" ")
        sym=messagebox.askokcancel("System","Kindly Fill atleast first two Symptoms")
        if sym:
            root.mainloop()
    else:
        from sklearn.ensemble import RandomForestClassifier
        clf4 = RandomForestClassifier(n_estimators=100)
        clf4 = clf4.fit(X,np.ravel(y))

        # calculating accuracy
        from sklearn.metrics import classification_report,confusion_matrix,accuracy_score
        y_pred=clf4.predict(X_test)
        print("Random Forest")
        print("Accuracy")
        print(accuracy_score(y_test, y_pred))
        print(accuracy_score(y_test, y_pred,normalize=False))
        print("Confusion matrix")
        conf_matrix=confusion_matrix(y_test,y_pred)
        print(conf_matrix)

        psymptoms = [Symptom1.get(),Symptom2.get(),Symptom3.get(),Symptom4.get(),Symptom5.get()]
```

```

om4.get(),Symptom5.get()])

    for k in range(0,len(l1)):
        for z in psymptoms:
            if(z==l1[k]):
                l2[k]=1

    inputtest = [l2]
    predict = clf4.predict(inputtest)
    predicted=predict[0]

    h='no'
    for a in range(0,len(disease)):
        if(predicted == a):
            h='yes'
            break
    if (h=='yes'):
        pred2.set(" ")
        pred2.set(disease[a])
    else:
        pred2.set(" ")
        pred2.set("Not Found")

    import sqlite3
    conn = sqlite3.connect('database.db')
    c = conn.cursor()
    c.execute("CREATE TABLE IF NOT EXISTS RandomForest(Name StringVar,Symtom1 StringVar,Symtom2 StringVar,Symtom3 StringVar,Symtom4 TEXT,Symtom5 TEXT,Disease StringVar)")
    c.execute("INSERT INTO RandomForest(Name,Symtom1,Symtom2,Symtom3,Symtom4,Symtom5,Disease) VALUES(?,?,?,?,?,?,?)", (NameEn.get(),Symptom1.get(),Symptom2.get(),Symptom3.get(),Symptom4.get(),Symptom5.get(),pred2.get()))
    conn.commit()
    c.close()
    conn.close()

    scatterplt(pred2.get())

```

KNearestNeighbour Algorithm

```
In [20]: pred4=StringVar()
def KNN():
    if len(NameEn.get()) == 0:
        pred1.set(" ")
        comp=messagebox.askokcancel("System","Kindly Fill the Name")
        if comp:
            root.mainloop()
    elif((Symptom1.get()=="Select Here") or (Symptom2.get()=="Select Here")):
        pred1.set(" ")
        sym=messagebox.askokcancel("System","Kindly Fill atleast first two Symptoms")
        if sym:
            root.mainloop()
    else:
        from sklearn.neighbors import KNeighborsClassifier
        knn=KNeighborsClassifier(n_neighbors=5,metric='minkowski',p=2)
        knn=knn.fit(X,np.ravel(y))

        from sklearn.metrics import classification_report,confusion_matrix,accuracy_score
        y_pred=knn.predict(X_test)
        print("KNearest Neighbour")
        print("Accuracy")
        print(accuracy_score(y_test, y_pred))
        print(accuracy_score(y_test, y_pred,normalize=False))
        print("Confusion matrix")
        conf_matrix=confusion_matrix(y_test,y_pred)
        print(conf_matrix)

        psymptoms = [Symptom1.get(),Symptom2.get(),Symptom3.get(),Symptom4.get(),Symptom5.get()]

        for k in range(0,len(l1)):
            for z in psymptoms:
                if(z==l1[k]):
```

```

l2[k]=1

inputtest = [l2]
predict = knn.predict(inputtest)
predicted=predict[0]

h='no'
for a in range(0,len(disease)):
    if(predicted == a):
        h='yes'
        break

if (h=='yes'):
    pred4.set(" ")
    pred4.set(disease[a])
else:
    pred4.set(" ")
    pred4.set("Not Found")

import sqlite3
conn = sqlite3.connect('database.db')
c = conn.cursor()
c.execute("CREATE TABLE IF NOT EXISTS KNearestNeighbour(Name StringVar,Symptom1 StringVar,Symptom2 StringVar,Symptom3 StringVar,Symptom4 TEXT,Symptom5 TEXT,Disease StringVar)")
c.execute("INSERT INTO KNearestNeighbour(Name,Symptom1,Symptom2,Symptom3,Symptom4,Symptom5,Disease) VALUES(?,?,?,?,?,?,?)", (NameEn.get(),Symptom1.get(),Symptom2.get(),Symptom3.get(),Symptom4.get(),Symptom5.get(),pred4.get()))
conn.commit()
c.close()
conn.close()

scatterplt(pred4.get())

```

Naive Bayes Algorithm

```

In [21]: pred3=StringVar()
def NaiveBayes():
    if len(NameEn.get()) == 0:
        pred1.set(" ")
        comp=messagebox.askokcancel("System","Kindly Fill the Name")
        if comp:
            root.mainloop()
    elif((Symptom1.get()=="Select Here") or (Symptom2.get()=="Select Here")):
        pred1.set(" ")
        sym=messagebox.askokcancel("System","Kindly Fill atleast first two Symptoms")
        if sym:
            root.mainloop()
    else:
        from sklearn.naive_bayes import GaussianNB
        gnb = GaussianNB()
        gnb=gnb.fit(X,np.ravel(y))

        from sklearn.metrics import classification_report,confusion_matrix,accuracy_score
        y_pred=gnb.predict(X_test)
        print("Naive Bayes")
        print("Accuracy")
        print(accuracy_score(y_test, y_pred))
        print(accuracy_score(y_test, y_pred,normalize=False))
        print("Confusion matrix")
        conf_matrix=confusion_matrix(y_test,y_pred)
        print(conf_matrix)

        psymptoms = [Symptom1.get(),Symptom2.get(),Symptom3.get(),Symptom4.get(),Symptom5.get()]
        for k in range(0,len(l1)):
            for z in psymptoms:
                if(z==l1[k]):
                    l2[k]=1

        inputtest = [l2]

```

```

predict = gnb.predict(inputtest)
predicted=predict[0]

h='no'
for a in range(0,len(disease)):
    if(predicted == a):
        h='yes'
        break
if (h=='yes'):
    pred3.set(" ")
    pred3.set(disease[a])
else:
    pred3.set(" ")
    pred3.set("Not Found")

import sqlite3
conn = sqlite3.connect('database.db')
c = conn.cursor()
c.execute("CREATE TABLE IF NOT EXISTS NaiveBayes(Name StringVar, Symtom1 StringVar, Symtom2 StringVar, Symtom3 StringVar, Symtom4 TEXT, Symtom5 TEXT, Disease StringVar)")
c.execute("INSERT INTO NaiveBayes(Name, Symtom1, Symtom2, Symtom3, Symtom4, Symtom5, Disease) VALUES(?,?,?,?,?,?,?)", (NameEn.get(), Symptom1.get(), Symptom2.get(), Symptom3.get(), Symptom4.get(), Symptom5.get(), pred3.get()))
conn.commit()
c.close()
conn.close()

scatterplt(pred3.get())

```

Building Graphical User Interface

```

In [22]: #Tk class is used to create a root window
root.configure(background='Ivory')
root.title('Smart Disease Predictor System')
root.resizable(0,0)

```

Out[22]: ''

```
In [23]: Symptom1 = StringVar()
Symptom1.set("Select Here")

Symptom2 = StringVar()
Symptom2.set("Select Here")

Symptom3 = StringVar()
Symptom3.set("Select Here")

Symptom4 = StringVar()
Symptom4.set("Select Here")

Symptom5 = StringVar()
Symptom5.set("Select Here")
Name = StringVar()
```

```
In [24]: prev_win=None
def Reset():
    global prev_win

    Symptom1.set("Select Here")
    Symptom2.set("Select Here")
    Symptom3.set("Select Here")
    Symptom4.set("Select Here")
    Symptom5.set("Select Here")
    NameEn.delete(first=0,last=100)
    pred1.set(" ")
    pred2.set(" ")
    pred3.set(" ")
    pred4.set(" ")
    try:
        prev_win.destroy()
        prev_win=None
    except AttributeError:
        pass
```



```
In [25]: from tkinter import messagebox
def Exit():
    qExit=messagebox.askyesno("System","Do you want to exit the system"
    )

    if qExit:
        root.destroy()
        exit()
```

```
In [26]: #Headings for the GUI written at the top of GUI
w2 = Label(root, justify=LEFT, text="Disease Predictor using Machine Le
arning", fg="Red", bg="Ivory")
w2.config(font=("Times",30,"bold italic"))
w2.grid(row=1, column=0, columnspan=2, padx=100)
w2 = Label(root, justify=LEFT, text="Contributors: Damodhar,Surya,Phan
i,Karthik", fg="Pink", bg="Ivory")
w2.config(font=("Times",30,"bold italic"))
w2.grid(row=2, column=0, columnspan=2, padx=100)
```

```
In [27]: #Label for the name
NameLb = Label(root, text="Name of the Patient *", fg="Red", bg="Ivory"
)
NameLb.config(font=("Times",15,"bold italic"))
NameLb.grid(row=6, column=0, pady=15, sticky=W)
```

```
In [28]: #Creating Labels for the symptoms
S1Lb = Label(root, text="Symptom 1 *", fg="Black", bg="Ivory")
S1Lb.config(font=("Times",15,"bold italic"))
S1Lb.grid(row=7, column=0, pady=10, sticky=W)

S2Lb = Label(root, text="Symptom 2 *", fg="Black", bg="Ivory")
S2Lb.config(font=("Times",15,"bold italic"))
S2Lb.grid(row=8, column=0, pady=10, sticky=W)

S3Lb = Label(root, text="Symptom 3", fg="Black",bg="Ivory")
S3Lb.config(font=("Times",15,"bold italic"))
S3Lb.grid(row=9, column=0, pady=10, sticky=W)
```

```

S4Lb = Label(root, text="Symptom 4", fg="Black", bg="Ivory")
S4Lb.config(font=("Times",15,"bold italic"))
S4Lb.grid(row=10, column=0, pady=10, sticky=W)

S5Lb = Label(root, text="Symptom 5", fg="Black", bg="Ivory")
S5Lb.config(font=("Times",15,"bold italic"))
S5Lb.grid(row=11, column=0, pady=10, sticky=W)

```

```

In [29]: #Labels for the different algorithms
lrLb = Label(root, text="DecisionTree", fg="white", bg="red", width = 20)
lrLb.config(font=("Times",15,"bold italic"))
lrLb.grid(row=15, column=0, pady=10,sticky=W)

destreeLb = Label(root, text="RandomForest", fg="Red", bg="Orange", width = 20)
destreeLb.config(font=("Times",15,"bold italic"))
destreeLb.grid(row=17, column=0, pady=10, sticky=W)

ranfLb = Label(root, text="NaiveBayes", fg="White", bg="green", width = 20)
ranfLb.config(font=("Times",15,"bold italic"))
ranfLb.grid(row=19, column=0, pady=10, sticky=W)

knnLb = Label(root, text="kNearestNeighbour", fg="Red", bg="Sky Blue", width = 20)
knnLb.config(font=("Times",15,"bold italic"))
knnLb.grid(row=21, column=0, pady=10, sticky=W)
OPTIONS = sorted(l1)

```

```

In [30]: #Taking name as input from user
NameEn = Entry(root, textvariable=Name)
NameEn.grid(row=6, column=1)

#Taking Symptoms as input from the dropdown from the user
S1 = OptionMenu(root, Symptom1,*OPTIONS)
S1.grid(row=7, column=1)

```

```

S2 = OptionMenu(root, Symptom2,*OPTIONS)
S2.grid(row=8, column=1)

S3 = OptionMenu(root, Symptom3,*OPTIONS)
S3.grid(row=9, column=1)

S4 = OptionMenu(root, Symptom4,*OPTIONS)
S4.grid(row=10, column=1)

S5 = OptionMenu(root, Symptom5,*OPTIONS)
S5.grid(row=11, column=1)

```

```

In [31]: #Buttons for predicting the disease using different algorithms
dst = Button(root, text="Prediction 1", command=DecisionTree,bg="Red",f
g="yellow")
dst.config(font=("Times",15,"bold italic"))
dst.grid(row=6, column=3,padx=10)

rnf = Button(root, text="Prediction 2", command=randomforest,bg="Light
green",fg="red")
rnf.config(font=("Times",15,"bold italic"))
rnf.grid(row=7, column=3,padx=10)

lr = Button(root, text="Prediction 3", command=NaiveBayes,bg="Blue",fg=
"white")
lr.config(font=("Times",15,"bold italic"))
lr.grid(row=8, column=3,padx=10)

kn = Button(root, text="Prediction 4", command=KNN,bg="sky blue",fg="re
d")
kn.config(font=("Times",15,"bold italic"))
kn.grid(row=9, column=3,padx=10)

rs = Button(root,text="Reset Inputs", command=Reset,bg="yellow",fg="pur
ple",width=15)
rs.config(font=("Times",15,"bold italic"))
rs.grid(row=10,column=3,padx=10)

ex = Button(root,text="Exit System", command=Exit,bg="yellow",fg="purpl

```

```
e",width=15)
ex.config(font=("Times",15,"bold italic"))
ex.grid(row=11,column=3,padx=10)
```

```
In [32]: #Showing the output of different algorithms
t1=Label(root,font=("Times",15,"bold italic"),text="Decision Tree",height=1,bg="Light green",width=40,fg="red",textvariable=pred1,relief="sunken").grid(row=15, column=1, padx=10)

t2=Label(root,font=("Times",15,"bold italic"),text="Random Forest",height=1,bg="Purple",width=40,fg="white",textvariable=pred2,relief="sunken").grid(row=17, column=1, padx=10)

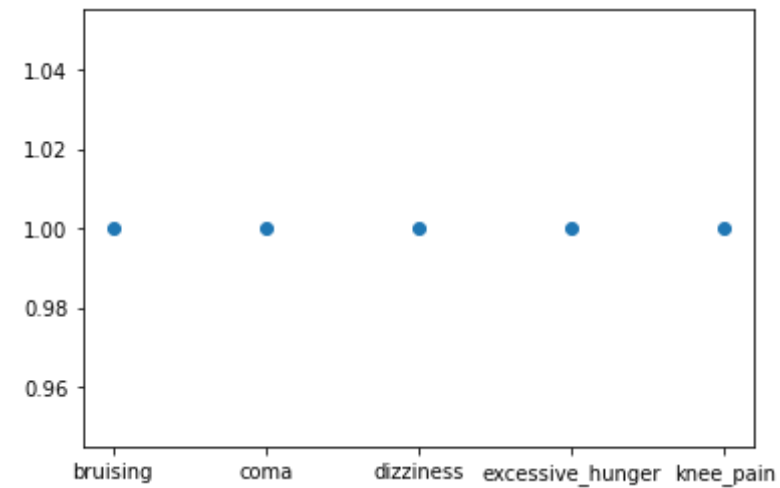
t3=Label(root,font=("Times",15,"bold italic"),text="Naive Bayes",height=1,bg="red",width=40,fg="orange",textvariable=pred3,relief="sunken").grid(row=19, column=1, padx=10)

t4=Label(root,font=("Times",15,"bold italic"),text="kNearest Neighbour",height=1,bg="Blue",width=40,fg="yellow",textvariable=pred4,relief="sunken").grid(row=21, column=1, padx=10)
```

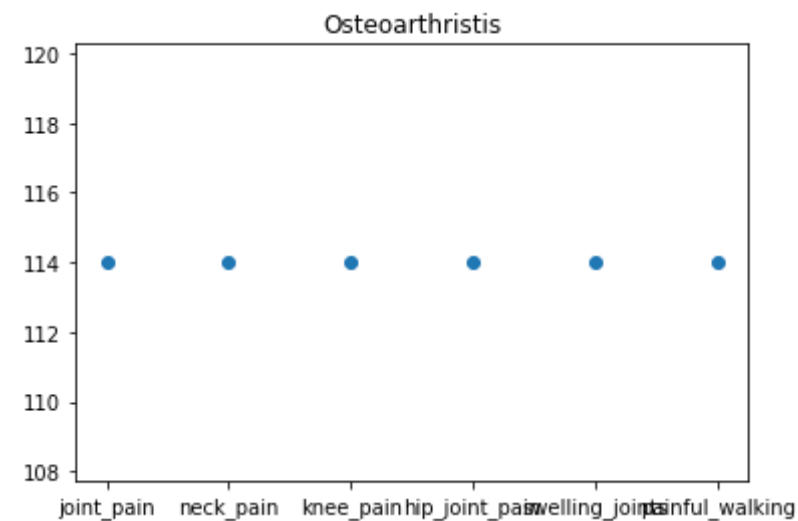
```
In [ ]: #calling this function because the application is ready to run
root.mainloop()
```

```
Decision Tree
Accuracy
0.9512195121951219
39
Confusion matrix
[[1 0 0 ... 0 0 0]
 [0 1 0 ... 0 0 0]
 [0 0 1 ... 0 0 0]
 ...
 [0 0 0 ... 1 0 0]
 [0 0 0 ... 0 1 0]
```

```
[0 0 0 ... 0 0 1]]
['bruising', 'coma', 'dizziness', 'excessive_hunger', 'knee_pain']
[1, 1, 1, 1, 1]
```



```
[114 114 114 114 114 114]
6
6
```

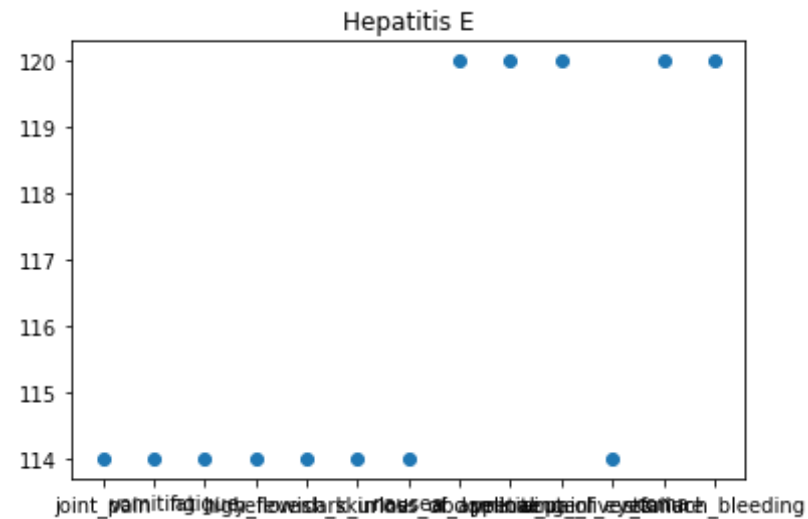


Random Forest

```

Accuracy
0.9512195121951219
39
Confusion matrix
[[1 0 0 ... 0 0 0]
 [0 1 0 ... 0 0 0]
 [0 0 1 ... 0 0 0]
 ...
 [0 0 0 ... 1 0 0]
 [0 0 0 ... 0 1 0]
 [0 0 0 ... 0 0 1]]
[114 114 114 114 114 114 114 120 120 120 114 120 120]
13
13

```



```

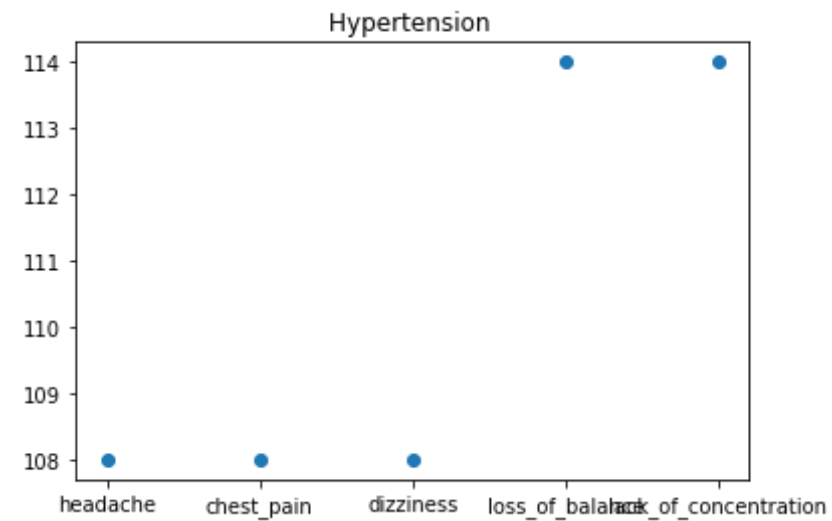
Naive Bayes
Accuracy
0.9512195121951219
39
Confusion matrix
[[1 0 0 ... 0 0 0]
 [0 1 0 ... 0 0 0]
 [0 0 1 ... 0 0 0]

```

```

...
[0 0 0 ... 1 0 0]
[0 0 0 ... 0 1 0]
[0 0 0 ... 0 0 1]]
[108 108 108 114 114]
5
5

```



```

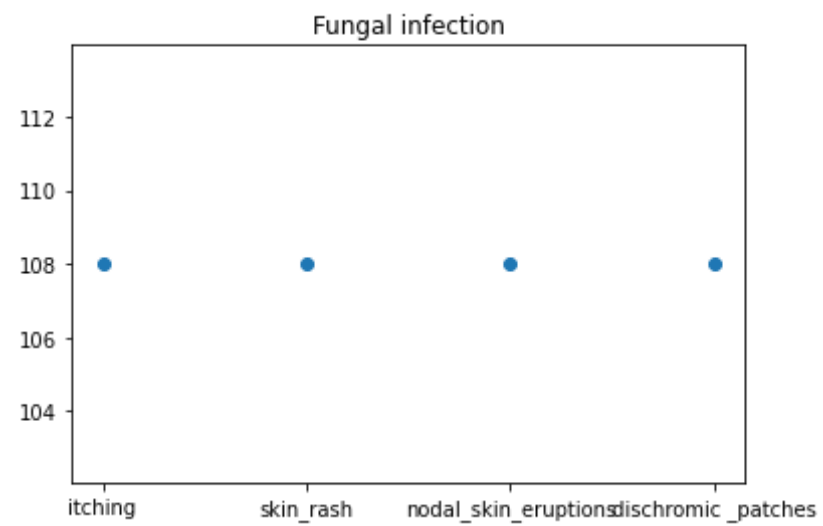
kNearest Neighbour
Accuracy
0.926829268292683
38

```

```

Confusion matrix
[[1 0 0 ... 0 0 0]
 [0 1 0 ... 0 0 0]
 [0 0 1 ... 0 0 0]
...
[0 0 0 ... 1 0 0]
[0 0 0 ... 0 1 0]
[0 0 0 ... 0 0 1]]
[108 108 108 108]
4
4

```



In []: