

# Title: *Personalized Medical Recommendation System with Machine Learning*

## Description:

Welcome to our cutting-edge Personalized Medical Recommendation System, a powerful platform designed to assist users in understanding and managing their health. Leveraging the capabilities of machine learning, our system analyzes user-input symptoms to predict potential diseases accurately.

### Importing required modules

```
In [6]: import pandas as pd
import numpy as np
import seaborn as sns
import matplotlib.pyplot as plt
from sklearn.utils import shuffle
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import LabelEncoder
from sklearn.svm import SVC
from sklearn.ensemble import GradientBoostingClassifier, RandomForestClassifier
from sklearn.naive_bayes import MultinomialNB
from sklearn.neighbors import KNeighborsClassifier
from sklearn.metrics import accuracy_score, confusion_matrix, f1_score
import warnings
warnings.filterwarnings("ignore")
```

### Loading Dataset

```
In [7]: df = pd.read_csv("Training.csv")
```

### Viewing samples to check the data

```
In [8]: df.head(5)
```

Out[8]:

	itching	skin_rash	nodal_skin_eruptions	continuous_sneezing	shivering	chills	joint_pain	stomach_pain	acidity	ulcers_on_tongue	...	blackheads	se
0	1	1		1	0	0	0	0	0	0	0	...	0
1	0	1		1	0	0	0	0	0	0	0	...	0
2	1	0		1	0	0	0	0	0	0	0	...	0
3	1	1		0	0	0	0	0	0	0	0	...	0
4	1	1		1	0	0	0	0	0	0	0	...	0

5 rows × 133 columns

In above dataframe, we can see the data is arranged label wise. So we need to shuffle it first\_

In [9]: df = shuffle(df)

In [10]: df.head()

Out[10]:

	itching	skin_rash	nodal_skin_eruptions	continuous_sneezing	shivering	chills	joint_pain	stomach_pain	acidity	ulcers_on_tongue	...	blackheads	se
1241	0	0		0	1	1	0	0	0	0	0	...	0
4377	0	0		0	0	0	0	0	0	0	0	...	0
4149	0	0		0	0	0	0	0	0	0	0	...	0
2212	1	0		0	0	0	0	0	0	0	0	...	0
376	0	1		0	0	0	0	0	0	0	0	...	1

5 rows × 133 columns

Checking unique values of target variable (Diseases which to be predict)\_

In [11]: df.prognosis.nunique()

```
Out[11]: 41
```

```
In [12]: X = df.drop("prognosis", axis=1)
y = df.prognosis
```

```
In [13]: le = LabelEncoder()
y = le.fit_transform(y)
```

### *\_Checking the number provided to each prognosis by label encoder\_*

```
In [14]: print("Label mapping:")
for label, integer_value in zip(le.classes_, le.transform(le.classes_)):
    print(f"{label}: {integer_value}")
```

Label mapping:

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

*Split the data into 4 Variables, i.e. X\_train, X\_test, y\_train, y\_test*

```
In [15]: X_train, X_test, y_train, y_test = train_test_split(X,y, test_size=0.3, random_state=30)
```

```
In [16]: X_train.shape, X_test.shape, y_train.shape, y_test.shape
```

```
Out[16]: ((3444, 132), (1476, 132), (3444,), (1476,))
```

### *\_Tried applying different classifiers to get the best out of it\_*

```
In [17]: models = {  
    "SVC": SVC(kernel="linear"),  
    "GradientBoost": GradientBoostingClassifier(n_estimators=120, random_state=42),  
    "RandomForest": RandomForestClassifier(n_estimators=120, random_state=42),  
    "Multinomial" : MultinomialNB(),  
    "KNearestNeighbors": KNeighborsClassifier(n_neighbors=5)  
}
```

```
In [18]: for model_name, model in models.items():  
    print(model_name, ":", model)  
  
SVC : SVC(kernel='linear')  
GradientBoost : GradientBoostingClassifier(n_estimators=120, random_state=42)  
RandomForest : RandomForestClassifier(n_estimators=120, random_state=42)  
Multinomial : MultinomialNB()  
KNearestNeighbors : KNeighborsClassifier()
```

### *\_Checking Accuracy, F1\_Score, and Confusion matrix for each model\_*

```
In [19]: for model_name, model in models.items():  
    model.fit(X_train, y_train)  
    pred = model.predict(X_test)  
  
    Accuracy = accuracy_score(y_test, pred)  
    F1_score = f1_score(y_test, pred, average="weighted")  
    cm = confusion_matrix(y_test, pred)  
  
    print(f"{model_name} Accuarcy {Accuracy}")  
    print(f"{model_name} F1 Score{F1_score}")  
    print(f"{model_name} Confusion Matrix:")  
    print(np.array2string(cm, separator=","))  
    print()
```

SVC Accuarcy 1.0  
SVC F1 Score1.0  
SVC Confusion Matrix:  
[[24, 0, 0,..., 0, 0, 0],  
 [ 0,37, 0,..., 0, 0, 0],  
 [ 0, 0,29,..., 0, 0, 0],  
 ...,  
 [ 0, 0, 0,...,42, 0, 0],  
 [ 0, 0, 0,..., 0,36, 0],  
 [ 0, 0, 0,..., 0, 0,41]]

GradientBoost Accuarcy 1.0  
GradientBoost F1 Score1.0  
GradientBoost Confusion Matrix:  
[[24, 0, 0,..., 0, 0, 0],  
 [ 0,37, 0,..., 0, 0, 0],  
 [ 0, 0,29,..., 0, 0, 0],  
 ...,  
 [ 0, 0, 0,...,42, 0, 0],  
 [ 0, 0, 0,..., 0,36, 0],  
 [ 0, 0, 0,..., 0, 0,41]]

RandomForest Accuarcy 1.0  
RandomForest F1 Score1.0  
RandomForest Confusion Matrix:  
[[24, 0, 0,..., 0, 0, 0],  
 [ 0,37, 0,..., 0, 0, 0],  
 [ 0, 0,29,..., 0, 0, 0],  
 ...,  
 [ 0, 0, 0,...,42, 0, 0],  
 [ 0, 0, 0,..., 0,36, 0],  
 [ 0, 0, 0,..., 0, 0,41]]

Multinomial Accuarcy 1.0  
Multinomial F1 Score1.0  
Multinomial Confusion Matrix:  
[[24, 0, 0,..., 0, 0, 0],  
 [ 0,37, 0,..., 0, 0, 0],  
 [ 0, 0,29,..., 0, 0, 0],  
 ...,  
 [ 0, 0, 0,...,42, 0, 0],  
 [ 0, 0, 0,..., 0,36, 0],  
 [ 0, 0, 0,..., 0, 0,41]]

KNearestNeighbors Accuarcy 1.0  
KNearestNeighbors F1 Score1.0

KNearestNeighbors Confusion Matrix:

```
[[24, 0, 0,..., 0, 0, 0],  
 [ 0,37, 0,..., 0, 0, 0],  
 [ 0, 0,29,..., 0, 0, 0],  
 ...,  
 [ 0, 0, 0,...,42, 0, 0],  
 [ 0, 0, 0,..., 0,36, 0],  
 [ 0, 0, 0,..., 0, 0,41]]
```

### *Chosed Random Forest Classifier among all the models*

```
In [20]: rfc = RandomForestClassifier(n_estimators=120, random_state=42)  
rfc.fit(X_train, y_train)  
predictions = rfc.predict(X_test)  
accuracy_score(y_test, predictions)
```

Out[20]: 1.0

### *Saving the model*

```
In [21]: import pickle  
pickle.dump(rfc, open("rfc.pkl","wb"))
```

### *Loading the model for prediction on the basis of user input*

```
In [22]: rfc = pickle.load(open("rfc.pkl","rb"))
```

```
In [23]: input1 = X_test.iloc[0].values.reshape(1,-1)
```

```
In [24]: print("Predicted Label: ", rfc.predict(input1))  
print("Actual Label: ", y_test[0])
```

Predicted Label: [40]  
Actual Label: 40

### *Loading the supportive datasets to give the complete solution on predicted disease*

```
In [25]: Symptoms_disease = pd.read_csv("symtoms_df.csv")  
precautions = pd.read_csv("precautions_df.csv")  
workout = pd.read_csv("workout_df.csv")  
description = pd.read_csv("description.csv")
```

```
medication = pd.read_csv("medications.csv")
diet = pd.read_csv("diets.csv")
```

```
In [26]: symptoms_dict = {'itching': 0, 'skin_rash': 1, 'nodal_skin_eruptions': 2, 'continuous_sneezing': 3, 'shivering': 4, 'chills': 5, 'joint_pai
diseases_list = {15: 'Fungal infection', 4: 'Allergy', 16: 'GERD', 9: 'Chronic cholestasis', 14: 'Drug Reaction', 33: 'Peptic ulcer diseae'
```

*Please select the number corresponding to the symptoms you're experiencing to determine the associated disease.*

```
In [27]: def get_predicted_value(patient_symptoms):
    input_vector = np.zeros(len(symptoms_dict))
    for item in patient_symptoms:
        input_vector[symptoms_dict[item]] = 1
    return diseases_list[rfc.predict([input_vector])[0]]

print("Select symptoms by entering their corresponding numbers separated by commas:")
for index, symptom in symptoms_dict.items():
    print(f"{symptom}: {index}")

selected_symptoms_input = input("Enter the numbers of the symptoms you're experiencing (separated by commas): ")

selected_symptom_nums = selected_symptoms_input.split(',')
selected_symptoms = []

for num in selected_symptom_nums:
    if num.strip().isdigit() and int(num.strip()) in range(len(symptoms_dict)):
        selected_symptoms.append(list(symptoms_dict.keys())[int(num.strip())])
    else:
        print(f"Invalid input: {num}. Skipping...")

print("Selected symptoms:")
for symptom in selected_symptoms:
    print(symptom)

predicted_disease = get_predicted_value(selected_symptoms)

print("Predicted Disease:", predicted_disease)
```

Select symptoms by entering their corresponding numbers separated by commas:

- 0: itching
- 1: skin\_rash
- 2: nodal\_skin\_eruptions
- 3: continuous\_sneezing
- 4: shivering
- 5: chills
- 6: joint\_pain
- 7: stomach\_pain
- 8: acidity
- 9: ulcers\_on\_tongue
- 10: muscle\_wasting
- 11: vomiting
- 12: burning\_micturition
- 13: spotting\_urination
- 14: fatigue
- 15: weight\_gain
- 16: anxiety
- 17: cold\_hands\_and\_feet
- 18: mood\_swings
- 19: weight\_loss
- 20: restlessness
- 21: lethargy
- 22: patches\_in\_throat
- 23: irregular\_sugar\_level
- 24: cough
- 25: high\_fever
- 26: sunken\_eyes
- 27: breathlessness
- 28: sweating
- 29: dehydration
- 30: indigestion
- 31: headache
- 32: yellowish\_skin
- 33: dark\_urine
- 34: nausea
- 35: loss\_of\_appetite
- 36: pain\_behind\_the\_eyes
- 37: back\_pain
- 38: constipation
- 39: abdominal\_pain
- 40: diarrhoea
- 41: mild\_fever
- 42: yellow\_urine
- 43: yellowing\_of\_eyes
- 44: acute\_liver\_failure

45: fluid\_overload  
46: swelling\_of\_stomach  
47: swelled\_lymph\_nodes  
48: malaise  
49: blurred\_and\_distorted\_vision  
50: phlegm  
51: throat\_irritation  
52: redness\_of\_eyes  
53: sinus\_pressure  
54: runny\_nose  
55: congestion  
56: chest\_pain  
57: weakness\_in\_limbs  
58: fast\_heart\_rate  
59: pain\_during\_bowel\_movements  
60: pain\_in\_anal\_region  
61: bloody\_stool  
62: irritation\_in\_anus  
63: neck\_pain  
64: dizziness  
65: cramps  
66: bruising  
67: obesity  
68: swollen\_legs  
69: swollen\_blood\_vessels  
70: puffy\_face\_and\_eyes  
71: enlarged\_thyroid  
72: brittle\_nails  
73: swollen\_extremeties  
74: excessive\_hunger  
75: extra\_marital\_contacts  
76: drying\_and\_tingling\_lips  
77: slurred\_speech  
78: knee\_pain  
79: hip\_joint\_pain  
80: muscle\_weakness  
81: stiff\_neck  
82: swelling\_joints  
83: movement\_stiffness  
84: spinning\_movements  
85: loss\_of\_balance  
86: unsteadiness  
87: weakness\_of\_one\_body\_side  
88: loss\_of\_smell  
89: bladder\_discomfort  
90: foul\_smell\_of\_urine

91: continuous\_feel\_of\_urine  
92: passage\_of\_gases  
93: internal\_itching  
94: toxic\_look\_(typhos)  
95: depression  
96: irritability  
97: muscle\_pain  
98: altered\_sensorium  
99: red\_spots\_over\_body  
100: belly\_pain  
101: abnormal\_menstruation  
102: dischromic\_patches  
103: watering\_from\_eyes  
104: increased\_appetite  
105: polyuria  
106: family\_history  
107: mucoid\_sputum  
108: rusty\_sputum  
109: lack\_of\_concentration  
110: visual\_disturbances  
111: receiving\_blood\_transfusion  
112: receiving\_unsterile\_injections  
113: coma  
114: stomach\_bleeding  
115: distention\_of\_abdomen  
116: history\_of\_alcohol\_consumption  
117: fluid\_overload.1  
118: blood\_in\_sputum  
119: prominent\_veins\_on\_calf  
120: palpitations  
121: painful\_walking  
122: pus\_filled\_pimples  
123: blackheads  
124: scurring  
125: skin\_peeling  
126: silver\_like\_dusting  
127: small\_dents\_in\_nails  
128: inflammatory\_nails  
129: blister  
130: red\_sore\_around\_nose  
131: yellow\_crust\_ooze

Selected symptoms:  
continuous\_sneezing  
joint\_pain  
ulcers\_on\_tongue  
nodal\_skin\_eruptions  
Predicted Disease: Fungal infection

```
In [32]: def helper(dis):
    # Get disease description
    desc = description[description['Disease'] == dis]['Description']
    desc = " ".join([w for w in desc])

    # Get precautions
    pre = precautions[precautions['Disease'] == dis][['Precaution_1', 'Precaution_2', 'Precaution_3', 'Precaution_4']]
    pre = [col for col in pre.values]

    # Get medication
    med = medication[medication['Disease'] == dis]['Medication']
    med = [med for med in med.values]

    # Get diet
    die = diet[diet['Disease'] == dis]['Diet']
    die = [die for die in die.values]

    # Get workout
    wrkout = workout[workout['disease'] == dis]['workout']

    return desc, pre, med, die, wrkout

desc, pre, med, die, wrkout = helper(predicted_disease)
```

*\_Here's the full solution for your illness. Take care!\_*

```
In [33]: desc, pre, med, die, wrkout = helper(predicted_disease)

print("=====predicted disease=====")
print(predicted_disease, "\n")
print("=====description=====")
print(desc, "\n")
print("=====precautions=====")
i = 1
for p_i in pre[0]:
    print(i, ": ", p_i)
    i += 1
print()
```

```
print("=====medications=====")  
for m_i in med:  
    print(i, ":", m_i)  
    i += 1  
print()  
print("=====workout=====")  
for w_i in wrkout:  
    print(i, ":", w_i)  
    i += 1  
print()  
print("=====diets=====")  
for d_i in die:  
    print(i, ":", d_i)  
    i += 1  
  
=====predicted disease=====  
Fungal infection  
  
=====description=====  
Fungal infection is a common skin condition caused by fungi.  
  
=====precautions=====  
1 : bath twice  
2 : use detol or neem in bathing water  
3 : keep infected area dry  
4 : use clean cloths  
  
=====medications=====  
5 : ['Antifungal Cream', 'Fluconazole', 'Terbinafine', 'Clotrimazole', 'Ketoconazole']  
  
=====workout=====  
6 : Avoid sugary foods  
7 : Consume probiotics  
8 : Increase intake of garlic  
9 : Include yogurt in diet  
10 : Limit processed foods  
11 : Stay hydrated  
12 : Consume green tea  
13 : Eat foods rich in zinc  
14 : Include turmeric in diet  
15 : Eat fruits and vegetables  
  
=====diets=====  
16 : ['Antifungal Diet', 'Probiotics', 'Garlic', 'Coconut oil', 'Turmeric']
```

**THANK YOU**