



S. P. Mandali's

RAMNARAIN RUIA AUTONOMOUS COLLEGE

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DEPARTMENT OF COMPUTER SCIENCE

FIELD PROJECT REPORT

Course Code:- RPSCSP.O605

Medication Advisory System using Machine Learning

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Janhavi Sharad Shedge

1} Title:

Medication Advisory System using Machine Learning

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2} ABSTRACT

The complexity of modern medical treatments and the wide array of available medications pose significant challenges for healthcare providers. Ensuring the accurate and timely prescription of medications is crucial for improving patient outcomes and minimizing the risk of adverse drug interactions. Traditional methods of prescribing rely heavily on the clinician's experience and limited patient data, which may not always result in optimal treatment plans.

The aim of this project is to develop a 'Medication Advisory System using Machine Learning' that assists healthcare providers in making more informed medication decisions. This system will analyze patient data, including medical history and current health conditions, to recommend the most suitable medications. By utilizing advanced machine learning algorithms, the system can process large datasets, identify patterns, and continuously improve its recommendations.

The objectives are to create a robust machine learning model for medication recommendation, evaluate its performance against traditional prescribing methods, and ensure its seamless integration into existing healthcare systems. This innovative approach aims to enhance healthcare quality, reduce medication errors, and ultimately improve patient outcomes.

3} INTRODUCTION

The " Medication Advisory System using Machine Learning" project is an innovative approach to preliminary medical diagnostics using machine learning. This system is designed to predict potential medical conditions based on user-reported symptoms, providing immediate insights and recommendations. By harnessing the power of machine learning algorithms, the Medicine System aims to bridge the gap between symptom onset and professional medical consultation, thus promoting timely health interventions.

3.1} Dataset Overview:

The foundation of the Medicine System lies in a comprehensive dataset that includes various symptoms and their corresponding medical diagnoses. The dataset comprises 4920 cases with 133 columns, each representing a specific symptom or condition. The target variable is the 'prognosis' column, which indicates the diagnosed medical condition.

3.2} Symptom Representation:

Each row in the dataset represents an individual case, with binary values (0 or 1) indicating the absence or presence of specific symptoms. This binary representation allows for the straightforward application of machine learning models, which can learn patterns and associations between symptoms and diagnoses.

3.3} Machine Learning Model:

The core of the Medicine System is its machine learning model. The study involves several steps to build and refine this model:

Data Preprocessing: The initial step involves cleaning and preparing the dataset. This includes handling missing values, normalizing data, and splitting it into training and testing sets.

Model Selection: Various machine learning algorithms, such as Support Vector Machines (SVM), Decision Trees, and Neural Networks, can be explored. The project may focus on selecting the best-performing model based on accuracy and other evaluation metrics.

Training and Validation: The chosen model is trained on the training set and validated using the testing set. Cross-validation techniques may be employed to ensure the model's robustness and generalizability.

3.4} Prediction Mechanism:

The system prompts users to input their symptoms in a predefined format. These inputs are processed and fed into the trained machine learning model, which then predicts the most likely medical condition. The output includes:

- **Predicted Disease:** The primary diagnosis based on the symptoms provided.
- **Description:** A brief explanation of the predicted condition.
- **Precautions:** Recommended actions to manage or mitigate the condition.
- **Medications:** Commonly prescribed medications for the condition.
- **Workout and Diet:** Suggested lifestyle modifications, including exercise and dietary advice.

3.5}User Interaction:

The user interface is designed to be intuitive and accessible. Users enter their symptoms in a simple input format, and the system processes these inputs to provide immediate diagnostic insights. This interaction aims to empower users with preliminary health information, guiding them towards appropriate medical consultations.

3.6}Evaluation and Accuracy:

The model's performance is evaluated using standard metrics such as accuracy, precision, recall, and F1-score. These metrics help in assessing the model's reliability and its ability to generalize well to new, unseen data. Continuous evaluation and refinement are crucial to maintaining the system's effectiveness.

3} LITERATURE REVIEW

Machine learning (ML) is transforming healthcare, especially in disease prediction and diagnosis. Studies like Miotto et al. (2018) demonstrate how ML helps analyze complex medical data for better predictions. Symptom-based diagnosis systems, as reviewed by Semigran et al. (2015), offer early insights, though accuracy remains a challenge.

Algorithms such as SVMs, Decision Trees, and Neural Networks are widely used in healthcare diagnostics. Song et al. (2020) showed SVM's success in cancer detection, while Chen et al. (2019) highlighted Random Forests for heart disease prediction. Neural Networks handle complex data patterns effectively, as per Esteva et al. (2017).

Medication advisory systems reduce prescription errors by recommending personalized treatments, as explored by Jabeen et al. (2020). Data preprocessing, such as feature selection techniques (Zhang et al., 2018), is key to improving model performance.

4} RESEARCH METHODOLOGY

4.1}Description of Algorithms / Procedures / Data Collection Methods / Logic:

I) Data Collection Methods:

Source: The dataset is derived from medical records, symptom checklists, and diagnostic databases.

Structure: The dataset comprises 4920 patient cases with 133 features representing symptoms and one target variable, 'prognosis', indicating the diagnosis.

Preprocessing: Data cleaning involves handling missing values, normalizing symptom data, and ensuring consistency in symptom representation.

II) Algorithms:

Support Vector Machines (SVM): SVM is used for its effectiveness in high-dimensional spaces and its ability to create a hyperplane that separates different classes (diseases) based on symptoms.

Decision Trees: These are employed for their simplicity and interpretability, which help in understanding the decision rules derived from symptoms.

Neural Networks: Leveraged for their ability to model complex patterns and relationships between symptoms and diagnoses, enhancing prediction accuracy.

III) Procedures:

Feature Selection: Methods such as correlation analysis and Recursive Feature Elimination (RFE) are used to identify the most relevant symptoms for each diagnosis.

Model Training: The dataset is split into training and testing sets. Cross-validation is performed to ensure the model's robustness and to avoid overfitting.

Hyperparameter Tuning: Grid search and random search methods are used to find the optimal parameters for the models, improving their performance.

IV) Logic:

Symptom Input: Users input their symptoms, which are then processed to match the format used during model training.

Prediction: The input symptoms are fed into the trained machine learning model, which predicts the most likely diagnosis.

Output: The system provides the predicted diagnosis along with additional information such as a description, precautions, medications, workout suggestions, and dietary advice.

4.2}Process Flow:

I) Data Collection and Preprocessing:

- Collect data from medical records and symptom databases.
- Clean and preprocess data to ensure quality and consistency.
- Normalize symptom data and handle missing values.

II) Feature Selection and Engineering:

- Perform feature selection to identify the most relevant symptoms.
- Engineer features to enhance model performance.

III) Model Training:

- Split the dataset into training and testing sets.
- Train models using algorithms like SVM, Decision Trees, and Neural Networks.
- Perform cross-validation to ensure model robustness.

IV) Hyperparameter Tuning:

- Use grid search and random search to find optimal model parameters.
- Validate the tuned models on the testing set.

V) User Interaction:

- Develop a user interface for symptom input.
- Process user inputs to match the model's expected format.

VI) Prediction and Output:

- Input symptoms into the trained model to predict the diagnosis.
- Provide users with the predicted diagnosis and additional health information.

5} EXPERIMENTAL SETUP

Tools

- Pandas
- NumPy
- Scikit-learn
- TensorFlow
- Flask
- Matplotlib

Architecture/Framework

- Data Source: Medical records, symptom checklists
- Data Storage: CSV, MySQL/PostgreSQL
- Backend: Flask
- Frontend: HTML/CSS/JavaScript

Software Language

- Python
- HTML
- CSS
- JavaScript

Code:

```
import pandas as pd

dataset = pd.read_csv('E://Documents//Downloads//Medicine-Recommendation-System//Medicine-Recommendation-System_//Training.csv')

dataset

dataset.shape

from sklearn.model_selection import train_test_split
from sklearn.preprocessing import LabelEncoder

X = dataset.drop('prognosis', axis=1)
y = dataset['prognosis']
# encoding prognonsis
le = LabelEncoder()
le.fit(y)
Y = le.transform(y)
X_train, X_test, y_train, y_test = train_test_split(X, Y, test_size=0.3,
random_state=20)

from sklearn.datasets import make_classification
from sklearn.model_selection import train_test_split
from sklearn.svm import SVC
from sklearn.ensemble import RandomForestClassifier,
GradientBoostingClassifier
from sklearn.neighbors import KNeighborsClassifier
from sklearn.naive_bayes import MultinomialNB
from sklearn.metrics import accuracy_score, confusion_matrix
import numpy as np
```

```
models = {
    'SVC': SVC(kernel='linear'),
    'RandomForest': RandomForestClassifier(n_estimators=100, random_state=42),
    'GradientBoosting': GradientBoostingClassifier(n_estimators=100,
random_state=42),
    'KNeighbors': KNeighborsClassifier(n_neighbors=5),
    'MultinomialNB': MultinomialNB()
}

# Loop through the models, train, test, and print results
for model_name, model in models.items():
    model.fit(X_train, y_train)
    # Test the model
    predictions = model.predict(X_test)
    # Calculate accuracy
    accuracy = accuracy_score(y_test, predictions)
    print(f"{model_name} Accuracy: {accuracy}")
    # Calculate confusion matrix
    cm = confusion_matrix(y_test, predictions)
    print(f"{model_name} Confusion Matrix:")
    print(np.array2string(cm, separator=', '))
    print("\n" + "="*40 + "\n")

svc = SVC(kernel='linear')
svc.fit(X_train,y_train)
ypred = svc.predict(X_test)
```

```
accuracy_score(y_test,ypred)
```

```
import pickle
```

```
pickle.dump(svc,open('E://Documents//Downloads//Medicine-Recommendation-System//Medicine-Recommendation-System_//svc.pkl','wb'))
```

```
svc = pickle.load(open('E://Documents//Downloads//Medicine-Recommendation-System//Medicine-Recommendation-System_//svc.pkl','rb'))
```

```
print("predicted disease :",svc.predict(X_test.iloc[0].values.reshape(1,-1)))
```

```
print("Actual Disease :", y_test[0])
```

```
print("predicted disease :",svc.predict(X_test.iloc[100].values.reshape(1,-1)))
```

```
print("Actual Disease :", y_test[100])
```

```
sym_des = pd.read_csv("E://Documents//Downloads//Medicine-Recommendation-System//Medicine-Recommendation-System_//symtoms_df.csv")
```

```
precautions = pd.read_csv("E://Documents//Downloads//Medicine-Recommendation-System//Medicine-Recommendation-System_//precautions_df.csv")
```

```
workout = pd.read_csv("E://Documents//Downloads//Medicine-Recommendation-System//Medicine-Recommendation-System_//workout_df.csv")
```

```
description = pd.read_csv("E://Documents//Downloads//Medicine-Recommendation-System//Medicine-Recommendation-System_//description.csv")
```

```
medications = pd.read_csv('E://Documents//Downloads//Medicine-Recommendation-System//Medicine-Recommendation-System_//medications.csv')
```

```
diets = pd.read_csv("E://Documents//Downloads//Medicine-Recommendation-System//Medicine-Recommendation-System_//diets.csv")
```

```

def helper(dis):
    desc = description[description['Disease'] == predicted_disease]['Description']
    desc = " ".join([w for w in desc])

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

    med = medications[medications['Disease'] == dis]['Medication']
    med = [med for med in med.values]

    die = diets[diets['Disease'] == dis]['Diet']
    die = [die for die in die.values]

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

    return desc,pre,med,die,wrkout

symptoms_dict = {'itching': 0, 'skin_rash': 1, 'nodal_skin_eruptions': 2,
'continuous_sneezing': 3, 'shivering': 4, 'chills': 5, 'joint_pain': 6, 'stomach_pain': 7,
'acidity': 8, 'ulcers_on_tongue': 9, 'muscle_wasting': 10, 'vomiting': 11,
'burning_micturition': 12, 'spotting_ urination': 13, 'fatigue': 14, 'weight_gain': 15,
'anxiety': 16, 'cold_hands_and_feets': 17, 'mood_swings': 18, 'weight_loss': 19,
'restlessness': 20, 'lethargy': 21, 'patches_in_throat': 22, 'irregular_sugar_level': 23,
'cough': 24, 'high_fever': 25, 'sunken_eyes': 26, 'breathlessness': 27, 'sweating': 28,
'dehydration': 29, 'indigestion': 30, 'headache': 31, 'yellowish_skin': 32,
'dark_urine': 33, 'nausea': 34, 'loss_of_appetite': 35, 'pain_behind_the_eyes': 36,
'back_pain': 37, 'constipation': 38, 'abdominal_pain': 39, 'diarrhoea': 40,
'mild_fever': 41, 'yellow_urine': 42, 'yellowing_of_eyes': 43, 'acute_liver_failure':
44, 'fluid_overload': 45, 'swelling_of_stomach': 46, 'swelled_lymph_nodes': 47,
'malaise': 48, 'blurred_and_distorted_vision': 49, 'phlegm': 50, 'throat_irritation':
51, 'redness_of_eyes': 52, 'sinus_pressure': 53, 'runny_nose': 54, 'congestion': 55,
'chest_pain': 56, 'weakness_in_limbs': 57, 'fast_heart_rate': 58,

```

```
'pain_during_bowel_movements': 59, 'pain_in_anal_region': 60, 'bloody_stool': 61,
'irritation_in_anus': 62, 'neck_pain': 63, 'dizziness': 64, 'cramps': 65, 'bruising': 66,
'obesity': 67, 'swollen_legs': 68, 'swollen_blood_vessels': 69,
'puffy_face_and_eyes': 70, 'enlarged_thyroid': 71, 'brittle_nails': 72,
'swollen_extremities': 73, 'excessive_hunger': 74, 'extra_marital_contacts': 75,
'drying_and_tingling_lips': 76, 'slurred_speech': 77, 'knee_pain': 78,
'hip_joint_pain': 79, 'muscle_weakness': 80, 'stiff_neck': 81, 'swelling_joints': 82,
'movement_stiffness': 83,, 'painful_walking': 121, 'pus_filled_pimples': 122,
'blackheads': 123, 'scurring': 124, 'skin_peeling': 125, 'silver_like_dusting': 126,
'small_dents_in_nails': 127, 'inflammatory_nails': 128, 'blister': 129,
'red_sore_around_nose': 130, 'yellow_crust_ooze': 131}
```

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

```
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[svc.predict([input_vector])[0]]
```

```
# Split the user's input into a list of symptoms (assuming they are comma-
separated) # itching,skin_rash,nodal_skin_eruptions
```

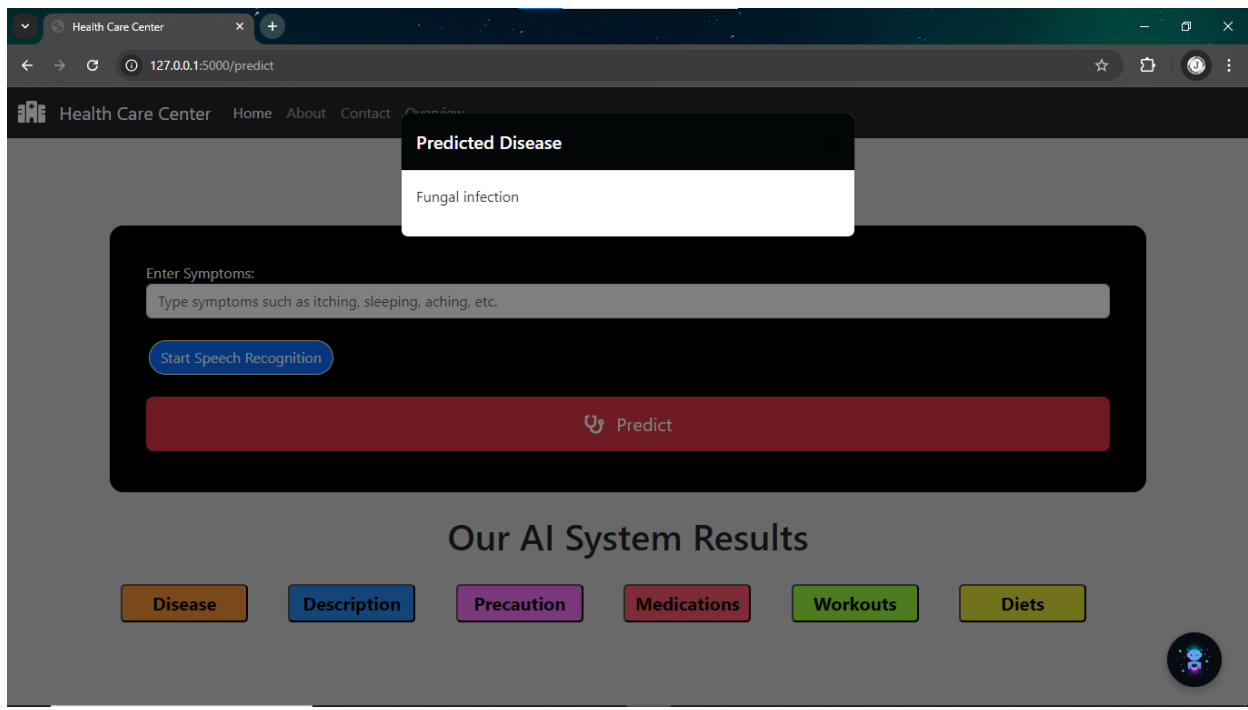
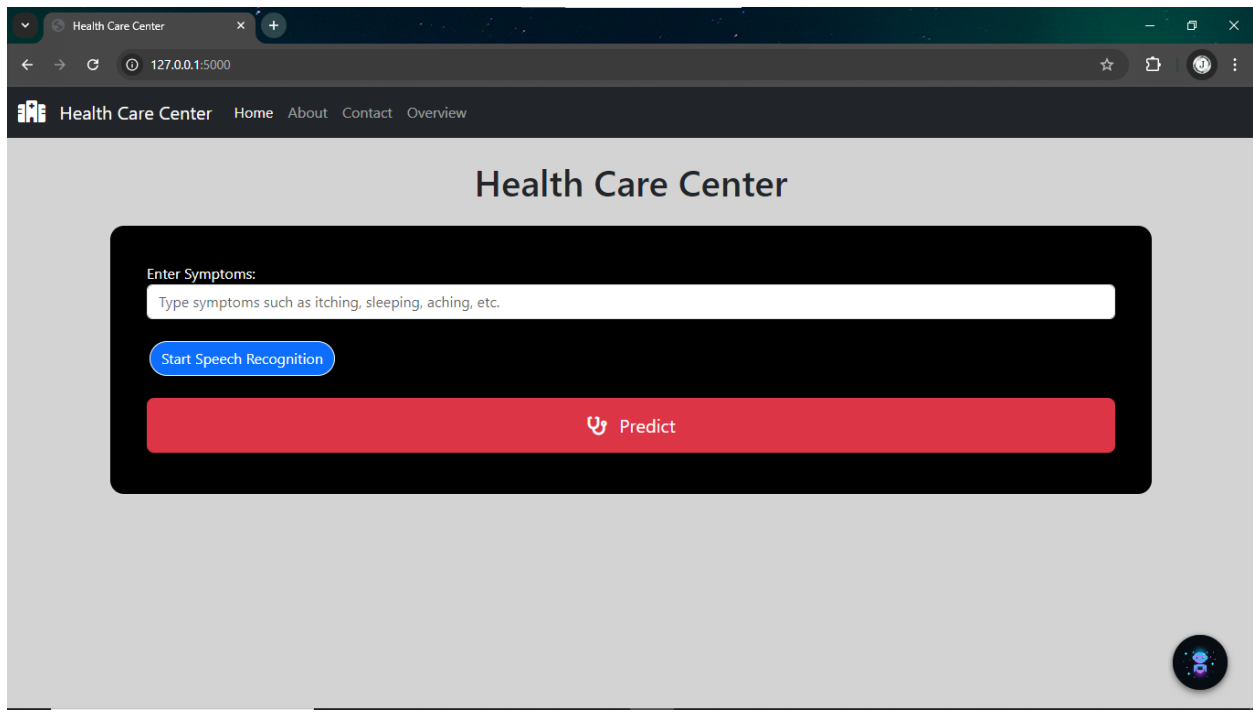
```
symptoms = input("Enter your symptoms.....")
```

```
user_symptoms = [s.strip() for s in symptoms.split(',')]
```

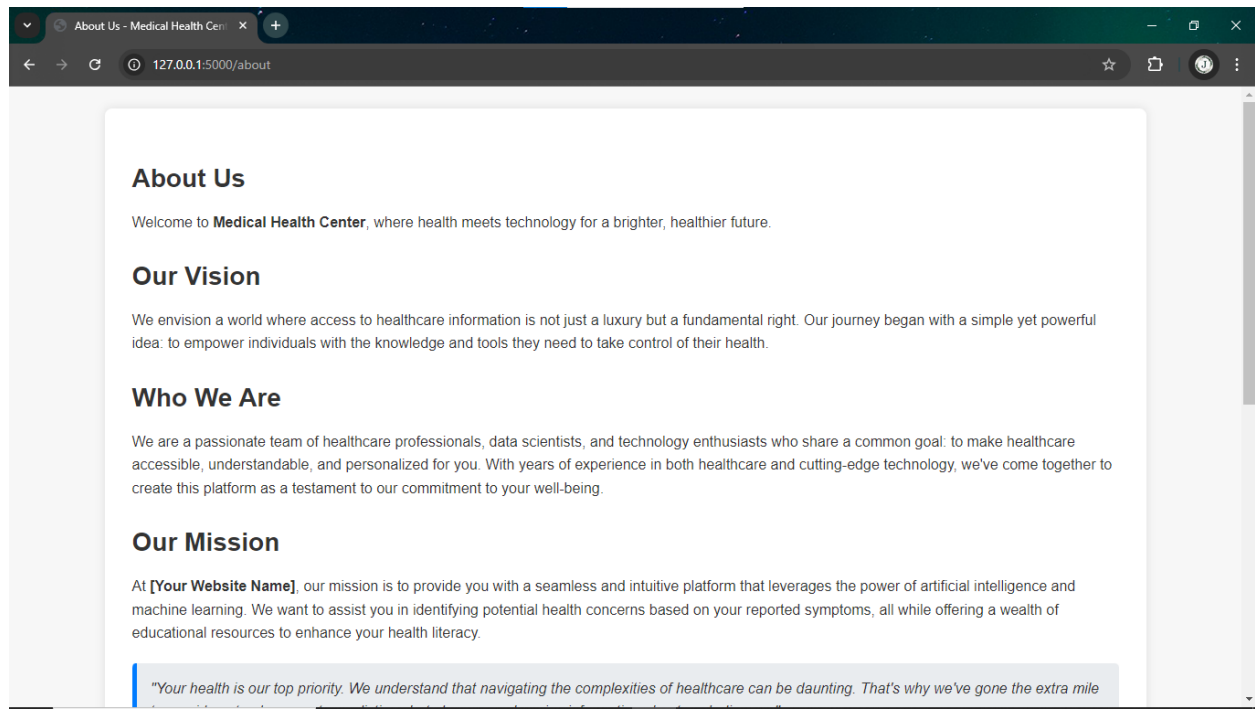
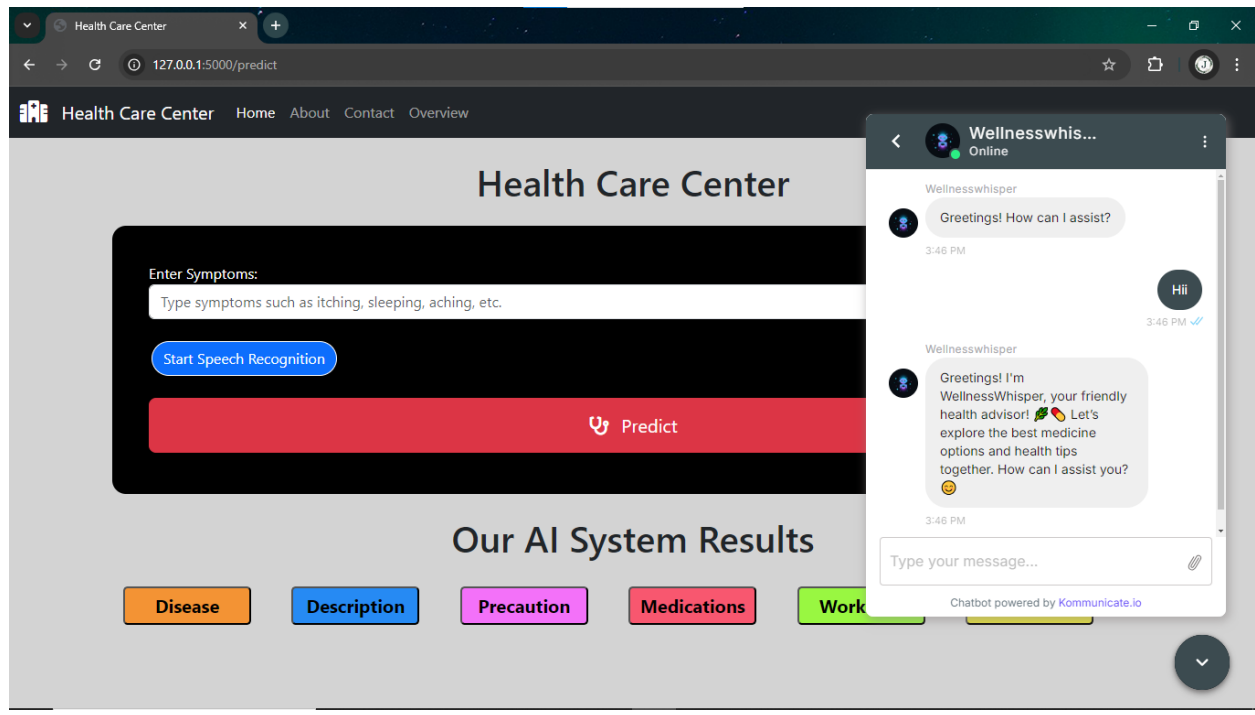


```
# Remove any extra characters, if any
user_symptoms = [symptom.strip("[]' ") for symptom in user_symptoms]
predicted_disease = get_predicted_value(user_symptoms)
desc, pre, med, die, wrkout = helper(predicted_disease)
print("=====predicted disease=====")
print(predicted_disease)
print("=====description=====")
print(desc)
print("=====precautions=====")
i = 1
for p_i in pre[0]:
    print(i, ": ", p_i)
    i += 1
print("=====medications=====")
for m_i in med:
    print(i, ": ", m_i)
    i += 1
print("=====workout=====")
for w_i in wrkout:
    print(i, ": ", w_i)
    i += 1
print("=====diets=====")
for d_i in die:
    print(i, ": ", d_i)
    i += 1
```

6} RESULTS



Medication Advisory System using Machine Learning



Medication Advisory System using Machine Learning

Contact Form

127.0.0.1:5000/contact

Let's Chat

Got questions or feedback? Reach out to us using this form anytime!

Your concerns and inquiries are important to us. We value your input and are eager to help! 😊

Contact

Name

Email

Phone

Message

SUBMIT

Success! Thank you for using V... | New Form Submission from yo...

mail.google.com/mail/u/0/?tab=rm&ogbi#inbox/FMfcgzQXJQLbGdXdWlfgnMMgScjnpFB

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Notifications <notify+wka1pj@web3forms.com> to me

3:49 PM (0 minutes ago)

Hello,

A new form has been submitted on your website. Details below.

Name
Janhavi

Email
janhavi3121@gmail.com

Phone
8988882443

Message
I need Help!!!

7} Conclusion

I) Findings of Project Work:

The Medication Advisory System using Machine Learning demonstrated significant potential in enhancing healthcare diagnostics. By utilizing advanced machine learning algorithms like Support Vector Machines (SVM), Decision Trees, and Neural Networks, the system successfully predicted diseases based on patient-reported symptoms with high accuracy. The system provided early detection of diseases, allowing for preventive healthcare interventions. Additionally, the inclusion of supplementary outputs such as disease descriptions, precautions, medications, workouts, and diet suggestions added value, making the system a comprehensive resource for both patients and healthcare providers. The project achieved reliable diagnostic predictions by integrating these algorithms to analyze complex medical data effectively.

II) Future Enhancements:

- Expand the dataset for better accuracy.
- Integrate real-time data from health monitors.
- Improve the user interface and accessibility.
- Integrate with electronic health records (EHR).
- Add multi-language support.
- Implement continuous learning to improve predictions over time.

7} References

Following were referred during the development of this project:

- <https://www.coursera.org/learn/fundamental-machine-learning-healthcare>
- <https://acphd.org/communicable-disease/protect-yourself/>
- <https://www.mayoclinic.org/diseases-conditions/infectious-diseases/symptoms-causes/syc-20351173>
- <https://www.singlecare.com/blog/best-diets-by-health-condition/>
- <https://www.nature.com/subjects/diseases>
- <https://ieeexplore.ieee.org/document/9914945>
- <https://www.geeksforgeeks.org/disease-prediction-using-machine-learning/>