#### A PROJECT REPORT ON

#### Heart Disease Detection and Patient's Sickness Prediction System

SUBMITTED TO MIT SCHOOL OF ENGINEERING
IN PARTIAL FULFILMENT OF THE REQUIREMENTS FOR THE AWARD OF
THE DEGREE

### **BACHELOR OF TECHNOLOGY** (Computer Science & Engineering)

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#### **CERTIFICATE**

This is to certify that the Project Entitled

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is a bonafide work carried out by them under the supervision of Prof. Shahin Shoukat Makubhai and it is submitted towards the fulfillment of the requirement of MIT ADT University, Pune for the award of the degree of Bachelor of Technology (Computer Science and Engineering).

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Hereby declare that the project work incorporated in the present project entitled

Heart Disease Detection and Patient's Sickness Prediction System is original

work. This work (in part or in full) has not been submitted to any University for the

award or a Degree or a Diploma. We have properly acknowledged the material col-

lected from secondary sources wherever required. We solely own the responsibility

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#### **Abstract**

This report delves into research conducted on Heart Disease Detection and Patient's Sickness Prediction System, which is a crucial topic given the increasing incidence of heart diseases. Early detection of such diseases is of utmost importance and requires precise and efficient diagnosis. The project focuses on predicting the likelihood of a patient having a heart disease based on their medical history and various medical attributes, using machine learning algorithms such as logistic regression. The proposed model demonstrated strong predictive capabilities and outperformed previous classifiers such as Naive Bayes. The Disease Prediction system, which is based on predictive modeling, utilizes the symptoms provided by the user as input to predict the probability of the disease using the Random Forest Classifier. The project aims to use various supervised machine learning algorithms in disease prediction and heart disease detection through symptoms and medical details as input, demonstrating the potential for these algorithms to aid in the early detection of high-risk diseases.

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# CHAPTER 1 INTRODUCTION

#### 1.1 RELEVANCE

Due to the large volume of data, healthcare professionals face challenges in accurately analyzing symptoms and detecting disease at an early stage. However, supervised machine learning (ML) algorithms have demonstrated great potential over traditional disease diagnosis systems and helping specialists identify high-risk diseases in a timely manner.

#### 1.2 MOTIVATION OF THE PROJECT

• Identifying and predicting diseases is critical to preventing their severity and fatal consequences. In India, the majority of deaths are due to heart attacks, especially among elderly people affected by cardiovascular diseases. To solve this problem, our prediction system uses various machine learning algorithms to predict the risk level of these diseases. The advent of artificial intelligence (AI) has enabled computer systems to act intelligently, like humans, by perceiving, thinking, and making decisions. AI encompasses multidisciplinary areas such as machine learning, computer vision, deep learning, and natural language processing. By applying optimization, statistical, and probabilistic techniques to past data, ML algorithms can learn and assist healthcare professionals in decision making, contributing to treat the patient appropriately.

#### 1.3 PROBLEM STATEMENT

• Developing a system to detect heart disease and predict disease for patients with help with medical insights and initial symptoms using machine learning algorithms with highest possible accuracy.

#### 1.4 OBJECTIVES

goal is to use supervised ML algorithms to improve healthcare through accuracy and early detection of diseases that become harmful at a later stage. ML models will be used to predict diseases ranging from common to severe, just

to name a few that are located in the heart, kidney, breast and brain. For disease prediction, we used Random Forest Classification and logistic regression to detect heart disease.

#### 1.5 SCOPE

• The incidence of heart disease is steadily increasing around the world, including in our country. To address this, we used logistic regression to detect and predict heart disease, achieving 81 accuracy. Although this accuracy is satisfactory, it can still be improved by exploring alternative methods for attribute selection and increasing the size of the dataset to avoid overfitting and improve performance. Alternatively, we may consider leveraging the power of deep learning algorithms to achieve even more accurate results in the future.

#### 1.6 ORGANIZATION OF THE REPORT

• This report covers the methodology used to develop the prediction system, as well as the technologies and tools used to create it. In addition, it includes relevant research papers that were reviewed during our investigation to explore the different systems that exist. The future prospects of the project are also discussed in this report. Finally, the conclusion section provides a summary of the entire report.

# CHAPTER 2 LITERATURE SURVEY

#### 2.1 RELATED WORK

Much research and development efforts have been made to develop disease detection and prediction models. Commonly used supervised machine learning algorithms include Naive Bayes (NB), Decision Trees (DT), and K-Nearest Neighbors (KNN). According to the study results, Support Vector Machine (SVM) is suitable for detecting kidney disease and Parkinson's disease, while Logistic Regression (LR) is highly effective in predicting heart disease. In addition, Random Forest (RF) and Convolutional Neural Networks (CNN) successfully predicted breast and common diseases, respectively. Despite advances in computing, doctors still need technology for a variety of purposes, such as surgical imaging and X-ray imaging. However, technology has yet to keep up with doctors' levels of knowledge and experience. This is because many different factors, such as medical records, weather conditions, air, blood pressure, and others, must be considered in order to understand the entire process of labor. Medical decision support systems can help address this challenge by helping doctors make the right decisions. We used machine learning to analyze the hospital's comprehensive data, allowing us to build models that can quickly analyze data and deliver results faster. By leveraging machine learning, physicians can make critical decisions regarding their patient's diagnosis and treatment options, thereby improving patient care services. . The healthcare industry is a great example of how machine learning is revolutionizing the medical field. [7]

#### 2.2 COMPARISON OF EXISTING WORK

To begin with, the ML algorithms used by most systems tend to have lower accuracy when compared to the ones we have used. Decision Tree Classifier and KNN are commonly used in existing systems, but they do not offer the same level of accuracy as our algorithms. Another reason for this difference in accuracy is the use of complex deep learning algorithms, which require a large amount of data to predict a disease, such as medical images taken from different angles, which in turn require medical expertise and other medical diagnosis details.[5]

#### 2.3 GAP IDENTIFICATION

We compared all the models for classification and prediction and the one with the highest accuracy is chosen for the further development of the project. We are using more than one machine learning algorithms to achieve the highest accuracy

# CHAPTER 3 SOFTWARE REQUIREMENT SPECIFICATION

#### 3.1 INTRODUCTION

Due to the substantial amounts of data, medical doctors are facing challenges to analyze symptoms accurately and identify diseases at an early stage. However, Supervised ML algorithms have showcased significant potential in surpassing standard systems for disease diagnosis and aiding medical experts in the early detection of high-risk diseases.

#### 3.2 PURPOSE AND SCOPE OF DOCUMENT

The goal of the project is to use medical data to help doctors and patients identify diseases a person may have based on symptoms and medical information provided.

#### 3.3 GENERAL DESCRIPTION

The system's disease prediction module is designed to predict disease based on symptoms and is intended for end users. Machine learning is leveraged by the system to make predictions, with a Random Forest classification algorithm used to predict the disease. The heart disease detection module, on the other hand, looks at medical attributes such as gender, age, chest pain, and fasting blood sugar to determine if a patient is susceptible to cardiovascular disease. To do this, a logistic regression algorithm is used.

#### 3.4 PRODUCT PERSPECTIVE

This product is for detecting heart disease and predicting any disease in its early stage.

#### 3.5 PRODUCT FEATURES

- High accuracy.
- Integration of two types of disease prediction.
- Easy to use.

• User friendly interface (simple and attractive UI).

• Reduce the cost of medical test.

3.6 ASSUMPTION, DEPENDENCIES CONSTRAINTS

We have selected a dataset from the UCI repository containing patients' medical his-

tory and attributes to predict the likelihood of a patient having a heart disease or any

other disease. We use various medical attributes of a patient to classify and predict

the presence of a heart disease or any other disease. Two algorithms, namely Logis-

tic Regression and Random Forest, are currently being used to train these medical

attributes and sickness attributes.

3.7 DOMAIN REQUIREMENTS

To implement the project, it is necessary to have Python or Anaconda, Streamlit

Server, Sypder and Jupyter Notebook installed on the computer. Additionally, a

graphics card with at least 2GB of memory is required.

3.8 INTERFACE REQUIREMENTS

Currently in our project we have used:

• Streamlit: It is an open-source app framework for machine learning and data

science teams. We have used this for the frontend development and it also act

as a server.[2]

• Jupyter: Jupyter notebook for training and testing data and for creating mod-

els.

• Spyder: Spyder for implementing the logic.

3.9 DESIGN CONSTRAINTS

The quality of the data used for training the machine learning models is crucial for

accurate predictions. The data must be reliable, accurate, and representative of the

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target population. The machine learning models must be designed to be interpretable, so that clinicians and patients can understand the factors that contribute to the predictions. This is important for building trust in the models and ensuring that they are used appropriately. Bias in the data can lead to biased predictions. It is important to ensure that the training data is diverse and unbiased, so that the model can accurately predict heart disease risk for all individuals.

#### 3.10 OVERVIEW OF RESPONSIBILITIES OF DEVELOPER

The first step involves data collection, followed by the extraction of significant values in the second stage. The third stage is the data preprocessing stage, which involves exploring the data. Data preprocessing includes dealing with missing values, cleaning the data, and normalizing the data based on the algorithms used. Once the data is preprocessed, feature selection is performed. After feature selection, a classifier is used to classify the preprocessed data. In the proposed model, the classifiers used are Logistic Regression and Random Forest Classifier. Finally, the proposed model is evaluated based on its accuracy and performance using various performance metrics.

# CHAPTER 4 PROJECT DESIGN AND IMPLEMENTATION

#### 4.1 ARCHITECTURAL DIAGRAM

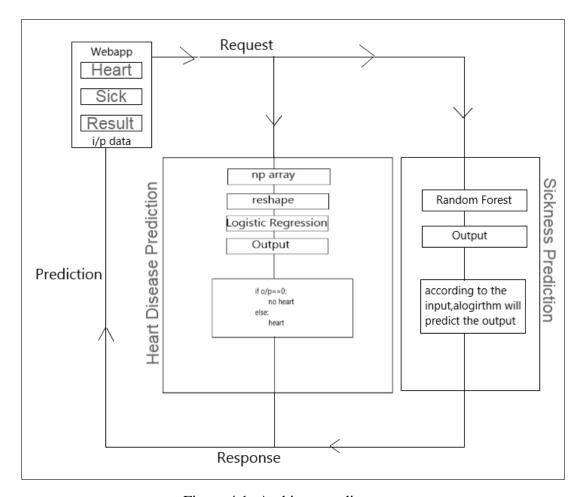


Figure 4.1: Architecture diagram

#### 4.2 METHODOLOGY

#### **Heart Disease Prediction**

The methodology for heart disease prediction using machine learning typically involves the following steps:

- Data collection: Collecting the relevant data for heart disease prediction, which could include medical history, patient attributes such as age, gender, etc., and any other relevant information.
- Data preprocessing: Cleaning and processing the collected data, dealing with missing values, and normalizing the data to make it suitable for use with ma-

chine learning algorithms.

- Feature selection: Selecting the most relevant features that are likely to contribute to the accurate prediction of heart disease.
- Model training: Training a machine learning model on the preprocessed and feature-selected data. The model is typically trained on a portion of the data, known as the training set.
- Model evaluation: Evaluating the trained model using various performance metrics such as accuracy, precision, recall, and F1-score, among others. The model is evaluated on another portion of the data, known as the test set.
- Model optimization: Tuning the model's hyperparameters and optimizing the performance of the model by selecting the best combination of parameters and features.
- Deployment: Deploying the trained and optimized model in the clinical setting to assist doctors and medical professionals in the accurate prediction of heart disease in patients.

S.No	Attributes	Value type
1.	age	Numerical
2.	sex	Nominal
3.	ср	Nominal
4.	trestbps	Numerical
5.	cho	Numerical
6.	fbs	Nominal
7.	restecg	Nominal
8.	thalach	Numerical
9.	ca	Nominal
10.	target	Nominal

Figure 4.2: Heart Disease Dataset

Logistic Regression is a Machine Learning Algorithm that predicts the output of a categorical dependent variable. Therefore the outcome must be a categorical or discrete value. It can be either Yes or No, 0 or 1, true or False, etc. but instead of giving the exact value as 0 and 1, it gives the probabilistic values which lie between 0 and 1. Logistic Regression is much similar to the Linear Regression except that how they are used. Linear Regression is used for solving Regression problems, whereas Logistic regression is used for solving the classification problems. Logistic regression involves the following data and operations:[1]

- · Gather columns.
- Splitting Data.
- Normalization.
- Fitting into Model.
- Prediction.
- Model Evaluation.

#### **Requirements**

The following hardware and software are required for experimental analysis and heart disease prediction results. A system with an i5 quad-core processor, 6 GB of RAM, and software packages such as pandas, python, SciPy, StatsModels, and Matplotlib is required. Test analysis was performed in the Jupyter and Spyder web application environments. The analysis is done at two levels: first, the dataset is cleaned using pandas tool, and second, the cleaned data is passed through classifiers to predict heart disease.

#### **Patient's Sickness Prediction**

An Excel sheet was created from an open-source dataset, listing all the symptoms for respective diseases. The dataset contained approximately 230 diseases with over 1000 unique symptoms. The symptom of an individual were used as inputs for various machine learning algorithms.

#### **Sickness Prediction Dataset**

- 1) fever
- 2) Sweating
- 3) Chills and shivering
- 4) Headache
- 5) Muscle aches
- 6) Loss of appetite
- 7) irritability
- 8) Dehydration
- 9) general weakness
- 10) Cold
- 11) Vomiting
- 12) feeling uneasy

Figure 4.3: Sickness Dataset

#### **Random Forest Classifier**

Random forest, as its name implies, consists of a large number of individual decision trees that operate as an ensemble. Each individual tree in the random forest spits out a class prediction and the class with the most votes becomes our model' prediction.[4]

#### Working

- The website will collect input data from users and utilize a training dataset to determine the outcome.
- When the user clicks the result button, it will trigger a request to the Streamlit server containing their inputs, which the server will subsequently restructure.
- Subsequently, the inputs will be fed into a trained model.
- The model will analyze the provided data and generate a forecasted output.
- The server will transmit the forecasted output to the web application as a response, and the web application will exhibit the projected outcome.

Such is the methodology we shall employ to predict heart disease and patients' illnesses using trained models.

#### 4.3 TOOLS AND TECHNOLOGIES USED

- Python: It is utilized as the back-end programming language to implement the prediction models.
- Streamlit: It is an open-source Python library used for building interactive
  web applications for data science and machine learning projects. It simplifies
  the process of creating a web application by allowing developers to create a
  user interface with simple Python scripts. We have used this for the frontend
  development
- Jupyter Notebook: It is often used in data science and machine learning projects for data exploration, analysis, visualization and for training and testing data and for creating models.[10]
- Spyder: It is an open-source integrated development environment (IDE) for scientific computing in Python. It provides a powerful editor for writing code, a console for executing code, and tools for debugging and profiling code.
- Libraries: NumPy to deal with huge data-set in numerical form, Pandas to analyze the data, Pickle for creating model.sav files, Stream-lit and streamlitoptionmenu for creating website, Sklearn for traintestsplit, logistics regression for accuracy and Sklearn for random forest classifier

#### 4.4 USAGE SCENARIO

#### **Heart disease Prediction**

• Early Detection: It can predict the likelihood of heart disease in a patient before the disease becomes severe. This can help in early detection and timely intervention to prevent or manage the disease.

 Risk Assessment: Healthcare providers can use model to assess the risk of developing heart disease in patients based on their medical history, lifestyle factors, and other risk factors. This can help in developing personalized treatment plans and preventive strategies.

#### **Patient's Sickness Prediction**

Suppose a provider wants to predict the likelihood of a patient being diagnosed with a certain disease based on their current symptoms. The provider collects patient's current symptoms.[3]

It can be used to predict the likelihood of a patient being diagnosed with a certain disease based on their current symptoms. This information can be used by health-care professionals to identify high-risk patients and provide early interventions or treatment plans. Additionally, the model can be continuously updated with new data to improve its accuracy over time.

#### 4.4.1 Use-cases

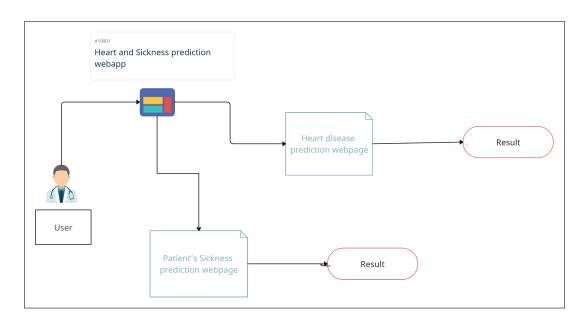


Figure 4.4: Webapp Use Case Diagram

### CHAPTER 5

**RESULTS** 

Our web-app is designed to predict both the sickness and heart disease of patients. The system is built using machine learning algorithms such as Random Forest Classifier and Logistic Regression.

Random Forest Classifier is used for predicting various types of diseases by inputting the symptoms of the patients. The system has an accuracy of 95 percent which makes it highly reliable.

For predicting heart disease, we have used Logistic Regression which is known for its effectiveness in training models with large datasets and getting a discrete value as an output. In comparison to other algorithms, the accuracy of our heart disease prediction model is better, especially when patients provide some essential medical details such as blood pressure, cholesterol, diabetes, LDL, etc.[6]

Our web-app is user-friendly and easy to use, doctors can input the medical details and symptoms of patients and the web-app will provide the predictions accordingly. Additionally, the it provides results in a timely manner which makes it highly useful for medical professionals and patients alike.

## CHAPTER 6 SYSTEM TESTING

We have used ITERATIVE MODEL. In this type of model possible key steps in the process start with a simple implementation of a subset of the system requirements and iteratively enhance the evolving versions until the full system is implemented.

#### 6.1 WHITE BOX TESTING

This type of testing ensures that:[9]

- All independent paths have been exercised at least once.
- All logical decisions have been exercised on their true and false sides.
- All loops are executed at their boundaries and within their operational bounds.
- All data structures have been exercised to assure their validity.

#### 6.2 CONDITIONAL TESTING

In this part of the testing each of the conditions were tested to both true and false aspects. And all the resulting paths were tested. So that each path that may be generate on particular condition is traced to uncover any possible errors.

#### 6.3 DATA FLOW TESTING

This type of testing selects the path of the program according to the location of definition and use of variables. This kind of testing was used only when some local variable was declared. The definition use chain method was used in this type of testing.

#### 6.4 TEST RESULTS

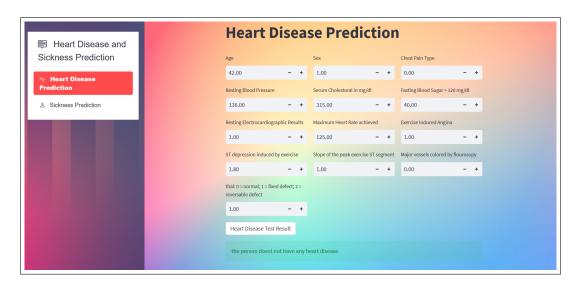


Figure 6.1: Patient With Healthy Heart

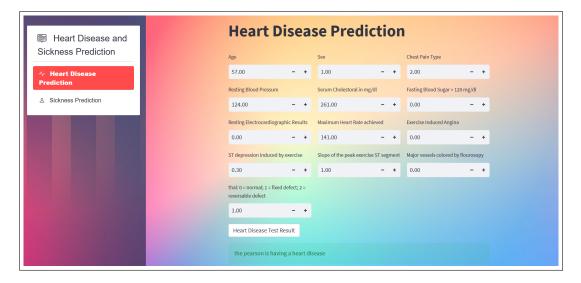


Figure 6.2: Patient With Unhealthy Heart



Figure 6.3: Patient's Sickness Prediction

### CHAPTER 7 CONCLUSION AND FUTURE SCOPE

The project aids in disease prediction at both personal and public levels. It provides information about the potential risk factors, thus reducing the need for unnecessary tests and associated costs. Moreover, the system can extract and analyze data from patients to reveal hidden patterns, which can be useful for future research and medical advancements.

During the training and testing phase of our project, we evaluated multiple machine learning models for predicting common diseases. The Random Forest Classifier model emerged as the most effective and accurate model with a precision of 95 percent. For the prediction and detection of heart disease, we utilized Logistic Regression which yielded an accuracy of 81 percent. There are various possibilities to further improve the accuracy of this model in the future.

The Heart disease detection model using Logistic Regression and the sickness prediction model using Random Forest Classifier showed superior results. These algorithms are not only more accurate but also cost-efficient and faster compared to the algorithms used by previous researchers. The Random Forest Classifier achieved a maximum accuracy of 95 percent, and Logistic Regression accuracy was 81 percent, which is either greater or almost equal to the accuracy's obtained from previous research. We can conclude that our accuracy has improved by using additional medical attributes from the dataset we used and can also improve more in future updates.

#### 7.1 CONCLUSION

We worked as a team, and gained some experience on how professional programmers work in the industry. There is always room for improvement, and this application we created can also be improved. This is especially because we had to create it within a limited time due to other projects, quizzes and Exams. The main goal of our system is to help detect any illness a person may be carrying early, allowing doctors to keep the treatment on track. This is especially helpful for people with heart problems because they can determine if they are at risk for heart disease. By using our system, individuals can benefit from early detection and treatment, which ultimately leads to better health outcomes.

#### 7.2 FUTURE SCOPE

There are several ways in which we can further enhance the accuracy of our system. For instance, we can explore the utilization of deep learning algorithms,[8] consider alternative techniques for attribute selection, and potentially expand the size of the dataset in the future to address issues related to overfitting and to improve overall performance.

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# ANNEXURE A LIST OF PUBLICATIONS AND RESEARCH PAPER

- 1. Paper Title: Heart Disease Detection and Patient's Sickness Prediction System
- 2. Name of the Conference/Journal where paper submitted : International Organization of Scientific Research
- 3. Paper accepted/rejected: Accepted
- 4. Review comments by reviewer: Quality of Manuscript is good.
- 5. Corrective actions if any: None

## ANNEXURE B PLAGIARISM REPORT

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