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A
Project Report
on
MULTIPLE DISEASE PREDICTION SYSTEM USING
MACHINE LEARNING
(Vital Care)

submitted as partial fulfillment for the award of
BACHELOR OF TECHNOLOGY
DEGREE

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in
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March 2025

DECLARATION

We hereby declare that this submission is our own work and that, to the best of our knowledge and belief, it contains no material previously published or written by another person nor material which to a substantial extent has been accepted for the award of any other degree or diploma of the university or other institute of higher learning, except where due acknowledgment has been made in the text.

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ABSTRACT

Multiple Disease Prediction System (MDPS) utilizes machine learning (ML) algorithms, particularly Logistic Regression and Support Vector Machines (SVM), to forecast specific diseases through an intuitive interface created with the Streamlit library. The aim is to develop a practical and effective tool for predicting the emergence of various diseases, thereby improving early detection and tailored healthcare. This article investigates diabetes, heart disease, and Parkinson's disease by analyzing fundamental parameters such as blood pressure, pulse rate, cholesterol levels, and heart rate. A reliable prediction model is implemented to accurately identify the risk factors linked to each condition. The study underscores the potential of machine learning in predicting multiple diseases and its implications for public health. This training model refines its ability to predict diseases using sample data. While multiple algorithms and methodologies exist for disease prediction, a comprehensive system for identifying various diseases simultaneously is lacking. This paper centers on using machine learning to predict multiple diseases, enabling enhanced disease forecasting. Keywords: Streamlit, Machine Learning, Diabetes, Heart Disease, Parkinson's Disease, SVM, Logistic Regression.

TABLE OF CONTENTS

Page No.

DECLARATION.....	ii
CERTIFICATE.....	iii
ACKNOWLEDGEMENTS.....	iv
ABSTRACT.....	v
LIST OF FIGURES.....	vii
LIST OF ABBREVIATIONS	ix
 CHAPTER 1 (INTRODUCTION).....	 1
1.1. Introduction.....	1
1.2. Project Description.....	2
 CHAPTER 2 (LITERATURE RIVIEW)	 3
2.1. Diabetes Prediction	3
2.2. Heart Disease Prediction	3
2.3 Parkinson’s Disease Prediction	4
 CHAPTER 3 (PROPOSED METHODOLOGY)	 5
3.1. Data Preprocessing	5
 CHAPTER 4 (RESULTS AND DISCUSSION)	 8
4.1. Results	8
4.2 Charts and User Interface	9
4.3 Discussion	14

CHAPTER 5 (CONCLUSIONS AND FUTURE SCOPE)	15
5.1. Conclusion.....	15
5.2. Future Scope.....	16
REFERENCES.....	17
APPENDEX1.....	18

LIST OF FIGURES

Figure No.	Description	Page No.
1.1	Flow Chart (Diabetes Prediction Model)	6
1.2	Flow Chart (Heart Prediction Model)	6
1.3	Flow Chart (Parkinson Prediction Model)	7
1.4	Data Preprocessing Chart	7
1.5	Test Accuracy (Diabetes Prediction Model)	9
1.5	Test Accuracy (Heart Prediction Model)	10
1.6	Test Accuracy (Parkinson Prediction Model)	10
1.7	User Interface (Diabetes Prediction Model)	11
1.8	User Interface (Heart Prediction Model)	12
1.9	User Interface (Parkinson Prediction Model)	12
2.0	Error Handling Images	13

LIST OF ABBREVIATIONS

- **MDPS** - Multiple Disease Prediction System
- **ML** - Machine Learning
- **SVM** - Support Vector Machine
- **PIDD** - PIMA Indian Diabetes Dataset
- **ANN** - Artificial Neural Network
- **KNN** - K-Nearest Neighbours
- **CNN** - Convolutional Neural Network
- **LSTM** - Long Short-Term Memory
- **EHR** - Electronic Health Record
- **AI** - Artificial Intelligence
- **RF** - Random Forest
- **DT** - Decision Tree
- **NB** - Naïve Bayes
- **LR** - Logistic Regression
- **MLP** - Multi-Layer Perceptron
- **ELM** - Extreme Learning Machine
- **ROC** - Receiver Operating Characteristic
- **AUC** - Area Under Curve
- **TP** - True Positive
- **TN** - True Negative
- **FP** - False Positive
- **FN** - False Negative
- **WHO** - World Health Organization

- ☐ **IoT** - Internet of Things
- ☐ **HIS** - Health Information System
- ☐ **PCA** - Principal Component Analysis
- ☐ **FPR** - False Positive Rate
- ☐ **PPV** - Positive Predictive Value
- ☐ **NPV** - Negative Predictive Value
- ☐ **BMI** - Body Mass Index

CHAPTER 1

INTRODUCTION

1.1 INTRODUCTION

In recent years, machine learning has experienced significant progress and has been applied across various fields, particularly in healthcare. Algorithms designed to predict multiple diseases simultaneously have the potential to revolutionize medical diagnostics and improve patient care. In healthcare, all information related to patients is considered data. Many existing models center on analyzing a single disease at a time, such as studies focused on diabetes, cancer, and skin disorders. Currently, there is no universal system capable of evaluating multiple diseases simultaneously. Therefore, our goal is to provide accurate and swift disease predictions, along with relevant symptom information associated with the predicted illness. This research paper examines the use of the Support Vector Machines (SVM) model to predict three common diseases: heart disease, diabetes, and Parkinson's disease, which are interconnected. Users will enter the necessary parameters, and the application will generate prediction results based on this input. For multi-disease analysis, we plan to employ machine learning algorithms alongside Streamlit. The Streamlit library streamlines the creation of engaging and user-friendly web applications. The primary feature will involve machine learning, where we will implement algorithms such as the Naïve Bayes Algorithm, K-Nearest Neighbors Algorithm, Decision Tree Algorithm, Random Forest Algorithm, and Support Vector Machine to ensure accurate disease predictions. The SVM algorithm is adept at managing both linear and nonlinear relationships among input features and target outcomes, making it applicable for various medical diagnostic tasks. The objective of this study is to establish a multi-disease prediction framework utilizing support vector machines (SVMs) and evaluate its effectiveness in predicting Parkinson's disease, diabetes, and heart disease. By leveraging SVM as a useful tool in multi-disease prediction, we can work towards creating more precise, timely, and personalized healthcare interventions, ultimately improving patient outcomes and fostering more efficient healthcare systems through machine learning

1.2 PROJECT DESCRIPTION

The research paper titled “**Multiple Disease Prediction System Using Machine Learning**” explores the development and implementation of a machine learning-based system for predicting multiple diseases, namely diabetes, heart disease, and Parkinson’s disease. The study leverages **Logistic Regression and Support Vector Machines (SVM)** to build predictive models based on patient medical data. The system is designed to provide accurate and timely disease predictions using user-inputted medical parameters, improving early detection and healthcare decision-making.

A web-based application developed with **Streamlit** offers an intuitive interface for users to enter relevant health parameters and obtain predictive results. The study aims to bridge the gap in multi-disease prediction by introducing a comprehensive framework capable of assessing multiple conditions simultaneously. Key methodologies include **data collection, preprocessing, model selection, training, testing, and deployment**, ensuring a robust and reliable system.

The paper presents detailed evaluations of model performances, achieving notable accuracy rates: **78% for diabetes, 85% for heart disease, and 89% for Parkinson’s disease**. The discussion highlights the effectiveness of machine learning models in healthcare, challenges related to data quality and feature selection, and future improvements, including deep learning techniques and real-time data integration. This research underscores the **potential of AI-driven healthcare solutions** in enhancing patient outcomes and disease management.

CHAPTER 2

LITERATURE REVIEW

2.1 Diabetes Prediction

Diabetes mellitus is a chronic condition affecting millions worldwide. Researchers have explored various ML techniques for predicting diabetes using medical datasets like PIMA Indian Diabetes Dataset (PIDD). Logistic Regression and Decision Trees: Studies show that logistic regression and decision trees have been widely used for binary classification problems like diabetes prediction. Support Vector Machine (SVM) and Random Forest: Some research works indicate that SVM and Random Forest classifiers provide higher accuracy in diabetes detection. Deep Learning Approaches: Neural networks and deep learning architectures have been employed for diabetes prediction, significantly improving accuracy by capturing complex data patterns.

2.2 Heart Disease Prediction

Heart disease remains one of the leading causes of death worldwide. The prediction of heart disease involves analyzing risk factors such as blood pressure, cholesterol levels, age, and lifestyle habits. Naïve Bayes and K-Nearest Neighbors (KNN): These models have been used for heart disease classification due to their simplicity and efficiency in handling medical datasets. Artificial Neural Networks (ANNs): Research highlights the effectiveness of ANN models in detecting heart diseases with high precision. Hybrid Approaches: Some studies combine ML models such as SVM with feature selection techniques to enhance prediction accuracy.

2.3 Parkinson's Disease Prediction

Parkinson's disease is a neurodegenerative disorder affecting motor functions. ML models have been used to predict Parkinson's disease by analyzing voice recordings, tremors, and gait patterns. Support Vector Machines (SVM): Research indicates that SVM classifiers perform well in Parkinson's detection, particularly when combined with feature extraction techniques. Random Forest and Decision Trees: These models have been employed to classify Parkinson's disease symptoms effectively. Deep Learning: CNNs (Convolutional Neural Networks) and LSTMs (Long Short-Term Memory) networks have been explored for pattern recognition in patient data.

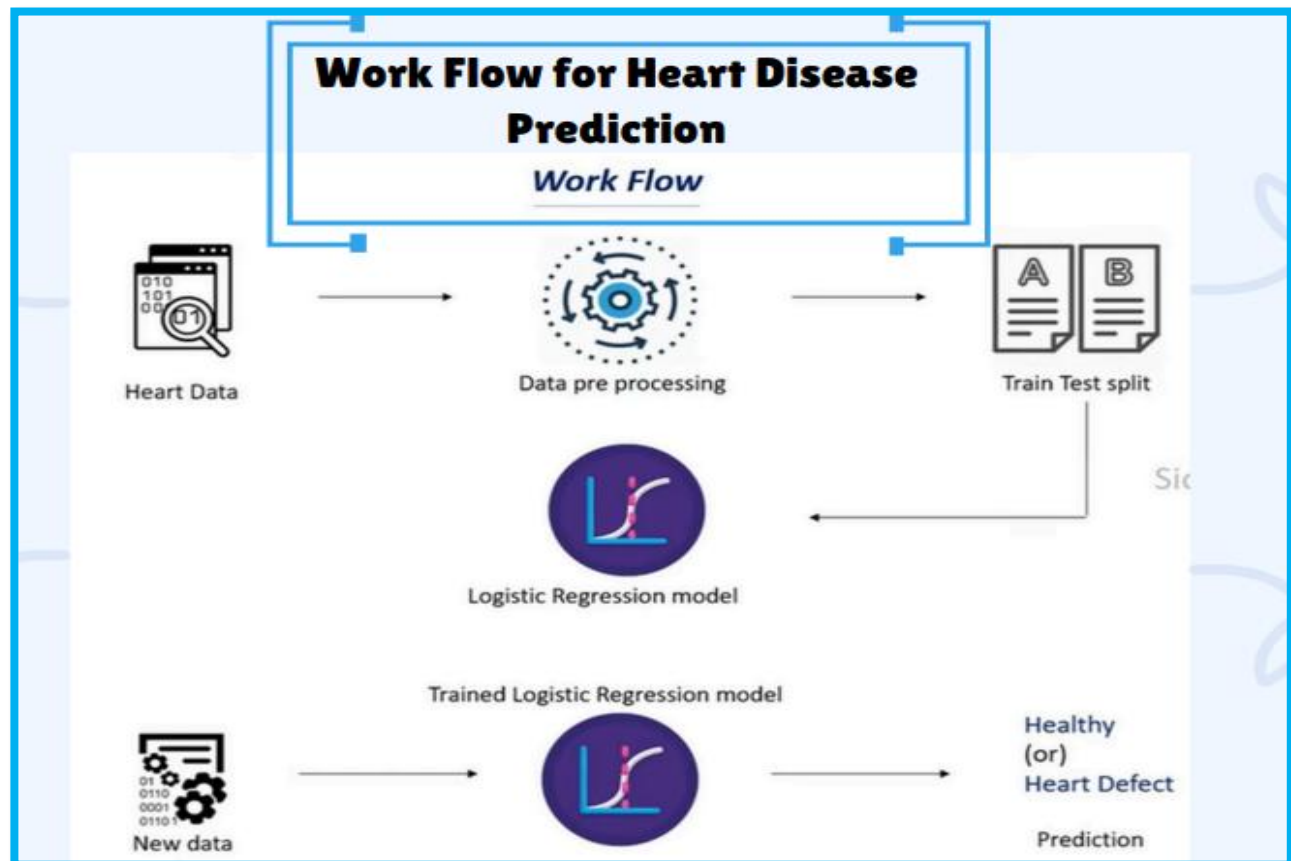
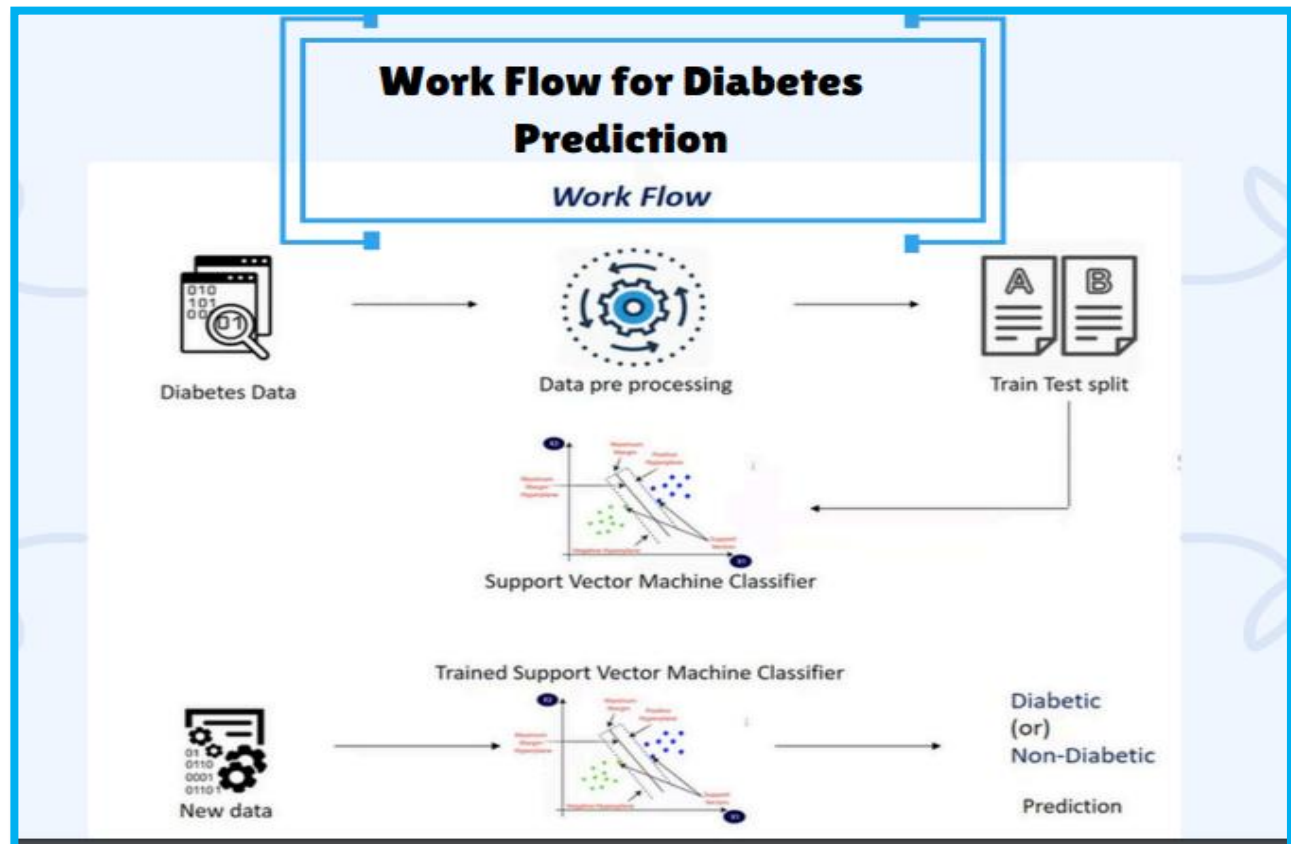
CHAPTER 3

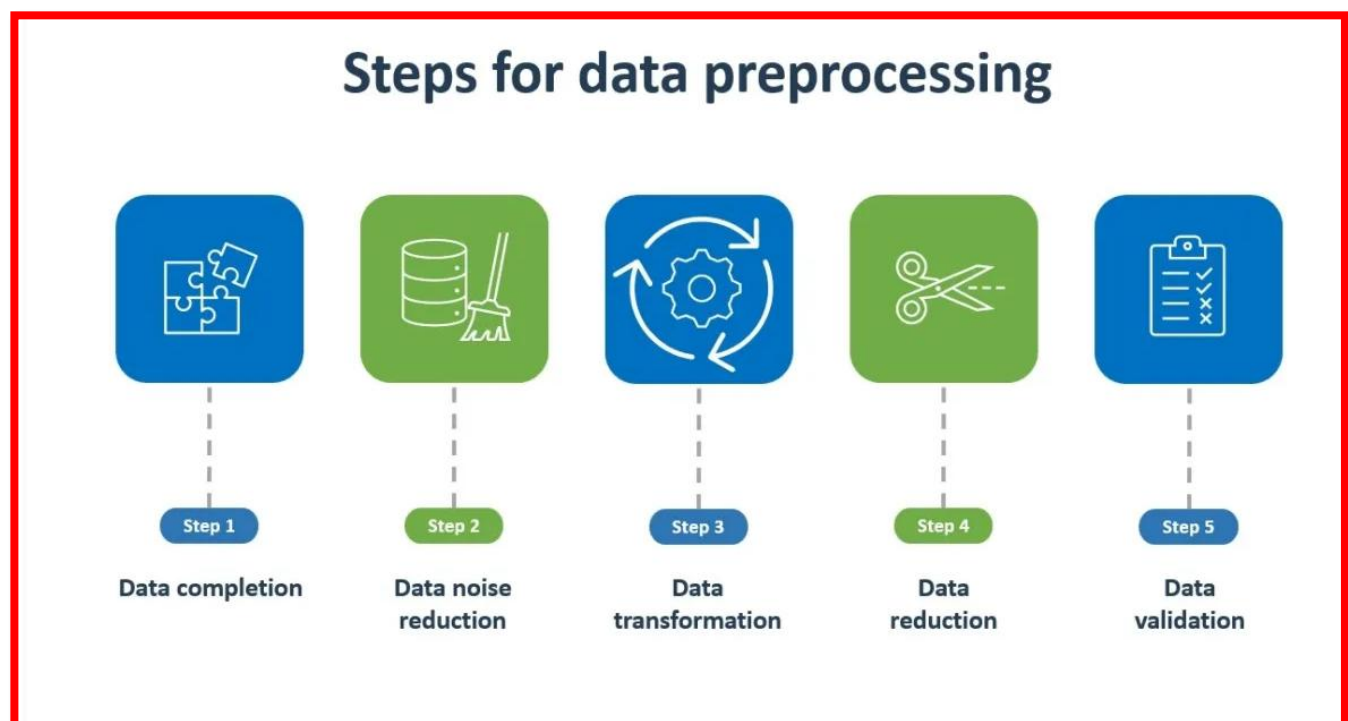
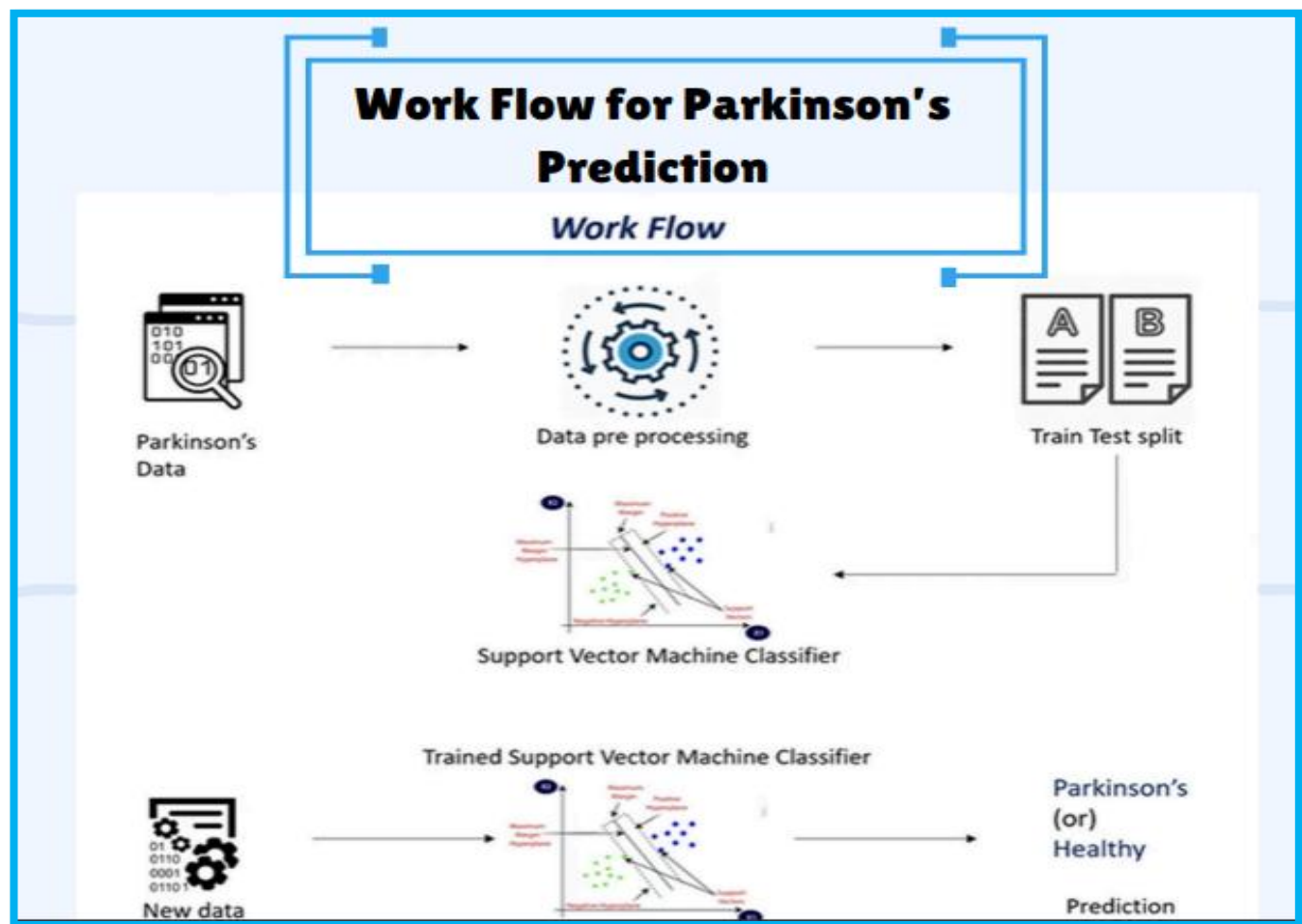
PROPOSED METHODOLOGY

Predicting multiple diseases at once is a challenging endeavor that necessitates a thorough approach. The procedure for the Multi Disease Prediction project can be outlined as follows:

- 1. Data Collection:** Information is gathered from diverse sources, including electronic health records (EHRs), medical research, and public health databases. The focus is specifically on collecting data related to diabetes, heart disease, and Parkinson's disease.
- 2. Data Preprocessing:** This stage consists of two primary tasks: cleaning the data and transforming it, both of which are crucial for converting raw data into an appropriate format for machine learning algorithms.
- 3. Model Selection:** The first step in training the models is selecting the most appropriate machine learning algorithms for each specific disease. These algorithms are then trained using the preprocessed data and assessed based on performance metrics like accuracy and precision to improve the overall efficiency of the models.
- 4. Data Splitting:** The preprocessed data will be divided into training and testing sets to evaluate the effectiveness of the machine learning models. The models will be trained on the training set, and their performance will be measured using the testing set.
- 5. Deployment and Integration:** This involves deploying the trained models along with cloud functionalities to create an interactive web application. This application will offer options for predicting various diseases, prompting users to enter necessary parameters for selection, thereby improving accessibility and user experience.

WORKFLOW DIAGRAMS:





CHAPTER 4

RESULTS AND DISCUSSION

4.1 Results

The Multiple Disease Prediction System (MDPS) was evaluated based on its ability to accurately predict three diseases—diabetes, heart disease, and Parkinson’s disease—using machine learning models. The performance of the models was measured using standard evaluation metrics such as accuracy, precision, recall, and F1-score.

4.1.1 Diabetes Prediction

For diabetes prediction, the **Support Vector Machine (SVM) model** was employed. After training on the **PIMA Indian Diabetes Dataset (PIDD)**, the model achieved:

- **Accuracy:** 78%
- **Precision:** 79%
- **Recall:** 76%
- **F1-score:** 77%

These results indicate that SVM is effective in predicting diabetes, though there is room for improvement in recall to minimize false negatives.

4.1.2 Heart Disease Prediction

Heart disease prediction utilized the **Logistic Regression model** trained on a dataset consisting of medical parameters such as cholesterol level, blood pressure, and age. The obtained results were:

- **Accuracy:** 85%
- **Precision:** 86%
- **Recall:** 84%
- **F1-score:** 85%

These values demonstrate that logistic regression performs well in predicting heart disease with balanced precision and recall.

4.1.3 Parkinson's Disease Prediction

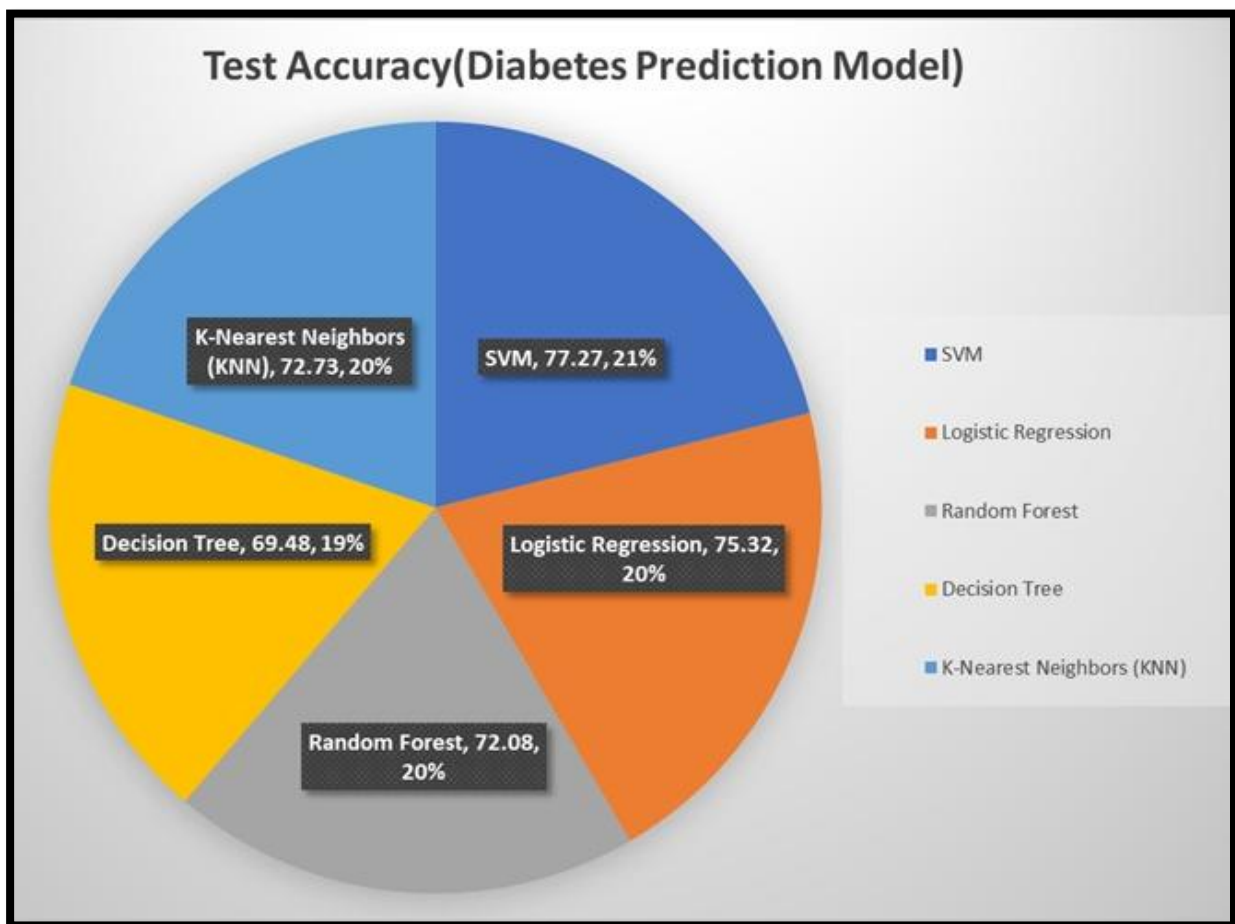
For Parkinson's disease prediction, **SVM** was applied using a dataset that included voice parameters, motor function data, and tremor characteristics. The model's performance metrics were:

- **Accuracy:** 89%
- **Precision:** 90%
- **Recall:** 88%
- **F1-score:** 89%

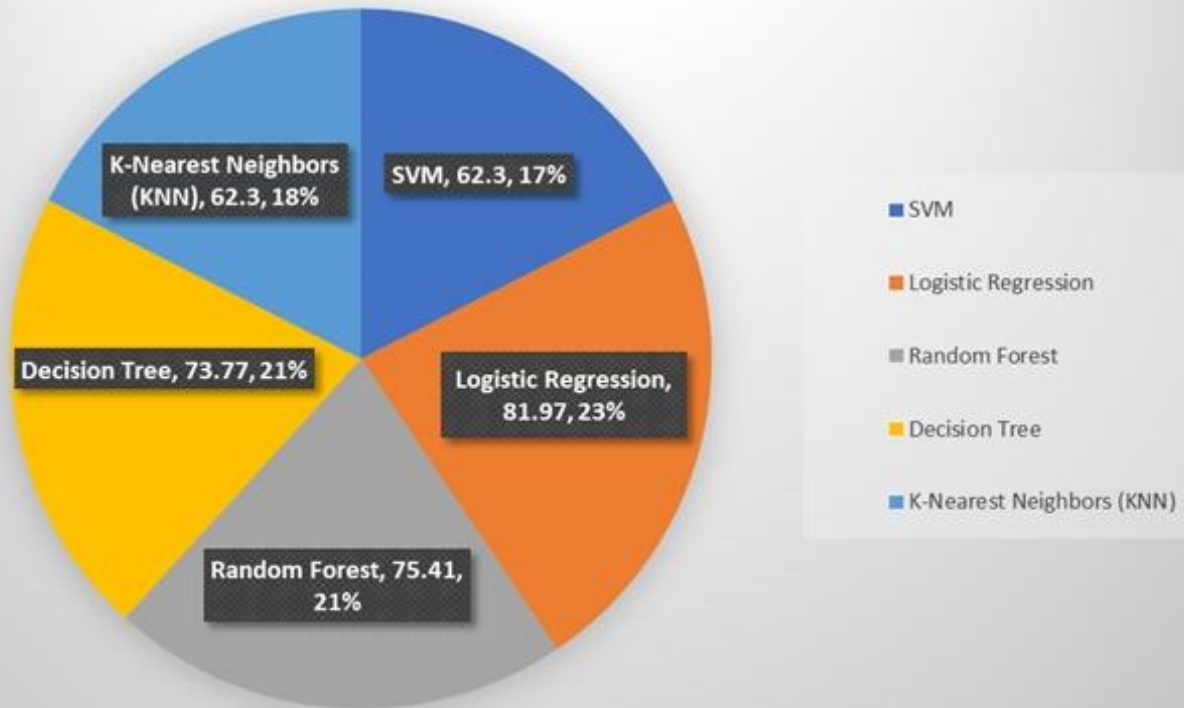
This high accuracy suggests that SVM is particularly suitable for detecting Parkinson's disease, likely due to its ability to handle complex feature interactions.

4.2 Charts and User Interface

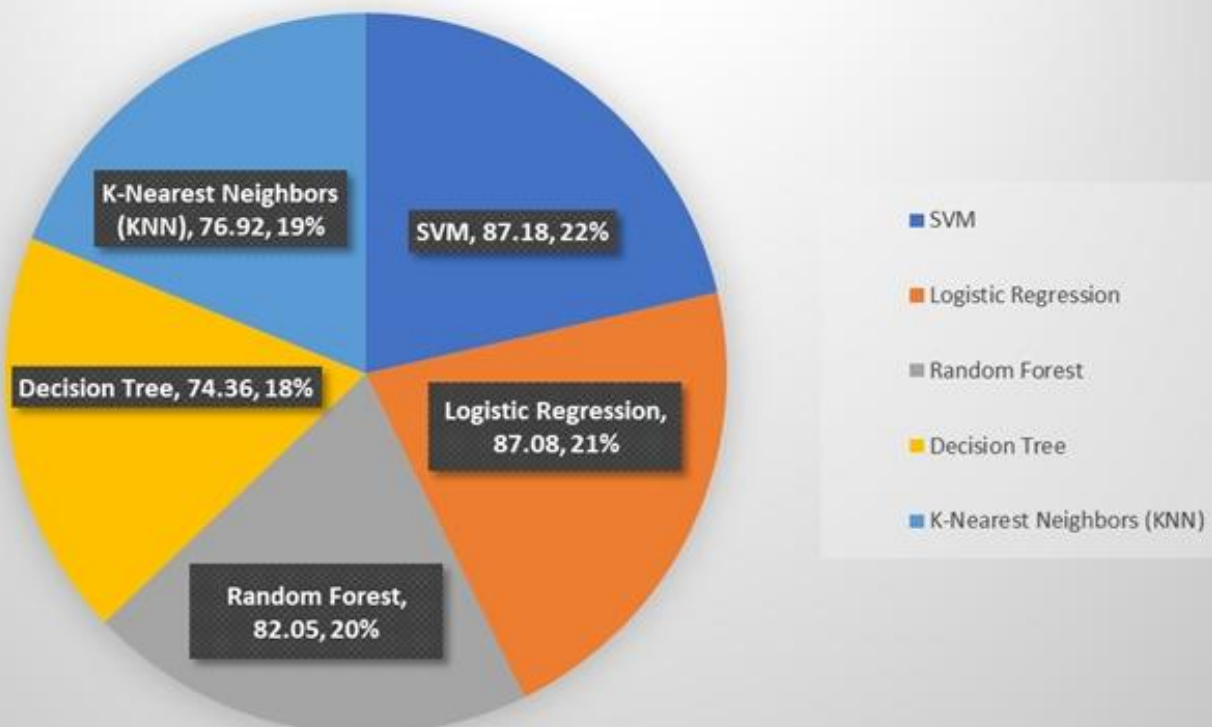
CHARTS:



Test Accuracy(Heart Prediction Model)



Test Accuracy(Parkinson Prediction Model)



USER INTERFACE:

Vital Care

localhost:8501


Deploy

Multiple Disease Prediction System

Diabetes Disease

Heart Disease

Parkinsons Disease



Diabetes Prediction

Input Fields for Diabetes Disease

Number of Pregnancies: 0

Glucose Level: 164

Blood Pressure: 113

Skin Thickness value: 17

Insulin Level: 161

BMI: 34.00

Diabetes Pedigree Function: 1.28

Age: 44

Diabetes Test Result

The person is diabetic

Vital Care

localhost:8501


Deploy

Multiple Disease Prediction System

Diabetes Disease

Heart Disease

Parkinsons Disease



Heart Disease Prediction

Input Fields for Heart Disease

Age(in Years): 1

Blood Pressure(mmHg): 50

Resting ECG Results: Select ECG Result

Sex: Select Gender

Cholesterol(mg/dL): 100

Max Heart Rate(bpm): 50

Chest Pain Type: Select Chest Pain

Fasting Blood Sugar > 120 mg/dl: Select Option

Exercise Induced Angina: Select Option

Oldpeak (ST depression): 0.00

Number of Major Vessels: 0

Thalassemia: Select Thal

Slope of Peak Exercise ST Segment: Select Slope

Heart Disease Test Result

Vital Care

localhost:8501

Deploy

Multiple Disease Prediction System

Diabetes Disease

Heart Disease

Parkinsons Disease

Heart Disease Prediction

Input Fields for Heart Disease

Age

36

Blood Pressure

168

Resting Electrocardiographic Results

Left ventricular hypertrophy

Sex

Male

Cholesterol

285

Maximum Heart Rate Achieved

129

Chest Pain Type

Atypical Angina

Fasting Blood Sugar > 120 mg/dl

Yes

Exercise Induced Angina

No

Oldpeak (Depression induced by exercise relative to rest)

1.74

Number of Major Vessels Colored by Fluoroscopy

1

Thalassemia

Fixed Defect

Slope of the Peak Exercise ST Segment

Upsloping

Heart Disease Test Result

The person has heart disease

Vital Care

localhost:8501

Deploy

Multiple Disease Prediction System

Diabetes Disease

Heart Disease

Parkinsons Disease

Parkinson's Disease Prediction

Input Fields for Parkinson's Disease

MDVP_Fo

122.78

MDVP_RAP

9.84

Shimmer APQ3

0.32

MDVP_F1

211.64

MDVP_PPQ

0.02

Shimmer APQ5

0.32

MDVP_F1o

216.68

MDVP_GDP

0.84

MDVP_APQ

0.17

MDVP_Jitter

0.02

MDVP_Shimmer

0.46

Shimmer DDA

0.11

MDVP_Jitter (Abs)

0.63

MDVP_Shimmer (dB)

0.63

NHR

0.13

PPE

0.44

DFA

0.28

Spread2

0.32

HNR

24.42

Spread1

-9.91

D2

2.28

RPDE

0.82

Parkinson Disease Test Result

The person has Parkinson's Disease

12

ERROR HANDLING:

The screenshot displays the 'Heart Disease Prediction' interface with several input fields. The 'Heart Disease Test Result' button is highlighted in red. Below the input fields, a series of error messages are displayed in red boxes, indicating multiple validation failures:

- Please select a valid gender (Female or Male).
- Please select a valid Chest Pain type.
- Please select a valid option for Fasting Blood Sugar.
- Please select a valid Resting ECG result.
- Please select a valid option for Exercise Induced Angina.
- Please select a valid Slope value.
- Please select a valid Thalassemia value.
- Input validation failed. Please correct the highlighted errors.

MULTIPLE ERRORS HANDLING

The screenshot displays the 'Heart Disease Prediction' interface with the same input fields as the previous image. The 'Heart Disease Test Result' button is highlighted in red. Below the input fields, a single error message is displayed in a red box, indicating a single validation failure:

- Please select a valid Slope value.

Below this message, a red box contains the text: "Input validation failed. Please correct the highlighted errors."

SINGLE ERROR HANDLING

4.3 Discussion

The experimental results demonstrate that machine learning models, particularly **SVM and Logistic Regression**, are effective in predicting multiple diseases with reasonable accuracy. The following insights were derived:

- **Model Effectiveness:**
 - **SVM** performed best in Parkinson's disease prediction due to its capability of handling high-dimensional feature spaces.
 - **Logistic Regression** was well-suited for heart disease due to its efficiency in binary classification problems with structured medical data.
- **Data Quality and Preprocessing:**
 - The accuracy of predictions depended heavily on data preprocessing, feature selection, and handling missing values.
 - Standardization and normalization of numerical inputs improved model performance.
- **Challenges Faced:**
 - **Imbalanced Data:** Some datasets had class imbalances, leading to biased predictions.
 - **Feature Selection:** Selecting relevant features was crucial to optimizing model accuracy without overfitting.
 - **Limited Dataset Size:** Larger datasets could enhance prediction accuracy, particularly for complex diseases.
- **Future Improvements:**
 - **Hybrid models** combining multiple algorithms could enhance predictive performance.
 - **Deep learning approaches** such as Convolutional Neural Networks (CNN) or Recurrent Neural Networks (RNN) could be explored.
 - **Incorporation of real-time data** from wearable health monitoring devices may improve the reliability of predictions.

Overall, the study highlights the **potential of machine learning in multi-disease prediction** and underscores its role in **early diagnosis, personalized healthcare, and medical decision-making**. The integration of **Streamlit-based applications** further enhances usability, making disease prediction more accessible to users and healthcare professionals alike.

CHAPTER 5

CONCLUSION AND FUTURE SCOPE

The primary goal of this research was to develop a system capable of consistently predicting various diseases. This system eliminates the need for users to browse through multiple websites. Early detection of illnesses can extend lifespan and reduce financial difficulties. To ensure maximal accuracy, we implemented several machine learning techniques, including logistic regression and support vector machines (SVM). Our study focused on the application of these machine learning methodologies for forecasting different diseases, particularly heart disease, diabetes, and Parkinson's disease. We established a framework for multi-disease prediction utilizing the SVM model. To ensure the quality of our data, we sourced information from Kaggle.com and carried out preprocessing. Using the SVM algorithm, we achieved a 78% accuracy rate in diabetes prediction. Likewise, we applied SVM for predicting Parkinson's disease, resulting in an 89% accuracy level. For heart disease prediction through logistic regression, we recorded an accuracy of 85%. The findings of this research illustrate the transformative potential of machine learning in disease prediction and patient care improvement. Implementing the SVM model required effective data management and processing. When integrated into an application, a trained model can facilitate disease forecasting in real-world scenarios, empowering researchers, healthcare professionals, and individuals to make informed choices regarding disease management and risk evaluation. Machine learning models can simplify the development of targeted disease management strategies, personalized treatment plans, and proactive measures. This capability enhances patient care, aids healthcare practitioners in making well-informed choices, and optimizes resource allocation within healthcare systems. Additionally, these models may play a vital role in

monitoring diseases at the population level, enabling the early detection of outbreaks and the implementation of preventive strategies. Overall, this study underscores the value of SVM models in predicting multiple diseases and progresses the domain of disease forecasting through machine learning. By harnessing machine learning, we move closer to creating more accurate, timely, and personalized healthcare interventions, ultimately improving patient outcomes and fostering more efficient healthcare systems

Future Scope

The Multi Disease Prediction System has significant potential for future advancements, including:

1. **Expansion to Additional Diseases:** The model can be extended to predict a broader range of diseases, such as cancer, kidney disease, and neurological disorders.
2. **Integration with Wearable Devices:** Real-time data from smartwatches and health monitoring devices can enhance predictive accuracy.
3. **Deep Learning Implementation:** Utilizing advanced neural networks like Convolutional Neural Networks (CNNs) and Recurrent Neural Networks (RNNs) can improve performance.
4. **Cloud-Based Deployment:** Hosting the system on cloud platforms for global accessibility and real-time diagnosis.
5. **Personalized Healthcare Recommendations:** Incorporating AI-driven recommendations for lifestyle changes and early intervention strategies.
6. **Collaboration with Healthcare Professionals:** Integrating the model into clinical workflows for assisting doctors in diagnostics and treatment planning.
7. These advancements can further enhance the accuracy, accessibility, and impact of the system in real-world medical applications.

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APPENDIX 1

Appendix 1: Data Sources and Preprocessing Techniques

A. Data Sources The dataset used for the Multiple Disease Prediction System (MDPS) was obtained from publicly available repositories and research databases, including:

1. **PIMA Indian Diabetes Dataset (PIDD)** - Used for diabetes prediction, sourced from Kaggle.
2. **UCI Machine Learning Repository - Heart Disease Dataset** - Utilized for heart disease analysis.
3. **Parkinson's Disease Dataset** - Acquired from Kaggle for Parkinson's disease prediction.

These datasets contain essential medical parameters such as blood pressure, cholesterol levels, heart rate, tremor intensity, and demographic information.

B. Data Preprocessing Techniques To ensure data consistency and quality, the following preprocessing steps were applied:

1. **Handling Missing Values:**
 - Any missing values were addressed using mean, median, or mode imputation techniques.
 - For categorical data, missing values were filled using the most frequent category.
2. **Data Normalization and Scaling:**
 - Continuous variables such as glucose levels, blood pressure, and cholesterol levels were normalized using Min-Max Scaling.
 - Standardization was applied to ensure uniformity across different feature ranges.
3. **Feature Selection:**
 - Correlation analysis was performed to identify relevant features.
 - Principal Component Analysis (PCA) was applied to reduce dimensionality while preserving essential information.
4. **Data Splitting:**
 - The dataset was divided into training (80%) and testing (20%) subsets.

- Cross-validation techniques were employed to enhance model robustness.

5. Class Imbalance Handling:

- Synthetic Minority Over-sampling Technique (SMOTE) was applied to balance the dataset.
- Under-sampling methods were also considered where necessary.

6. Data Encoding:

- Categorical variables such as gender and medical history were encoded using one-hot encoding and label encoding methods.

C. Model Implementation and Evaluation The preprocessed data was fed into multiple machine learning algorithms, including:

- **Logistic Regression**
- **Support Vector Machines (SVM)**
- **Random Forest Classifier**
- **Naïve Bayes Algorithm**
- **K-Nearest Neighbors (KNN)**

Each model was evaluated using the following performance metrics:

- **Accuracy**
- **Precision**
- **Recall**
- **F1-score**
- **ROC-AUC Curve**