

## Phase-1 Submission

**Student Name:** Jeevikasri R

**Register Number:** 410723104029

**Institution:** Dhanalakshmi college of engineering

**Department:** Computer Science and Engineering

**Date of Submission:** 30-04-2025

---

### 1.Problem Statement

#### AI – POWERED DISEASE PREDICTION

Transforming healthcare with AI-powered disease prediction based on patient data.

### 2.Objectives of the Project

#### Project Objective:

To develop an AI-powered system that predicts the risk of diseases using patient data, enabling early diagnosis, personalized care, and better clinical outcomes.

#### Key Outcomes:

- Train machine learning models to predict diseases based on patient demographics, medical history, and lifestyle factors.
- Classify patients into risk categories for targeted intervention.
- Identify key predictors influencing disease risk.
- Evaluate model accuracy using metrics like precision, recall, and AUC.
- Provide a decision-support tool for clinicians with risk insights and recommendations.

### **3.Scope of the Project**

#### **Features to Analyse/Build:**

##### **1. Demographics:**

- Age, gender, ethnicity, location

##### **2. Medical History:**

- Past diagnoses, family history of diseases, previous hospitalizations

##### **3. Vital Signs & Clinical Metrics:**

- Blood pressure, heart rate, BMI, cholesterol, glucose levels

##### **4. Lab Test Results:**

- Blood work, urinalysis, liver/kidney function tests

##### **5. Lifestyle Factors:**

- Smoking status, alcohol consumption, physical activity, diet

##### **6. Medication & Treatment History:**

- Current/past prescriptions, treatment adherence

##### **7. Genetic or Genomic Data (if available)**

##### **8. Time-Series Data (for longitudinal analysis):**

- Trends in vitals or lab values over time

## Limitations & Constraints:

### 1. Data Constraints:

- Use of publicly available or anonymized datasets (e.g., MIMIC-III, UCI Health datasets)
- Missing or imbalanced data may affect model performance

### 2. Model Constraints:

- Limited to interpretable models if required by clinical partners (e.g., logistic regression, decision trees)
- Avoid black-box models unless explainability techniques (e.g., SHAP, LIME) are applied

### 3. Deployment Constraints:

- Prototype may not be deployed in real clinical settings without regulatory approval (e.g., FDA clearance)
- Compliance with data privacy laws (e.g., HIPAA, GDPR)

### 4. Tool Constraints:

- Development limited to Python-based frameworks (e.g., scikit-learn, TensorFlow, PyTorch)
- Visualization using tools like Streamlit, Dash, or Power BI

## 4.Data Sources

### Dataset Description:

- **Dataset Name :** Healthcare Dataset
- **Source :** Kaggle ([dataset](#))
- **Accessibility :** Public
- **Type :** Static

## 5.High-Level Methodology

- **Data Collection** – The dataset will be obtained through direct download from publicly available source **Kaggle** for disease diagnosis.
- **Data Cleaning** – Identify potential issues such as **missing values, duplicates, or inconsistent formats**.
- **Exploratory Data Analysis (EDA)** –  
**Predictive Modeling & Risk Analysis** •  
**Techniques:**
  1. **Logistic regression, decision trees, or random forests** – for predicting disease risk.
  2. **Survival analysis (e.g., Kaplan-Meier curves)** – for analysing time to event (e.g., time until readmission).
- **Visualizations:**
  1. **ROC curves / AUC plots** – to evaluate model performance.
  2. **Survival curves** – to compare patient outcomes by treatment groups.

- **Model Building :**

**Supervised Learning**

1. **Logistic Regression** – Simple, interpretable, great for binary outcomes.
2. **Random Forest** – Handles non-linear data, robust to noise.
3. **XGBoost/LightGBM** – High accuracy, handles complex patterns well.
4. **SVM** – Good for high-dimensional classification.
5. **Neural Networks** – Flexible, good for large and complex datasets.

**Unsupervised Learning**

6. **K-Means** – Fast and effective for patient clustering.
7. **Hierarchical Clustering** – Useful for exploring group hierarchies.

**Deep Learning (for Images/Text)**

**CNNs** – Best for medical image analysis.

**Transformers (e.g., BERT)** – Excellent for clinical text mining.

- **Model Evaluation**

**Metrics**

1. **Classification:** Accuracy, Precision, Recall, F1 Score, ROC-AUC
2. **Regression:** MAE, RMSE,  $R^2$
3. **Survival:** C-index, Log-rank Test
4. **Clustering:** Silhouette Score, Clinical relevance

**Validation Strategies**

5. **Train-Test Split** – Simple, quick check
6. **k-Fold CV / Stratified k-Fold** – Robust, keeps class balance
7. **Time Series Split** – For time-dependent data
8. **Bootstrapping** – Good for uncertainty estimation

- **Visualization & Interpretation**

**Visualization & Interpretation**

1. **Charts:** Line, bar, scatter, boxplots, heatmaps
2. **Dashboards:** Interactive summaries (e.g., Power BI, Tableau)
3. **Model Explainers:** SHAP, LIME, feature importance
4. **Reports:** Clear visuals + insights for stakeholders

## 6.Tools and Technologies

- **Programming Language** – The main language we use is Python.
- **Notebook/IDE** –The platform we use is Google Colab.
- **Libraries** – The libraries we use is pandas, NumPy, seaborn, matplotlib.

## 7.Team Members and Roles

S.No	NAME	ROLE
1	Agnes Selestina S	Data Collection, Data Cleaning
2	Christina Ryka S	Visualization & Interpretation
3	Jeevikasri R	Exploratory Data Analysis (EDA), Feature Engineering
4	Keerthana R	Model Building, Model Evaluation