Module Code & Module Title

CU6051NI Artificial Intelligence

25% Individual Coursework

Submission: Milestone 1

Academic Semester: Autumn Semester 2025

Credit: 15 credit semester long module

Student Name: Unison Raj Tuladhar

London Met ID: 23050414

College ID: NP01CP4A230141

Assignment Due Date: 01/07/2026

Assignment Submission Date: 07/07/2026

Submitted To: Er. Roshan Shrestha

GitHub Link	https://github.com/UnisonTuladhar/AI_Project.git

I confirm that I understand my coursework needs to be submitted online via MST Classroom under the relevant module page before the deadline for my assignment to be accepted and marked. I am fully aware that late submissions will be treated as non-submission and a mark of zero will be awarded.

Table of Contents

Table of Figures

1. Introduction

1.1 Introduction to Artificial Intelligence and Machine Learning

Artificial Intelligence (AI) is a technology that enables computers and machines to perform cognitive functions we associate with human minds, such as perceiving, reasoning, learning, interacting and problem solving. We all have interacted with AI technology even if you don’t realize it. Voice assistants like Siri, Alexa or some customer chat bots are some examples of AI (McKinsey, 2024). AI encompasses the fields of computer and data science focused on building machines with human intelligence to perform various tasks. Instead of relying instructions from a individual AI systems can learn from the data which lets them handle the hard complex problems by their own and improve their accuracy overtime (Tech, 2025).

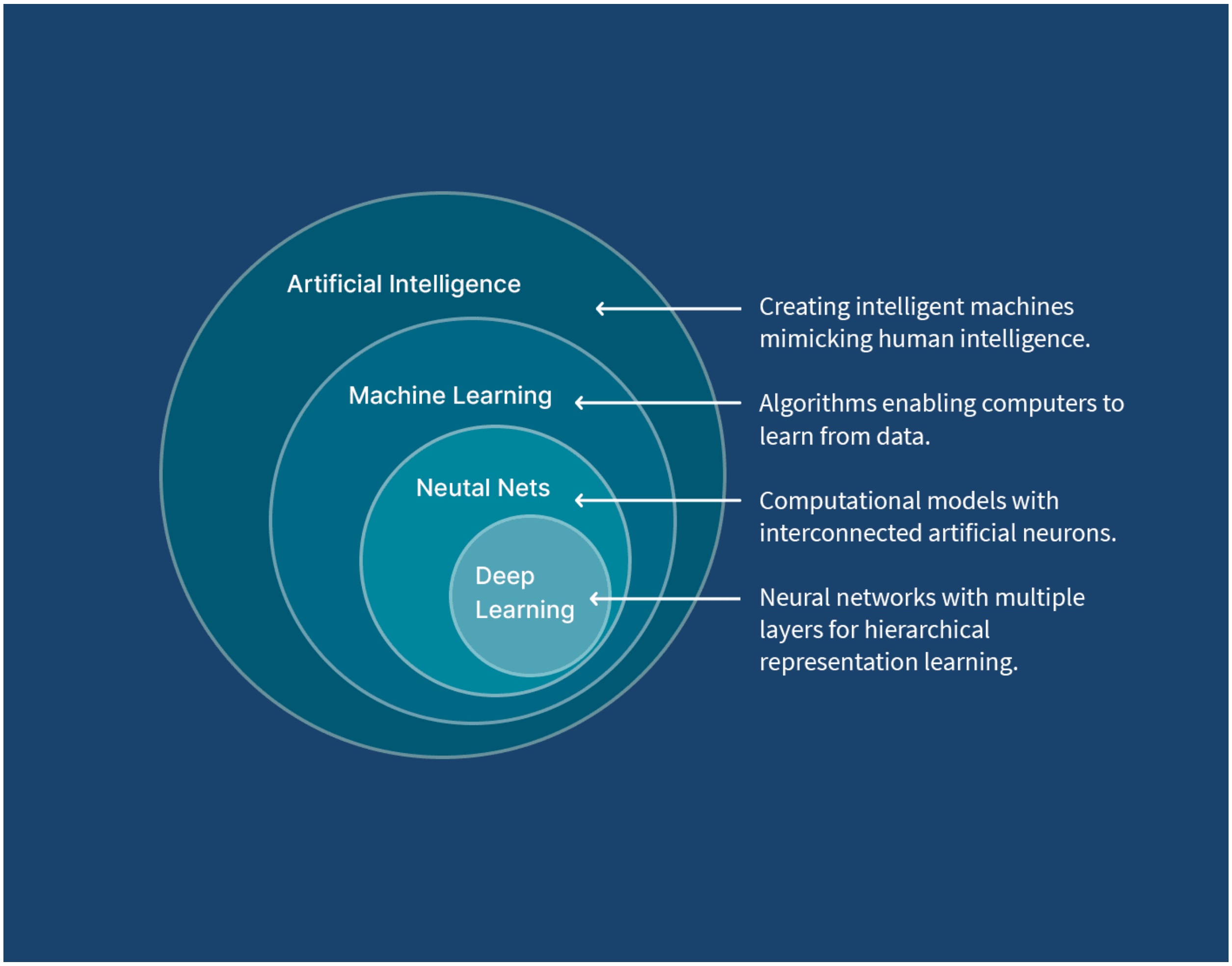


Figure 1: Artificial Intelligence and Machine Learning.

Machine Learning (ML) is a form of AI that enables machines to learn data patterns automatically and improve their performance over time without being explicitly programmed. It does this by optimizing model parameters through calculations. The learning algorithm then continuously updates the parameter values allowing the model to make predictions or decisions (ISO, 2022). Machine Learning is categorized into three types:

- Supervised Learning
- Unsupervised Learning
- Reinforcement Learning

Supervised Learning (classification) has been chosen for this project. This model uses machine learning classification to analyse the healthcare data from the diabetes dataset. The diabetes dataset contains data such as age, gender, BMI, Cholesterol, Triglycerides, HDL, LDL, Creatinine and Blood Urea Nitrogen. These data will be pre-processed and converted into suitable numerical format that can be learned from the ML algorithms. For this model three best classification algorithms were selected (Logistic Regression, Random Forest Classifier and SVM).

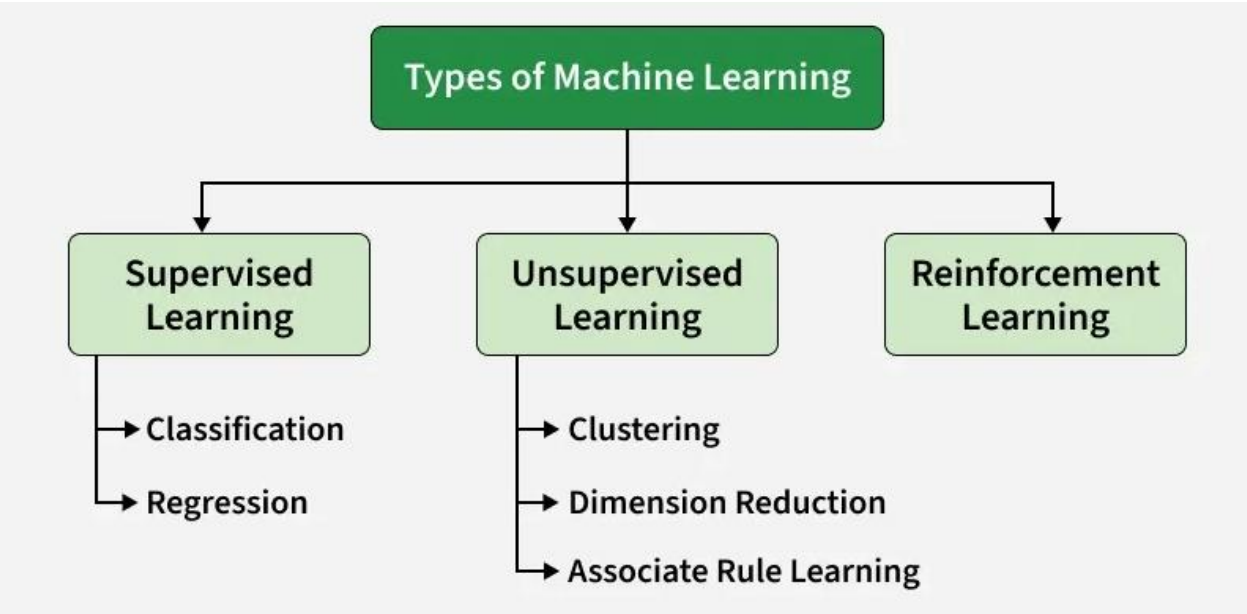


Figure 2: Types of Machine Learning (ML).

1.2 Problem Domain

The chosen problem domain for this project lies under healthcare sector. The topic of my project is Diabetes Prediction using supervised learning classification algorithm. Diabetes is a chronic disease that occurs when the pancreas does not produce enough insulin or the body cannot effectively use the insulin it produces. Diabetes is a disease that has affected millions of people worldwide and if not treated well it leads to serious damage to the body overtime. Around 14% of the adults aged 18 and older were having diabetes and more than 59% of adults 30 or older are living with diabetes according to the research done in 2022. In 2021 diabetes was the cause of death of around 1.6 million and 47% of them occurred before the age of 70 (WHO, 2024).

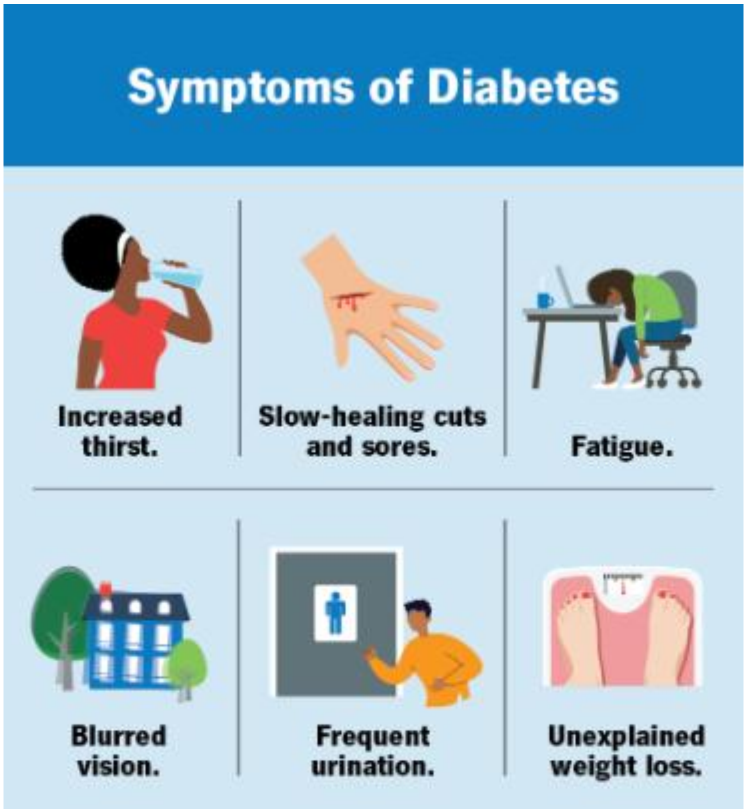


Figure 3: Symptoms of Diabetes.

2. Background

2.1 Research on Diabetes Prediction

Diabetes is a common disease that affected around 537 million adults around the world and this number might go up to 643 million by 2030 and 783 million in 2025 people of all ages. There are many types of diabetes but Type 2 diabetes are the most common ones, Type 1 diabetes is mostly diagnosed to children and teenagers and Gestational diabetes are the once developed during pregnancy (Clinic, 2023). Many individuals tend to delay their medical tests due to lack of time, laziness, lack of awareness even the symptoms are visible. As a result, diabetes reaches at later stages and the later stages conditions becomes more severe.

Since many individuals don't even realize they have diabetes until later stages which could be harmful so early prediction of diabetes using medical data can help individuals to know about the disease earlier. The main goal of this project is to build a classification model using algorithms like logistic regression, random forest classifier, support vector machine (SVM) to predict whether an individual is likely to have diabetes based on the persons attributes such as age, gender, body mass index, cholesterol, creatine and so on.

2.2 Existing Research Work

Several studies have been done and explored in the medical field related to diabetes as it is one of the most common disease. Here below are the three previous researches done in the medical field related to diabetes.

- **Research 1 (Diabetes Prediction Using Machine Learning Classification)**

This research is based on the supervised learning using classification algorithms to predict diabetes using medical data. Algorithms like Logistic regression, KNN, SVM, Decision Tree algorithms are using in this research. As the research the prevalence of diabetes for all the age group was estimated to be 171 million (2.8%) in 2000 A.D and increases up to 366 million (4.4%) in 2030 A.D. So, the fact that these many individuals are affected from the disease but still many of them don't even realize diabetes in them until later stages which could be harmful. The final result displayed the SVM and the Decision Tree algorithm performed better than the other algorithms with an accuracy rate of 80% which is considered to be a great amount. The study demonstrates that ML based classification can be a great help for the healthcare domain in early diabetes prediction (B.V., 2017).

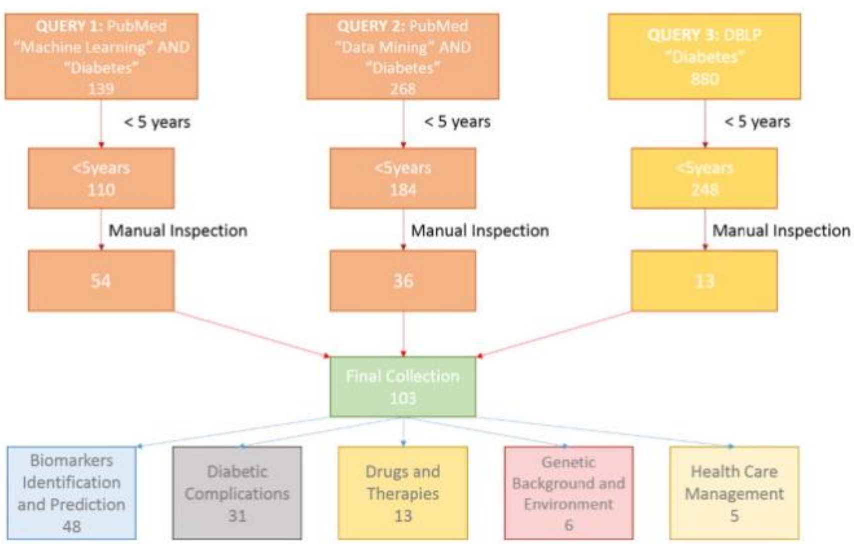


Figure 4: Workflow diagram of research 1.

- **Research 2 (National Library of Medicine Diabetes Prediction)**

This research is based on the supervised learning using classification algorithms to predict diabetes using medical data. According to the IDF (International Diabetes Federation) statistics in the research about 537 million people had diabetes all around the world in 2021. The most suffered country according to the statistics were Bangladesh with around 7.10 million people affected with this disease. The open source Pima Indian and a private data set of female Bangladeshi patients' data were used for this research. This research used various algorithms with accuracy around 60% to 70% but the highest accuracy was obtained by the XG boost classifier with an accuracy rate of 81% and an AUC and F1 score of 0.84 and 0.81 respectively. The study concludes that ML classification models can assist significantly the healthcare professionals by providing a fast and reliable data. This research also highlights the challenges faced during the research like the data imbalance and the model interpretability (Khan, 2022).

Performance metrics of various classifiers using adasyn in the merged dataset

Classifier	Precision	Recall	F1 Score	Accuracy	Auc
Logistic regression	0.76	0.75	0.75	75%	0.84
KNN	0.76	0.73	0.73	73%	0.82
Random forest	0.76	0.76	0.76	76%	0.84
Decision tree	0.81	0.72	0.72	72%	0.78
Bagging	0.80	0.79	0.79	79%	0.84
AdaBoost	0.75	0.76	0.76	76%	0.84
XGBoost	**0.81**	**0.81**	**0.81**	**81%**	**0.84**
Voting	0.77	0.77	0.77	77%	0.84
SVM	0.78	0.78	0.77	78%	0.83

Figure 5: Performance metrics of research 2.

- **Research 3 (Machine Learning Based Approaches for Diabetes Prediction)**

According to the MDPI research as per reports 8.5% of the world population in 2014 as suffering from diabetes where type-2 were for 95% of them. The diabetes patients were increasing day by day. till the date 2019 1.5 million deaths were caused by diabetes. Although diabetes is affected to both men and women but according to the research women has a higher chance of developing diabetes. The Pima Indian Diabetes dataset is used in this research. Many algorithms were used in this project like Decision Tree with an accuracy rate of 72% similarly Naive Bayes with an accuracy rate of 77% but the highest accuracy rate among all the algorithms was Logistic Regression with an accuracy rate of 80%. At last in the research it says early detection is crucial for preventing or delaying the development of serious diabetes cases. This research collected the dataset and using the help of the dataset created heatmap analytical tools, ML based prediction model which would be helpful for the health sectors to predict early diabetes (Ahmed, 2024).

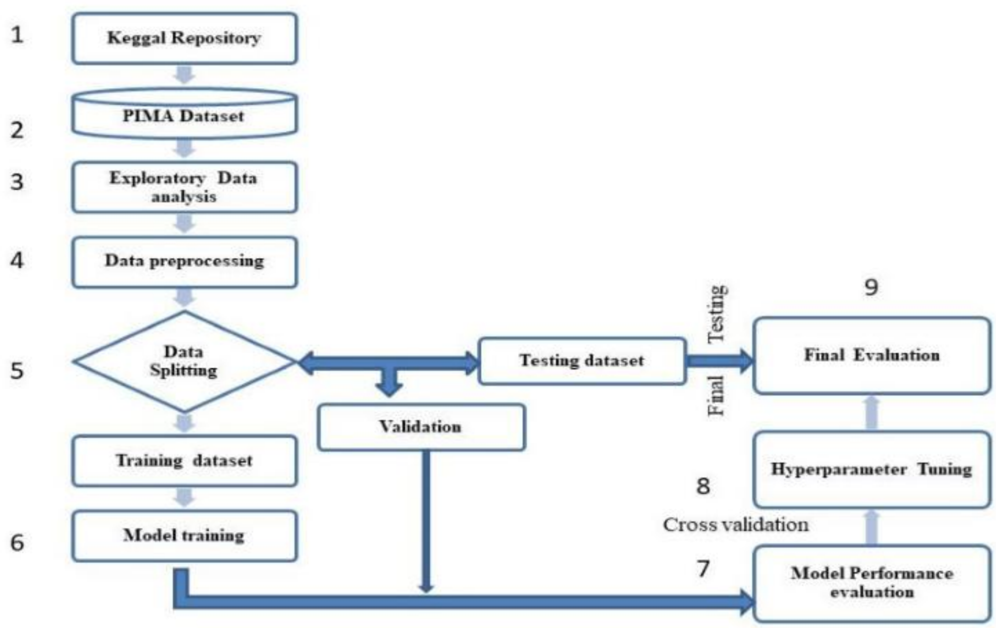


Figure 6: Flowchart of the methodology used for the ML model development.

2.3 Framework and tools used

- Python
- Pandas, NumPy, sklearn
- Matplotlib/ seaborn
- Jupiter Notebook
- GitHub

2.4 Advantages, Drawbacks and Issues

Advantages:

- Diabetes ML classification model can help detect early diabetes which might help preventing or delaying the development of serious diabetes cases.
- Algorithms selected in this project like Logistic Regression, Random Forest Classifier, Support Vector Machine (SVM) provide reliable performance in predicting diabetes.
- The selected dataset contains relevant clean medical features making the predictions accurate and applicable to the real-world health care scenarios.

Drawbacks:

- The performance of the ML learning models depends on the data quality and proper pre-processing.
- Complex models may lack some interpretability making this a bit difficult for healthcare professionals.
- Training advanced models may need better resources.

Issues:

- The database models mostly contain more non-diabetic patient cases rather than diabetic patient cases which can be bias model prediction if not trained properly.
- Models trained on a specific dataset may not perform the same or with the same accuracy on different regions.
- Without proper validation the model may perform with better accuracy on the testing but different on unseen data.

2.5 Dataset Information and Background

The dataset used in this project is a Diabetes Classification Dataset. The dataset was obtained from Opendatabay.

https://www.opendatabay.com/data/healthcare/4ea2d478-76c7-42eb-9c80-0d69050d2e40

The dataset contains 5132 rows and 11 columns with the following attributes.

- Age
- Gender
- Body Mass Index (BMI)
- Cholesterol (Chol)
- Triglycerides (TG)
- High Density Lipoprotein (HDL)
- Low Density Lipoprotein (LDL)
- Creatinine (Cr)
- Blood Urea Nitrogen (BUN)
- Diagnosis

```
df = pd.read_csv('Diabetes_Classification.csv')
df
```

	Unnamed: 0	Age	Gender	BMI	Chol	TG	HDL	LDL	Cr	BUN	Diagnosis
0	0	50	F	24	4.20	0.90	2.40	1.40	46.0	4.70	0
1	1	26	M	23	3.70	1.40	1.10	2.10	62.0	4.50	0
2	2	33	M	21	4.90	1.00	0.80	2.00	46.0	7.10	0
3	3	45	F	21	2.90	1.00	1.00	1.50	24.0	2.30	0
4	4	50	F	24	3.60	1.30	0.90	2.10	50.0	2.00	0
...	...	...	...	...	...	...	...	...	...	...	...
5127	5127	54	M	23	5.00	1.50	1.24	2.98	77.0	3.50	1
5128	5128	50	F	22	4.37	2.09	1.37	2.29	47.3	4.40	1
5129	5129	67	M	24	3.89	1.38	1.14	2.17	70.6	4.73	1
5130	5130	60	F	29	5.91	1.29	1.73	2.85	50.2	7.33	1
5131	5131	37	M	34	5.42	2.66	1.08	2.87	75.5	4.61	1

5132 rows × 11 columns

Figure 7: Diabetes Dataset.

The dataset contains both input values X and target values y which is Diagnosis.

3. Solution

3.1 Proposed Solution

The project uses a ML based supervised learning module using classification approach to predict whether a person is diabetic or non-diabetic based on their medical data provided. The prediction process begins with the loading the dataset and pre-processing the data such as handling the categorical variables like gender. The pre-processed data is then divided into training and test sets. For this model three best classification algorithms were selected (Logistic Regression, Random Forest Classifier and SVM). These three algorithms are trained and are evaluated using performance metrics like accuracy, precision to measure their effectiveness in the diabetes prediction. The proposed solution helps in identifying early diabetes cases which might help prevent the individuals from serious diabetic cases.

3.2 Algorithms Used

- **Random Forest Classifier**

Random Forest Classifier is a supervised learning algorithm that builds or combines multiple decision trees to produce an accurate single result. Random forest classifier can handle both numerical and categorical data. This algorithm can handle both numerical and categorical data and is capable of capturing complex relationships between functions (Kavlakoglu, 2025). This algorithm is suitable for this project because medical data mostly contains non-linear interactions between variables such as BMI, age and the ensemble of this algorithm reduces overfitting.

- **Logistic Regression**

Logistic regression is a supervised machine learning algorithm widely used for binary classification problems. Logistic regression is a type of classification algorithm that predicts discrete binary outcomes (Lee, 202). Logistic regression is suited for this project because it is simple, efficient and also allows easy understanding of how medical data contributes to the diabetes prediction.

- **Support Vector Machine (SVM)**

Support Vector Machine (SVM) is a supervised learning algorithm that classifies data by finding an optimal line or hyperplane that separates different classes with maximum margin. SVM is effective in high dimensional spaces and can also model both linear and non-linear decision boundaries (Kavlakogl, 2025). SVM is perfect for this project because it performs best with medium sized datasets and can handle complex decisions. SVM is robust to overfitting making it effective for medical projects and diabetes prediction.

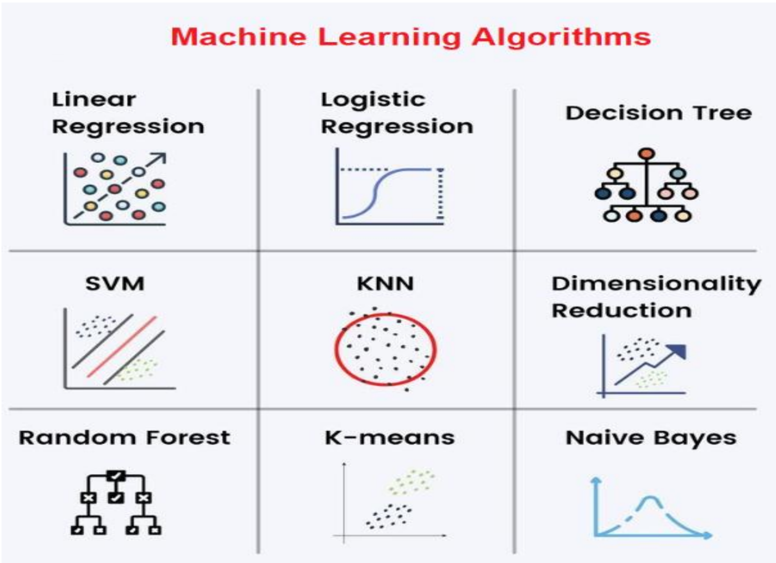


Figure 8: Machine Learning Algorithms.

3.3 Pseudocode

Pseudocode is a detailed description of what a program or algorithm should do. It is written in a formal and readable natural syntax and formatting state so that programmers can easily understand the development process. It basically serves as a blueprint for translating codes logic into an actual programming language (Sheldon, 2023).

3.3.1 Pseudocode for overall system

START

IMPORT required libraries

LOAD diabetes dataset

DISPLAY dataset information and check data type

IF missing values exists

THEN handle missing values

END IF

SELECT input features (Age, Gender, BMI, Chol, TG, HDL, LDL, Creatinine, BUN)

SELECT target variable (Diagnosis)

GENERATE heatmap, bar graph and scatter plot

INITIALIZE Logistic Regression model

TRAIN Logistic Regression model

PREDICT test results

CALCULATE accuracy score

INITIALIZE SVM model

TRAIN SVM model

PREDICT test results

CALCULATE accuracy score

INITIALIZE random forest model

TRAIN random forest model

PREDICT test results

CALCULATE accuracy score

COMPARE accuracy scores of all the models

SELECT model with highest accuracy

DISPLAY best performing algorithm

IF new patient data is provided

  APPLY same pre-processing steps

  PREDICT diabetes status

  DISPLAY result

END IF

END

3.3.2 Pseudocode for Logistic Regression

START

IMPORT required libraries

LOAD diabetes dataset

SELECT input features (Age, Gender, BMI, Chol, TG, HDL, LDL, Creatinine, BUN)

SELECT target variable (Diagnosis)

PREPROCESS dataset

SPLIT X and y into training and testing sets

INITIALIZE Logistic Regression model

TRAIN model using training data

CALCULATE predicted values for test data

FOR EACH prediction

APPLY sigmoid decision boundary

CLASSIFY as Diabetic or Non-Diabetic

END FOR

EVALUATE model performance using Accuracy score and Classification Report

RETURN Logistic Regression accuracy

END

3.3.3 Pseudocode for Random Forest Classifier

START

IMPORT required libraries

LOAD diabetes dataset

SELECT input features (Age, Gender, BMI, Chol, TG, HDL, LDL, Creatinine, BUN)

SELECT target variable (Diagnosis)

PREPROCESS dataset

SPLIT X and y into training and testing sets

INITIALIZE Random Forest with N decision trees

TRAIN model using training data

FOR EACH tree in forest

DO

SELECT random sample of data

BUILD decision tree

END FOR

CALCULATE predicted values for test data

EVALUATE model performance using Accuracy score, Precision, Recall, F1-score

RETURN random forest accuracy

END

3.3.4 Pseudocode for Support Vector Machine (SVM)

START

IMPORT required libraries

LOAD diabetes dataset

SELECT input features (Age, Gender, BMI, Chol, TG, HDL, LDL, Creatinine, BUN)

SELECT target variable (Diagnosis)

PREPROCESS dataset

SPLIT X and y into training and testing sets

INITIALIZE SVM classifier with kernel function

MAP input data

FIND optional hyperplane that maximizes margin

SEPARATE diabetic and non-diabetic classes

PREDICT class labels for test data

EVALUATE model performance using Accuracy score, classification report

RETURN SVM accuracy

END

3.4 Flowchart

3.4.1 Flowchart Diagram of overall system

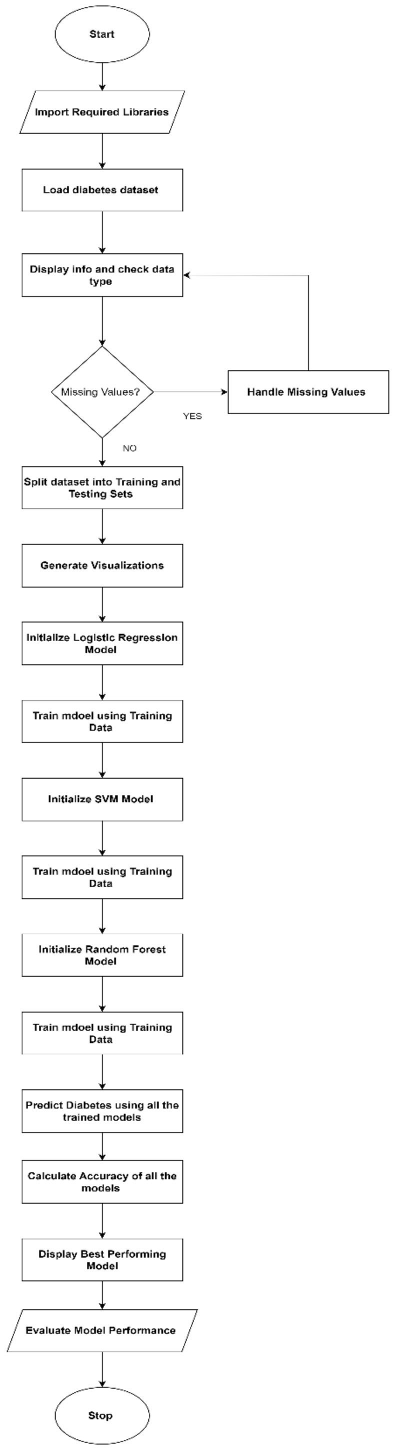


Figure 9: Flowchart Diagram of Overall System.

3.4.1 Flowchart Diagram of Logistic Regression

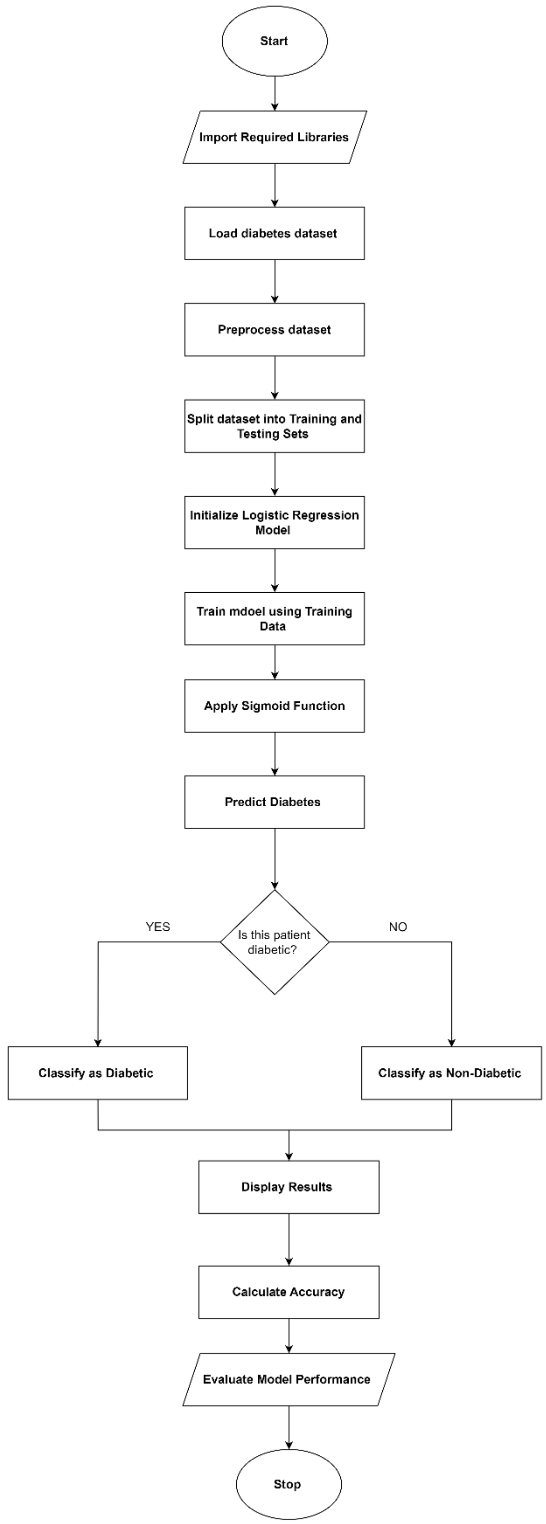


Figure 10: Flowchart Diagram of Logistic Regression.

3.4.2 Flowchart Diagram of Random Forest Classifier

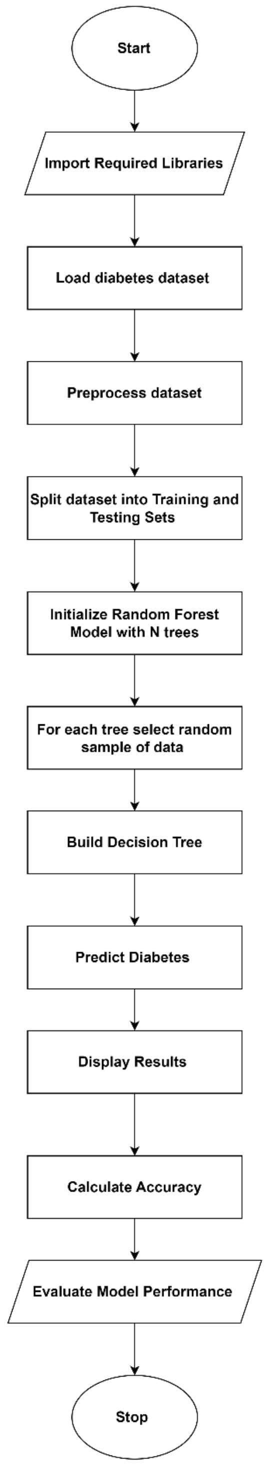


Figure 11: Flowchart Diagram of Random Forest Classifier.

3.4.3 Flowchart Diagram of Support Vector Machine (SVM)

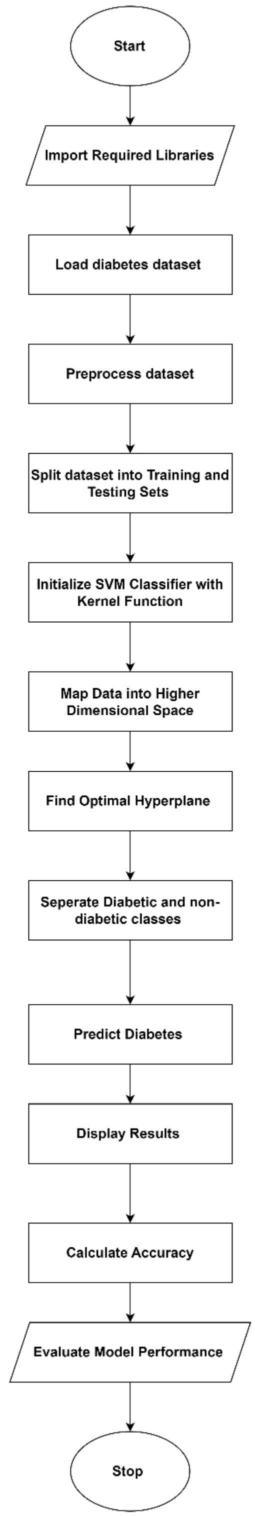


Figure 12: Flowchart Diagram of Support Vector Machine.

3.5 Development Process

The development of this project was carried out using the python programming language for proper data analytics and machine learning. This project was implemented on the Jupiter notebook platform.

The development process began with the data cleaning, fixing nan values and pre-prepossessing the data. Data visualization was done with the help of the heat map, bar graph and scatter plot to understand the data properly. After the data preparation supervised learning models were implemented such as Logistic regression, SVM and Random Forest Classifier. The models were evaluated using the performance tests such as recall, F1-score, precision and accuracy to know about the prediction effectiveness.

The following key libraries were used for the development process such as pandas for the data loading and pre-processing of the diabetes dataset, NumPy for the numerical computations, Scikit-learn used for implementing machine learning algorithms for train test split, matplotlib and seaborn for the data visualization.

3.6 Achieved Results

3.6.1 Libraries Import

Libraries Import

[234]:

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns

from sklearn.model_selection import train_test_split
from sklearn.linear_model import LogisticRegression
from sklearn.svm import SVC
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import accuracy_score

from sklearn.metrics import accuracy_score, classification_report, confusion_matrix
```

Figure 13: Libraries Import.

All the required libraries are being imported in this phase at the beginning of the development process. Libraries such as Pandas and NumPy are being used for data handling and numerical operations. Matplotlib and Seaborn are used for data visualization process.

3.6.2 Version Check

Version Check

```
[237]: import sys
       print(sys.version)

       3.12.7 | packaged by Anaconda, Inc. | (main, Oct  4 2024, 13:17:27) [MSC v.1929 64 bit (AMD64)]

[239]: pd.__version__

[239]: '2.2.2'

[241]: np.__version__

[241]: '1.26.4'
```

Figure 14: Version Check.

This image displays the version information of the python key libraries.

3.6.3 CSV file read

Diabetes.csv file read

[10]:
```
df = pd.read_csv('Diabetes_Classification.csv')
df
```

[10]:

	Unnamed: 0	Age	Gender	BMI	Chol	TG	HDL	LDL	Cr	BUN	Diagnosis
0	0	50	F	24	4.20	0.90	2.40	1.40	46.0	4.70	0
1	1	26	M	23	3.70	1.40	1.10	2.10	62.0	4.50	0
2	2	33	M	21	4.90	1.00	0.80	2.00	46.0	7.10	0
3	3	45	F	21	2.90	1.00	1.00	1.50	24.0	2.30	0
4	4	50	F	24	3.60	1.30	0.90	2.10	50.0	2.00	0
...	...	...	...	...	...	...	...	...	...	...	...
5127	5127	54	M	23	5.00	1.50	1.24	2.98	77.0	3.50	1
5128	5128	50	F	22	4.37	2.09	1.37	2.29	47.3	4.40	1
5129	5129	67	M	24	3.89	1.38	1.14	2.17	70.6	4.73	1
5130	5130	60	F	29	5.91	1.29	1.73	2.85	50.2	7.33	1
5131	5131	37	M	34	5.42	2.66	1.08	2.87	75.5	4.61	1

5132 rows × 11 columns

Figure 15: Diabeties.csv file read.

This image displays the “Diabetes_Classification.csv” file being read by the Jupiter notebook environment.

3.6.4 Unique Values

Unique Values

```
[11]: df['Diagnosis'].unique()

[11]: array([0, 1], dtype=int64)

[70]: df['Age'].unique()

[70]: array([50, 26, 33, 45, 48, 43, 32, 31, 30, 49, 42, 39, 41, 44, 47, 36, 38,
             46, 35, 40, 59, 51, 57, 63, 25, 60, 77, 54, 34, 55, 28, 56, 52, 69,
             73, 61, 58, 53, 66, 68, 62, 64, 67, 70, 79, 76, 65, 75, 20, 71, 37,
             27, 85, 29, 78, 22, 74, 72, 24, 84, 80, 83, 82, 86, 93, 23, 87, 88,
             81, 90, 91], dtype=int64)

[12]: df['Gender'].unique()

[12]: array(['F', 'M', 'f'], dtype=object)
```

Figure 16: Unique Values.

This image displays the unique values of the Diagnosis column and the unique values of the Gender attribute in the dataset.

3.6.5 Data Pre-processing

Data Preprocessing

```
[13]: df['Gender'] = df['Gender'].astype(str).str.strip().str.upper()
      df['Gender'] = df['Gender'].map({'F': 0, 'M': 1})
      df['Gender'] = df['Gender'].astype(int)
```

Figure 17: Data Preprocessing.

This image displays the pre-processing stage where the dataset is prepared for the model training converting the values into upper case.

3.6.6 Data Visualization

Data visualization

```
[168]: corr = df.corr()

# Plot heatmap
plt.figure(figsize=(10, 8))
sns.heatmap(
    corr,
    annot=True,
    cmap='coolwarm',
    fmt=".2f",
    linewidths=0.5
)

plt.title("Correlation Heatmap of Diabetes Dataset")
plt.show()
```

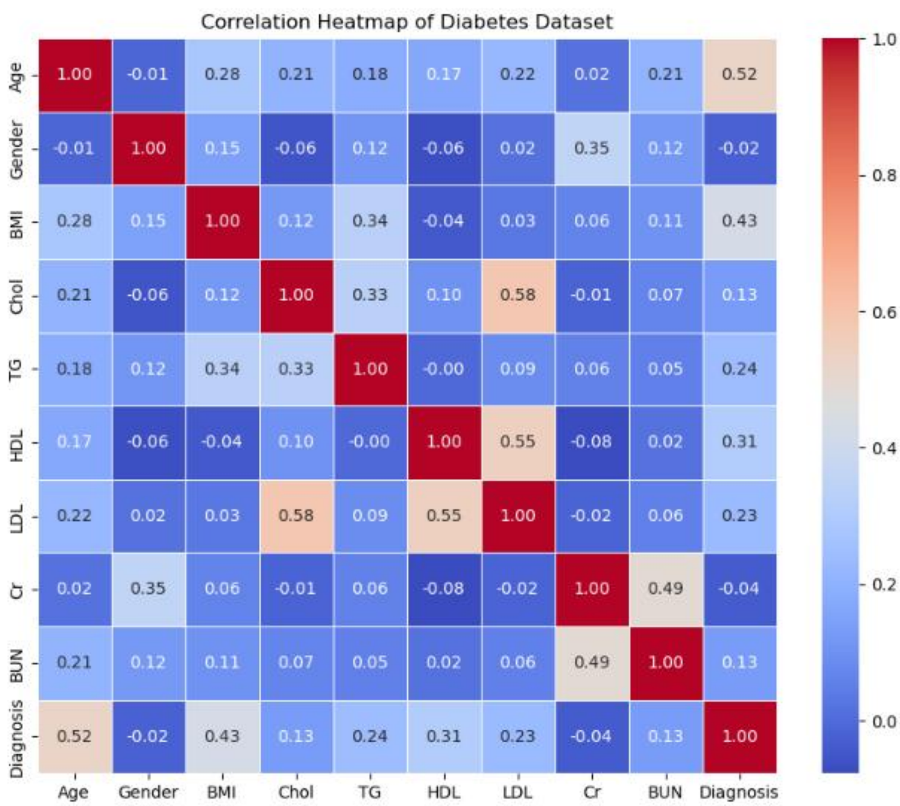
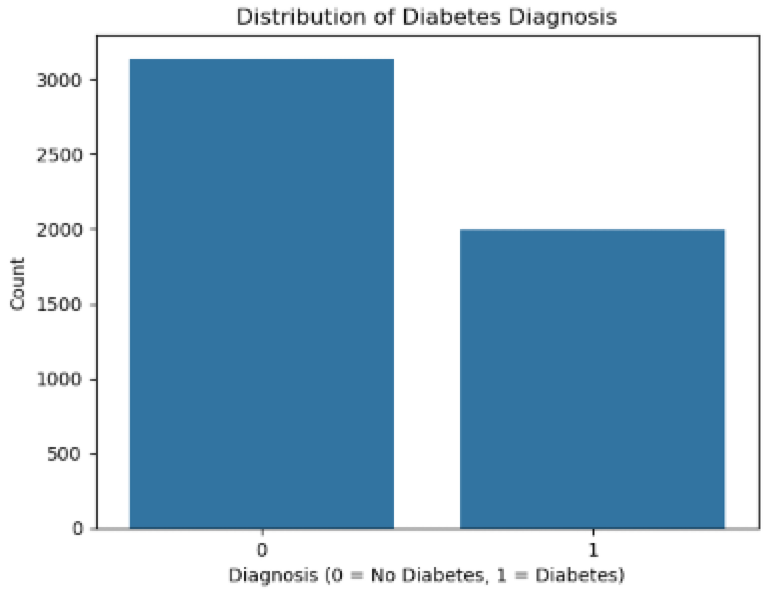


Figure 18: Data Visualization (Heatmap).

```
[169]: plt.figure()
       sns.countplot(x='Diagnosis', data=df)

       plt.xlabel('Diagnosis (0 = No Diabetes, 1 = Diabetes)')
       plt.ylabel('Count')
       plt.title('Distribution of Diabetes Diagnosis')
       plt.show()
```




```
[170]: plt.figure()
       sns.barplot(x='Diagnosis', y='BMI', data=df)

       plt.xlabel('Diagnosis')
       plt.ylabel('Average BMI')
       plt.title('Average BMI by Diabetes Diagnosis')
       plt.show()
```

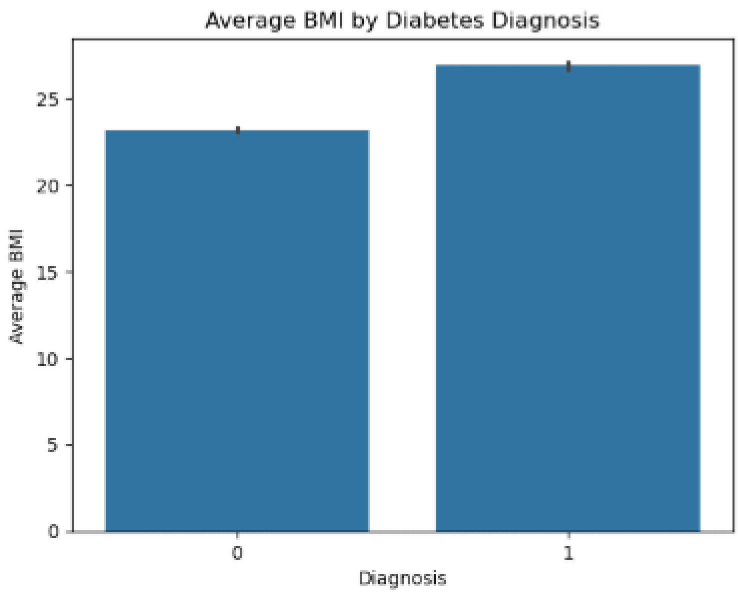


Figure 19: Data Visualization (Bar graph).

[171]:

```
plt.figure()

for label in df['Diagnosis'].unique():
    subset = df[df['Diagnosis'] == label]
    plt.scatter(subset['Age'], subset['BMI'], label=f'Diagnosis {label}')

plt.xlabel('Age')
plt.ylabel('BMI')
plt.title('Age vs BMI by Diabetes Diagnosis')
plt.legend()
plt.show()
```

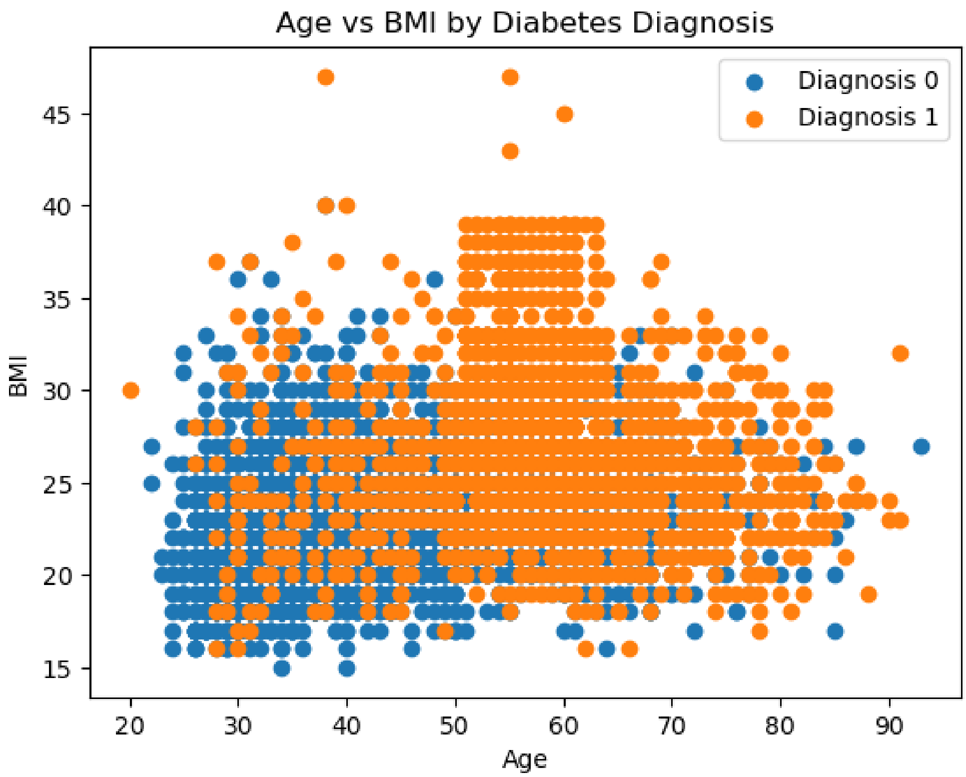


Figure 20: Data Visualization (Scatterplot).

3.6.7 Dataset Information

Dataset Info

[154]:
```
df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 5132 entries, 0 to 5131
Data columns (total 11 columns):
 #   Column      Non-Null Count  Dtype
---  ------      --------------  -----
 0   Unnamed: 0  5132 non-null   int64
 1   Age         5132 non-null   int64
 2   Gender      5132 non-null   int32
 3   BMI         5132 non-null   int64
 4   Chol        5132 non-null   float64
 5   TG          5132 non-null   float64
 6   HDL         5132 non-null   float64
 7   LDL         5132 non-null   float64
 8   Cr          5132 non-null   float64
 9   BUN         5132 non-null   float64
 10  Diagnosis   5132 non-null   int64
dtypes: float64(6), int32(1), int64(4)
memory usage: 421.1 KB
```

[156]:
```
df.isnull().sum()
```

[156]:
```
Unnamed: 0    0
Age           0
Gender        0
BMI           0
Chol          0
TG            0
HDL           0
LDL           0
Cr            0
BUN           0
Diagnosis     0
dtype: int64
```

[158]:
```
df['Gender'].unique()
```

[158]:
```
array([0, 1])
```

[160]:
```
print(df['Diagnosis'].value_counts())
```

```
Diagnosis
0    3139
1    1993
Name: count, dtype: int64
```

[162]:
```
df.shape
```

[162]:
```
(5132, 11)
```

Figure 21: Dataset Info.

This image displays the information about the diabetes dataset.

3.6.8 Drop Table

Drop Tables

```
[29]: df = df.drop(columns=['Unnamed: 0'])
      df
```

[29]:

	Age	Gender	BMI	Chol	TG	HDL	LDL	Cr	BUN	Diagnosis
0	50	0	24	4.20	0.90	2.40	1.40	46.0	4.70	0
1	26	1	23	3.70	1.40	1.10	2.10	62.0	4.50	0
2	33	1	21	4.90	1.00	0.80	2.00	46.0	7.10	0
3	45	0	21	2.90	1.00	1.00	1.50	24.0	2.30	0
4	50	0	24	3.60	1.30	0.90	2.10	50.0	2.00	0
...	...	...	...	...	...	...	...	...	...	...
5127	54	1	23	5.00	1.50	1.24	2.98	77.0	3.50	1
5128	50	0	22	4.37	2.09	1.37	2.29	47.3	4.40	1
5129	67	1	24	3.89	1.38	1.14	2.17	70.6	4.73	1
5130	60	0	29	5.91	1.29	1.73	2.85	50.2	7.33	1
5131	37	1	34	5.42	2.66	1.08	2.87	75.5	4.61	1

5132 rows × 10 columns

Figure 22: Drop Tables.

Unnamed: 0 column is being dropped.

3.6.9 X and y train

X and y train

```
[174]: X = df.drop(columns=['Diagnosis'])
       X
```

[174]:

	Age	Gender	BMI	Chol	TG	HDL	LDL	Cr	BUN
0	50	0	24	4.20	0.90	2.40	1.40	46.0	4.70
1	26	1	23	3.70	1.40	1.10	2.10	62.0	4.50
2	33	1	21	4.90	1.00	0.80	2.00	46.0	7.10
3	45	0	21	2.90	1.00	1.00	1.50	24.0	2.30
4	50	0	24	3.60	1.30	0.90	2.10	50.0	2.00
...	...	...	...	...	...	...	...	...	...
5127	54	1	23	5.00	1.50	1.24	2.98	77.0	3.50
5128	50	0	22	4.37	2.09	1.37	2.29	47.3	4.40
5129	67	1	24	3.89	1.38	1.14	2.17	70.6	4.73
5130	60	0	29	5.91	1.29	1.73	2.85	50.2	7.33
5131	37	1	34	5.42	2.66	1.08	2.87	75.5	4.61

5132 rows × 9 columns

```
[175]: y = df['Diagnosis']
       y
```

```
[175]: 0       0
       1       0
       2       0
       3       0
       4       0
              ..
       5127    1
       5128    1
       5129    1
       5130    1
       5131    1
       Name: Diagnosis, Length: 5132, dtype: int64
```

Figure 23: X and y Train 1.

```
[176]: X_train, X_test, y_train, y_test = train_test_split(X,y,test_size=0.2,random_state=42)
X_train
```

[176]:

	Age	Gender	BMI	Chol	TG	HDL	LDL	Cr	BUN
3740	36	0	20	4.87	0.79	1.580000	2.980000	54.1	3.68
949	43	0	22	4.08	0.61	1.740000	2.030000	49.3	4.59
3826	25	1	20	4.60	1.00	1.330000	2.630000	80.0	4.25
19	33	0	24	4.20	1.50	1.200000	2.300000	62.0	5.30
3049	72	0	21	5.33	1.55	1.200000	2.940000	65.0	3.56
...	...	...	...	...	...	...	...	...	...
4426	58	1	27	4.98	2.48	4.860753	4.860753	68.3	6.24
466	63	1	30	3.60	5.10	0.900000	2.500000	63.0	5.90
3092	40	1	16	4.00	0.00	1.000000	2.000000	67.0	4.00
3772	42	1	23	4.00	0.80	1.120000	2.460000	96.0	5.49
860	78	1	25	5.83	2.10	1.080000	3.500000	81.4	4.30

4105 rows × 9 columns

```
[178]: y_train
```

```
[178]: 3740    0
949     0
3826    0
19      0
3049    0
       ..
4426    1
466     1
3092    0
3772    0
860     0
Name: Diagnosis, Length: 4105, dtype: int64
```

Figure 24: X and y Train 2.

This image displays the separation of the X and y variables. All the medical attributes such as age, BMI, cholesterol, TG, HDL, LDL, Cr, BUN is the know variables X and the Diagnosis is the y variable.

[179]:
```
X_test
```

[179]:

	Age	Gender	BMI	Chol	TG	HDL	LDL	Cr	BUN
5106	68	1	29	5.22	3.55	0.87	2.46	93.8	4.89
2186	28	0	22	4.38	1.17	1.39	1.87	46.0	4.00
2589	54	1	18	4.09	0.96	1.43	2.50	81.7	7.68
831	40	0	19	5.87	1.29	1.75	3.37	61.1	4.10
1421	41	0	22	4.50	0.50	1.75	1.94	52.0	3.12
...	...	...	...	...	...	...	...	...	...
1662	53	1	23	4.03	1.57	1.03	2.56	72.4	6.00
833	36	1	26	6.69	3.49	0.91	3.64	67.5	3.86
366	69	0	32	5.30	3.80	1.40	2.30	243.0	14.50
3778	30	1	19	4.11	1.27	1.27	2.40	88.8	6.11
1235	78	0	27	4.87	1.40	1.05	3.03	47.4	5.60

1027 rows × 9 columns

[180]:
```
y_test
```

[180]:
```
5106    1
2186    0
2589    0
831     0
1421    0
       ..
1662    0
833     0
366     1
3778    0
1235    0
Name: Diagnosis, Length: 1027, dtype: int64
```

Figure 25: X and y Train 3.

3.6.10 Logistic Regression Model Training

Logistic Regression Model Training

```
[182]: lr = LogisticRegression(max_iter=5000)
       lr

[182]: LogisticRegression
       LogisticRegression(max_iter=5000)

[190]: lr.fit(X_train, y_train)

[190]: LogisticRegression
       LogisticRegression(max_iter=5000)

[274]: y_pred_lr = lr.predict(X_test)
       y_pred_lr

[274]: array([1, 0, 0, ..., 1, 0, 1], dtype=int64)

[280]: # Evaluation
       print("Logistic Regression Accuracy:", accuracy_score(y_test, y_pred_lr))
       print(confusion_matrix(y_test, y_pred_lr))
       print(classification_report(y_test, y_pred_lr))

       Logistic Regression Accuracy: 0.8023369036027264
       [[529  75]
        [128 295]]
                     precision    recall  f1-score   support

                  0       0.81      0.88      0.84       604
                  1       0.80      0.70      0.74       423

           accuracy                           0.80      1027
          macro avg       0.80      0.79      0.79      1027
       weighted avg       0.80      0.80      0.80      1027
```

Figure 26: Logistic Regression Model Training.

The training process is done for the logistic regression where the machine learns the patterns from the training data to predict the diabetic and non-diabetic patients. The evaluation includes the classification report including the precision, recall, f1-score and support.

3.6.11 SVM Classifier Model Training

SVM Classifier Model Training

```
[282]: svm = SVC(kernel='rbf', probability=True)
       svm
```

```
[282]: SVC
       SVC(probability=True)
```

```
[284]: svm.fit(X_train, y_train)
```

```
[284]: SVC
       SVC(probability=True)
```

```
[286]: y_pred_svm = svm.predict(X_test)
        y_pred_svm
```

```
[286]: array([1, 0, 0, ..., 1, 0, 1], dtype=int64)
```

```
[312]: # Evaluation
       print("SVM Accuracy:", accuracy_score(y_test, y_pred_svm))
       print(confusion_matrix(y_test, y_pred_svm))
       print(classification_report(y_test, y_pred_svm))
```

```
SVM Accuracy: 0.8003894839337877
[[510  94]
 [111 312]]
              precision    recall  f1-score   support

           0       0.82      0.84      0.83       604
           1       0.77      0.74      0.75       423

    accuracy                           0.80      1027
   macro avg       0.79      0.79      0.79      1027
weighted avg       0.80      0.80      0.80      1027
```

Figure 27: SVM Classifier Model Training.

The training process is done for the SVM classifier where the machine learns the patterns from the training data to predict the diabetic and non-diabetic patients. The evaluation includes the classification report including the precision, recall, f1-score and support.

3.6.12 Random Forest Model Training

Random Forest Model Training

```
[296]: rf = RandomForestClassifier(n_estimators=100,random_state=42)
       rf
```

```
[296]: RandomForestClassifier
       RandomForestClassifier(random_state=42)
```

```
[298]: rf.fit(X_train, y_train)
```

```
[298]: RandomForestClassifier
       RandomForestClassifier(random_state=42)
```

```
[300]: y_pred_rf = rf.predict(X_test)
       y_pred_rf
```

```
[300]: array([1, 0, 0, ..., 1, 0, 1], dtype=int64)
```

```
[314]: # Evaluation
       print("Random Forest Accuracy:", accuracy_score(y_test, y_pred_rf))
       print(confusion_matrix(y_test, y_pred_rf))
       print(classification_report(y_test, y_pred_rf))
```

```
Random Forest Accuracy: 0.8208373904576436
[[537  67]
 [117 306]]
              precision    recall  f1-score   support

           0       0.82      0.89      0.85       604
           1       0.82      0.72      0.77       423

    accuracy                           0.82      1027
   macro avg       0.82      0.81      0.81      1027
weighted avg       0.82      0.82      0.82      1027
```

Figure 28: Random Forest Model Training.

The training process is done for the SVM classifier where the machine learns the patterns from the training data to predict the diabetic and non-diabetic patients. The evaluation includes the classification report including the precision, recall, f1-score and support.

3.6.13 Diabetic Patient Predictor

Diabetic Patient Predictor

[308]:
```python
# Example diabetic patient data
new_patient = pd.DataFrame([{
    'Age': 45,
    'Gender': 1,       # 1 = Male, 0 = Female
    'BMI': 28.4,
    'Chol': 210,
    'TG': 160,
    'HDL': 45,
    'LDL': 130,
    'Cr': 1.1,
    'BUN': 18
}])
```

[324]:
```python
# Predictions with Yes/No
print("Logistic Regression Prediction:", "Yes" if lr.predict(new_patient) == [1] else "No")
print("SVM Prediction:", "Yes" if svm.predict(new_patient) == [1] else "No")
print("Random Forest Prediction:", "Yes" if rf.predict(new_patient) == [1] else "No")
```

```
Logistic Regression Prediction: Yes
SVM Prediction: Yes
Random Forest Prediction: Yes
```

Figure 29: Diabetic Patient Predictor.

A sample of diabetic dataset is created using random patient medical report. The data is then passed through all the trained models (Logistic, SVM, Random Forest). Each model evaluates the input features and predicted data is then displayed.

3.6.14 Non-Diabetic Patient Predictor

Non diabetic Patient Predictor

```
[204]: non_diabetic_patient = pd.DataFrame([{
           'Age': 50,
           'Gender': 0,
           'BMI': 24,
           'Chol': 4.20,
           'TG': 0.90,
           'HDL': 2.40,
           'LDL': 1.40,
           'Cr': 46.0,
           'BUN': 4.70
       }])
```

```
[316]: # Predictions
       print("Logistic Prediction:", "Yes" if lr.predict(non_diabetic_patient) == [1] else "No")
       print("SVM Prediction:", "Yes" if svm.predict(non_diabetic_patient) == [1] else "No")
       print("Random Forest Prediction:", "Yes" if rf.predict(non_diabetic_patient) == [1] else "No")
```

```
Logistic Prediction: Yes
SVM Prediction: No
Random Forest Prediction: No
```

Figure 30: Non-diabetic Patient Predictor.

A sample of non-diabetic dataset is created using random patient medical report. The data is then passed through all the trained models (Logistic, SVM, Random Forest). Each model evaluates the input features and predicted data is then displayed.

3.6.15 Accuracy Score

Accuracy Score

```
[206]: lr_accuracy = accuracy_score(y_test, y_pred_lr)
       svm_accuracy = accuracy_score(y_test, y_pred_svm)
       rf_accuracy = accuracy_score(y_test, y_pred_rf)

       print("Logistic Regression Accuracy:", lr_accuracy)
       print("SVM Accuracy:", svm_accuracy)
       print("Random Forest Accuracy:", rf_accuracy)
```

```
Logistic Regression Accuracy: 0.8023369036027264
SVM Accuracy: 0.8003894839337877
Random Forest Accuracy: 0.8208373904576436
```

Figure 31: Accuracy Score Results.

The accuracy score displays the accuracy of all the algorithms and represents the percentage of the accuracy made by all the used algorithms which helps in identifying the best algorithm for the prediction.

[218]:

```
models = ['Logistic Regression', 'SVM', 'Random Forest']
accuracies = [lr_accuracy, svm_accuracy, rf_accuracy]

plt.figure()
plt.bar(models, accuracies)
plt.xlabel('Machine Learning Models')
plt.ylabel('Accuracy')
plt.title('Accuracy Comparison of Machine Learning Models')
plt.ylim(0, 1)
plt.show()
```

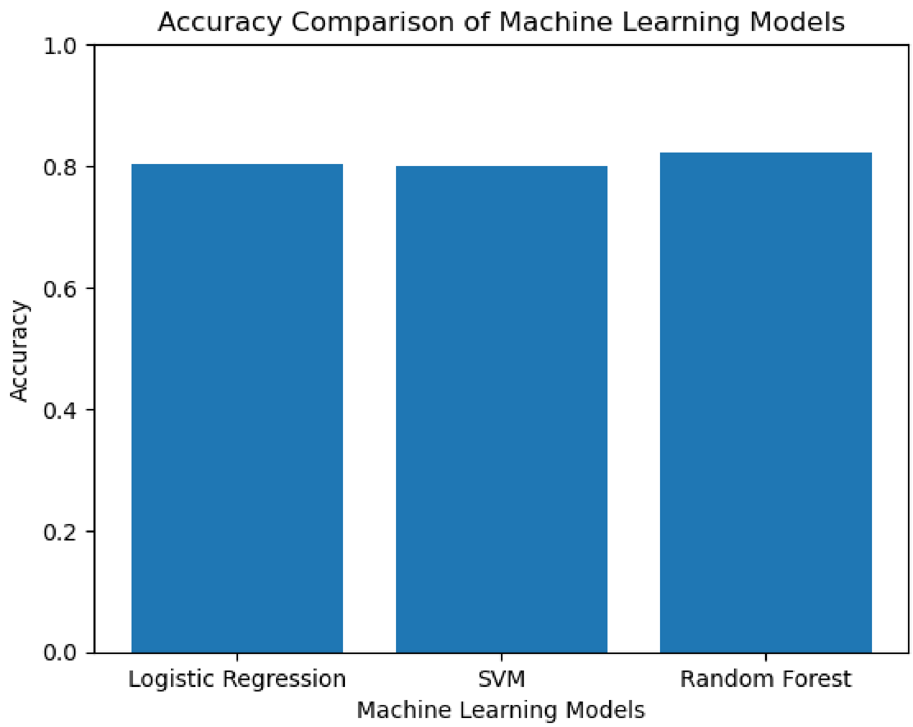


Figure 32: Accuracy Score Bar Graph.

The algorithm score bar graph displays the accuracy graph of all the used algorithms.

3.6.16 Best Performing Algorithm

Best Performing Algorithm

```
[228]: best_index = accuracies.index(max(accuracies))
       best_model = models[best_index]
       best_score = accuracies[best_index]

       print("Best Performing Model:", best_model)
       print("Best Accuracy Score:", best_score)

       Best Performing Model: Random Forest
       Best Accuracy Score: 0.8208373904576436
```

Figure 33: Best Performing Algorithm.

This image displays the best algorithm for this diabetes prediction model using the accuracy score.

Best Performing Score Graph

[230]:
```
plt.figure()
plt.bar(models, accuracies)
plt.xlabel('Machine Learning Models')
plt.ylabel('Accuracy')
plt.title('Best Performing Model Based on Accuracy')
plt.ylim(0, 1)

# Highlight best model with text
plt.text(best_index, best_score,
         f'Best Model\nAccuracy = {best_score:.2f}',
         ha='center', va='bottom')

plt.show()
```

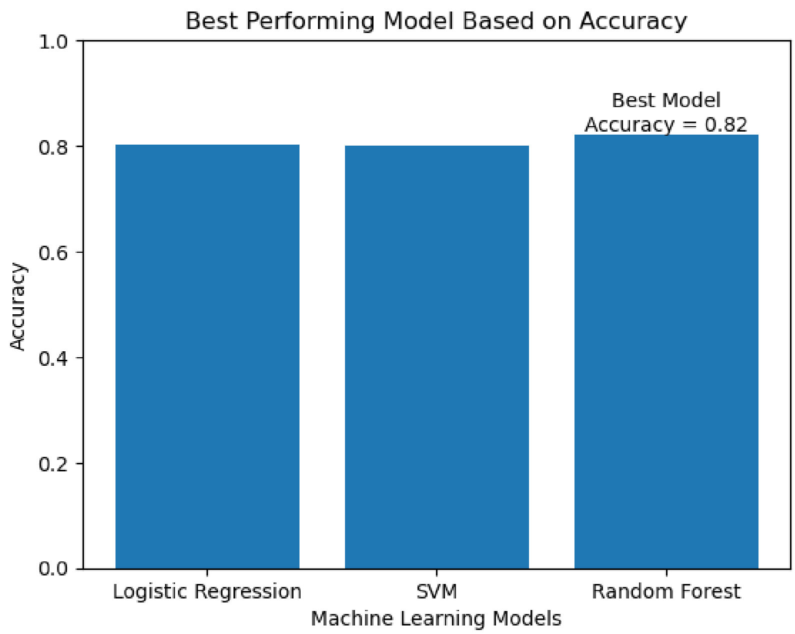


Figure 34: Best Performing Algorithm Bar Graph.

The image displays the bar graph of the best performing algorithm.

4. Conclusion

This diabetes predictor project is focused on developing a predicting system using supervised ML classification. This project focused on analysing structured medical diabetes dataset containing all the needed attributes to build a diabetes prediction model. All the needed data pre-processing was applied to prepare the dataset for the model training. This project was done using three best classification algorithms. Logistic Regression, Random Forest and SVM were implemented to predict whether an individual is diabetic or non-diabetic.

This milestone completed all the necessary data preparation tasks and model training tasks. These included all the data cleaning process, handling missing values and removing the irrelevant columns for suitable machine learning. Data visualization process is done to display the data information such as heatmap, bar graph and scatter plot. These steps help in effective model training

Three supervised learning classification model algorithms (logistic regression, random forest classifier, support vector machine) are being used to train the dataset. The predicted data is then evaluated using performance metrics to calculate the accuracy, precision, recall, f1-score and support. At last, the best performing algorithm is displayed to visualize using bar graph identifying the best performing algorithm. After the successful model training the system was tested using both diabetic and non-diabetic random medical report. The predicted data is then displayed for model evaluation.

5. References

Ahmed, A. (2024) *MDPI* [Online]. Available from: https://www.mdpi.com/2227-9032/13/1/37#healthcare-13-00037-t006 [Accessed 15 December 2025].

B.V., E. (2017) *ScienceDirect* [Online]. Available from: https://www.sciencedirect.com/science/article/pii/S2001037016300733 [Accessed 15 December 2025].

Clinic, C. (2023) *Cleveland Clinic* [Online]. Available from: https://my.clevelandclinic.org/health/diseases/7104-diabetes [Accessed 17 December 2025].

ISO. (2022) *ISO* [Online]. Available from: https://www.iso.org/artificial-intelligence/machine-learning [Accessed 13 December 2025].

Kavlakogl, E. (2025) *IBM* [Online]. Available from: https://www.ibm.com/think/topics/support-vector-machine#684929714 [Accessed 16 December 2025].

Kavlakoglu, E. (2025) *IBM* [Online]. Available from: https://www.ibm.com/think/topics/random-forest#684929713 [Accessed 16 December 2025].

Khan, R. (2022) *National Library of Medicine* [Online]. Available from: https://pmc.ncbi.nlm.nih.gov/articles/PMC10107388/#htl212039-bib-0003 [Accessed 15 December 2025].

Lee, F. (202) *IBM* [Online]. Available from: https://www.ibm.com/think/topics/logistic-regression [Accessed 16 December 2025].

McKinsey. (2024) *McKinsey & Company* [Online]. Available from: https://www.mckinsey.com/featured-insights/mckinsey-explainers/what-is-ai [Accessed 13 December 2025].

Sheldon, R. (2023) *TechTarget* [Online]. Available from: https://www.techtarget.com/whatis/definition/pseudocode [Accessed 16 December 2025].

Tech, M. (2025) *Michigan Tech* [Online]. Available from: https://www.mtu.edu/computing/ai/ [Accessed 13 December 2025].

WHO. (2024) *World Health Organization (WHO)* [Online]. Available from: https://www.who.int/news-room/fact-sheets/detail/diabetes [Accessed 13 December 2025].