



**SAMSUNG**

Samsung Innovation Campus

# PREDICTION OF DIABETES HEALTH INDICATORS

Artificial Intelligence Course

# OUR TEAM

The project presented by:



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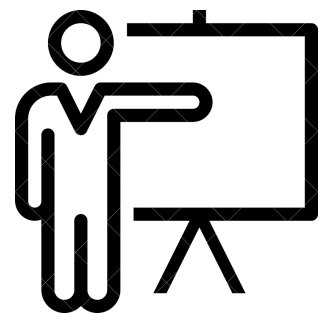
**Ahmad Muhammad**

**Abdullah Saad**

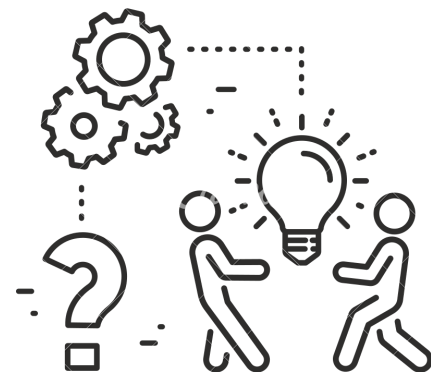
**Facilitator**  
**Eng Ziad omer**

Data used: <https://www.kaggle.com/datasets/alexteboul/diabetes-health-indicators-dataset>

# AGENDA



**Introduction**



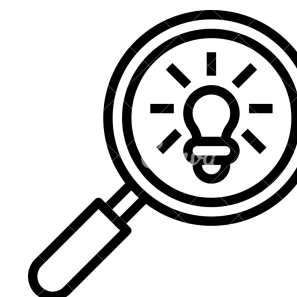
**Problem and Solution**



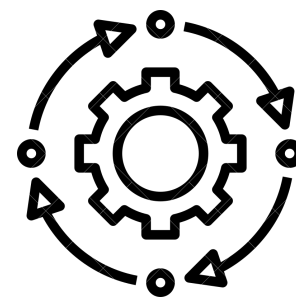
**Data**



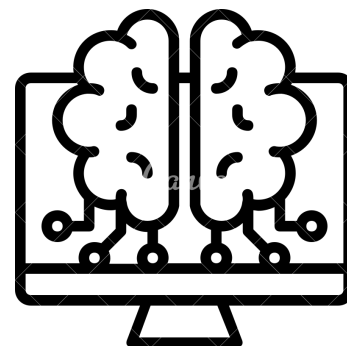
**Data Cleaning**



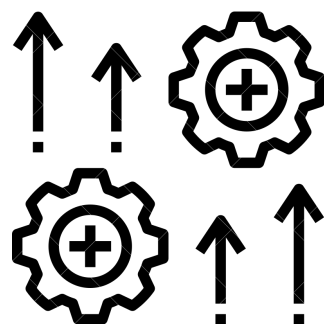
**Exploratory Data Analysis**



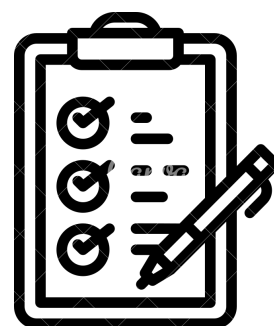
**Data Preprocessing**



**Data Modeling**



**Model Development**



**conclusion**

# INTRODUCTION

Diabetes is a serious condition where your blood glucose level is too high. It can happen when your body doesn't produce enough insulin or the insulin it produces isn't effective. Or, when your body can't produce any insulin at all.

It is a clean dataset of 253,680 survey responses to the CDC's BRFSS2015.

The Behavioral Risk Factor Surveillance System (BRFSS) is a system of health-related telephone surveys that collect state data about U.S. residents the largest continuously conducted health survey system in the world.

The dataset originally has 330 features collected but only 22 features relevant to diabetes are used in our based for machine learning algorithms



## PROBLEM

In this technological and competitive era, our daily lives have been greatly affected and influenced as we are busy with studying and working without concerning our health and fitness.

This resulted in the increase of pre-diabetes and diabetes patients over the years and the rate of prevalence of this disease has never been decreasing. Factors of this problem arise from the varieties of food available in our country, the convenience of ordering food via the super app such as Grab, as well as the lack of regular exercise due to the many workloads and laziness.



## SOLUTION

### Trianing Models for dataset

Logistic Regression

Decision Tree  
Classifier

Random Forest  
Classifier

KNN

To obtain the best model for predicting diabetes and thus reducing the diabetes prevalence rate

Overview

Overview

Alerts 6

Reproduction

Dataset statistics

Number of variables	22
Number of observations	253680
Missing cells	0
Missing cells (%)	0.0%
Duplicate rows	11369
Duplicate rows (%)	4.5%
Total size in memory	42.6 MiB
Average record size in memory	176.0 B

Variable types

Categorical	16
Numeric	6



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# DATA

## Categorical features

- Diabetes binary
- HighBP
- HighChol
- CholCheck
- Smoker
- Stroke
- Fruits
- Veggies
- DiffWalk
- Sex
- HeartDiseaseorAttack
- PhysActivity
- AnyHealthCare
- NoDocbcCost
- HvyAlcoholConsump

## Numerical features

- Age
- BMI
- Income
- MentHlth
- GenHlth
- PhysHlth
- Education

#	Column	Non-Null	Count	Dtype
0	Diabetes_binary	253680	non-null	float64
1	HighBP	253680	non-null	float64
2	HighChol	253680	non-null	float64
3	CholCheck	253680	non-null	float64
4	BMI	253680	non-null	float64
5	Smoker	253680	non-null	float64
6	Stroke	253680	non-null	float64
7	HeartDiseaseorAttack	253680	non-null	float64
8	PhysActivity	253680	non-null	float64
9	Fruits	253680	non-null	float64
10	Veggies	253680	non-null	float64
11	HvyAlcoholConsump	253680	non-null	float64
12	AnyHealthcare	253680	non-null	float64
13	NoDocbcCost	253680	non-null	float64
14	GenHlth	253680	non-null	float64
15	MentHlth	253680	non-null	float64
16	PhysHlth	253680	non-null	float64
17	DiffWalk	253680	non-null	float64
18	Sex	253680	non-null	float64
19	Age	253680	non-null	float64
20	Education	253680	non-null	float64
21	Income	253680	non-null	float64

dtypes: float64(22)

# DATA CLEANING

## Handling Missing values

Diabetes_binary	0
HighBP	0
HighChol	0
CholCheck	0
BMI	0
Smoker	0
Stroke	0
HeartDiseaseorAttack	0
PhysActivity	0
Fruits	0
Veggies	0
HvyAlcoholConsump	0
AnyHealthcare	0
NoDocbcCost	0
GenHlth	0
MentHlth	0
PhysHlth	0
DiffWalk	0
Sex	0
Age	0
Education	0
Income	0

There is no missing values in our dataset

## Checking duplicate values

# Number of duplicates

24206

# drop duplicates

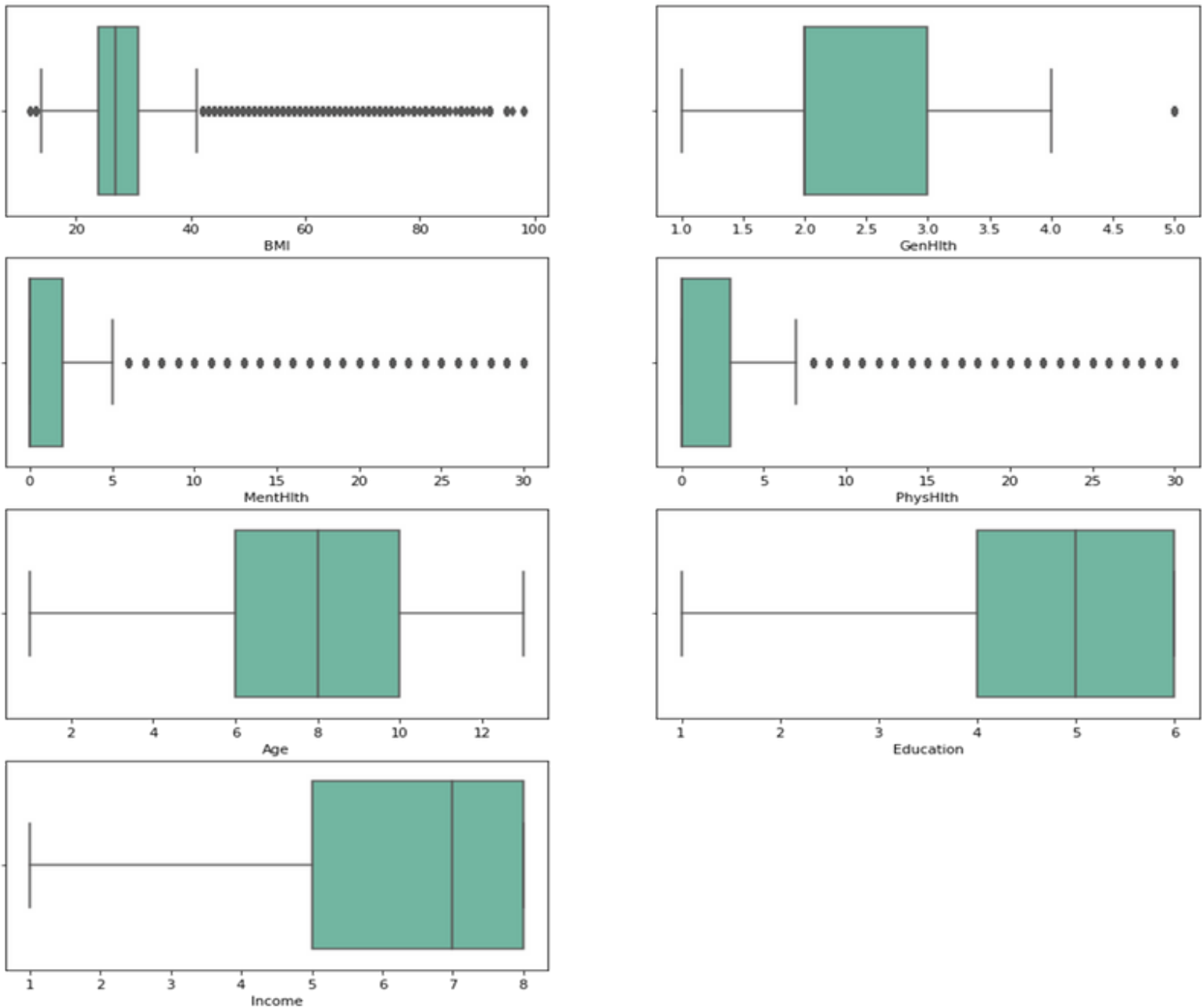
0

There are 24206 duplicated rows in our dataset, it could potentially introduce bias into the model. Therefore, it's a good idea to delete any duplicate rows before training the model.



# DATA CLEANING

## Checking outliers



The box plot shows us that there are outliers, but these values are real data and we must take them into account

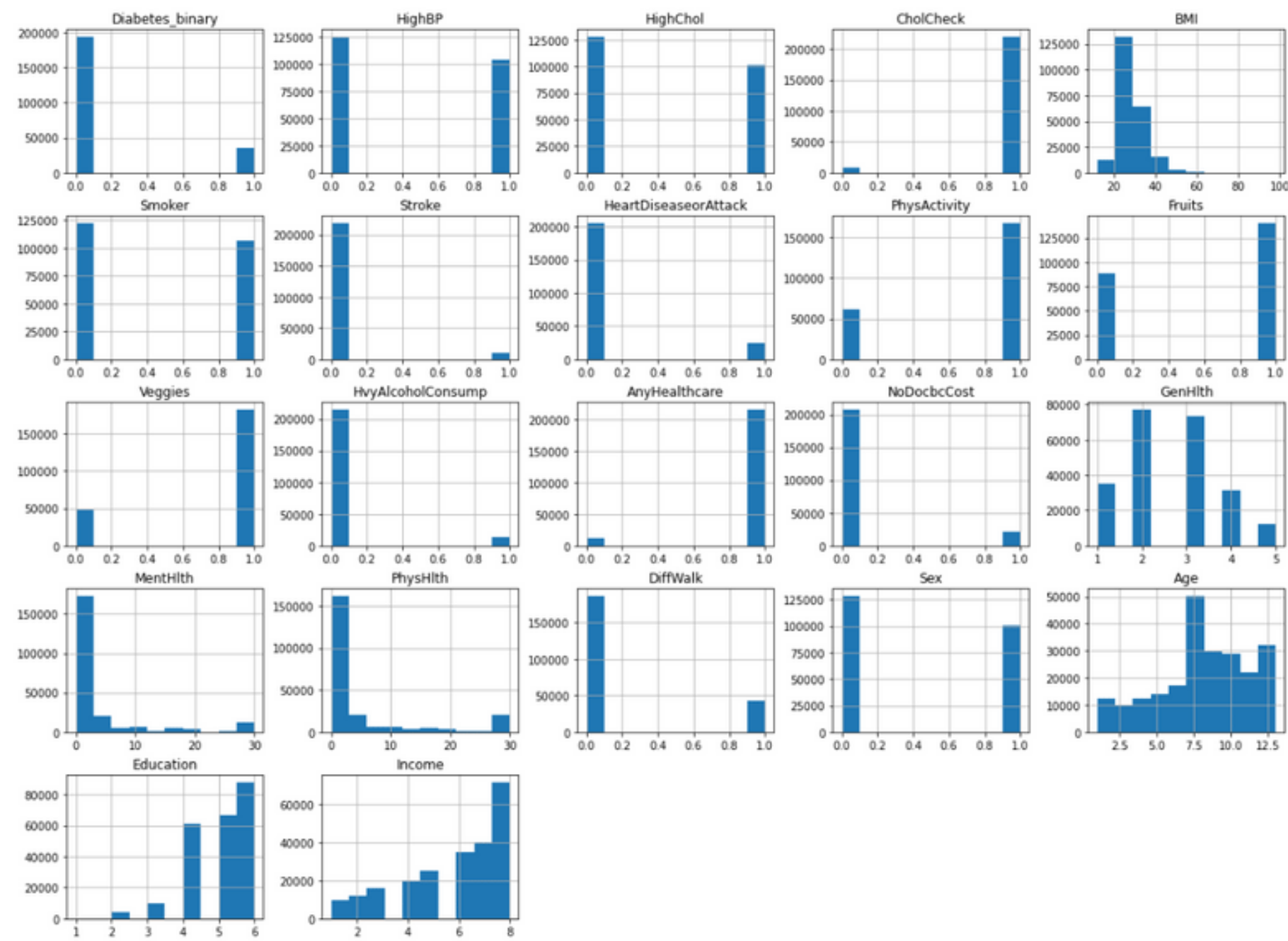
## transform the features type

#	Column	Non-Null Count	Dtype		#	Column	Non-Null Count	Dtype
0	Diabetes_binary	253680 non-null	float64		0	Diabetes_binary	253680 non-null	int32
1	HighBP	253680 non-null	float64		1	HighBP	253680 non-null	int32
2	HighChol	253680 non-null	float64		2	HighChol	253680 non-null	int32
3	CholCheck	253680 non-null	float64		3	CholCheck	253680 non-null	int32
4	BMI	253680 non-null	float64		4	BMI	253680 non-null	int32
5	Smoker	253680 non-null	float64		5	Smoker	253680 non-null	int32
6	Stroke	253680 non-null	float64		6	Stroke	253680 non-null	int32
7	HeartDiseaseorAttack	253680 non-null	float64		7	HeartDiseaseorAttack	253680 non-null	int32
8	PhysActivity	253680 non-null	float64		8	PhysActivity	253680 non-null	int32
9	Fruits	253680 non-null	float64		9	Fruits	253680 non-null	int32
10	Veggies	253680 non-null	float64		10	Veggies	253680 non-null	int32
11	HvyAlcoholConsump	253680 non-null	float64		11	HvyAlcoholConsump	253680 non-null	int32
12	AnyHealthcare	253680 non-null	float64		12	AnyHealthcare	253680 non-null	int32
13	NoDocbcCost	253680 non-null	float64		13	NoDocbcCost	253680 non-null	int32
14	GenHlth	253680 non-null	float64		14	GenHlth	253680 non-null	int32
15	MentHlth	253680 non-null	float64		15	MentHlth	253680 non-null	int32
16	PhysHlth	253680 non-null	float64		16	PhysHlth	253680 non-null	int32
17	DiffWalk	253680 non-null	float64		17	DiffWalk	253680 non-null	int32
18	Sex	253680 non-null	float64		18	Sex	253680 non-null	int32
19	Age	253680 non-null	float64		19	Age	253680 non-null	int32
20	Education	253680 non-null	float64		20	Education	253680 non-null	int32
21	Income	253680 non-null	float64		21	Income	253680 non-null	int32
dtypes: float64(22)					dtypes: int32(22)			

Here we transform the features type to an integer to speed the model and the analysis

# EXPLORATORY DATA ANALYSIS

## Distribution of numerical features



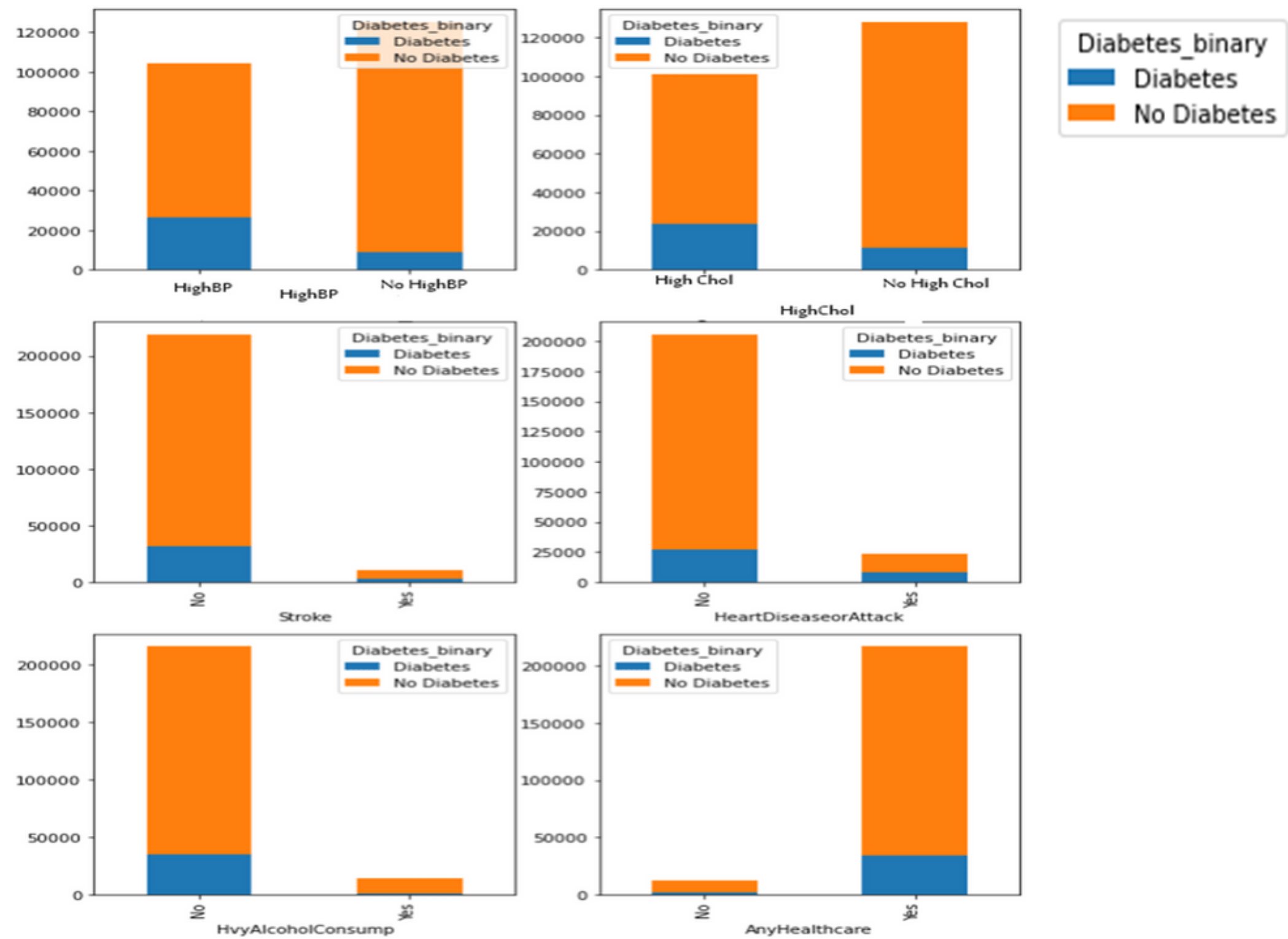
we can see here the value counts of all 22 columns some of them columns are continuous columns and some of them are discrete columns, here is the Frequency of values in different columns.

# EXPLORATORY DATA ANALYSIS

## Distribution of categorical features

People with HighBP and HighChol are diabetics.

In Other columns, according to the percentage of yes and no in each column, it is a normal amount.



# EXPLORATORY DATA ANALYSIS

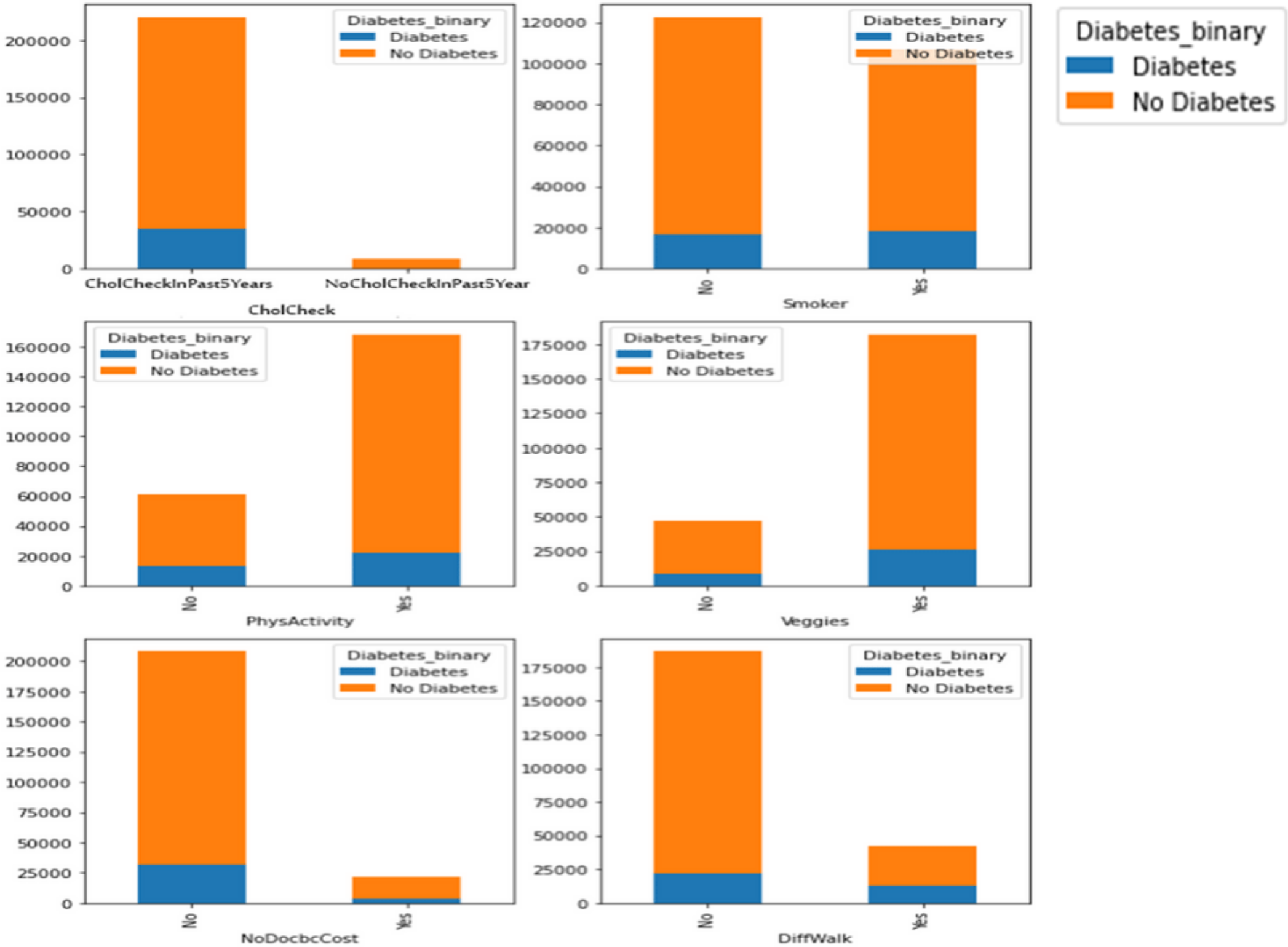
## Distribution of categorical features

People who have cholesterol in the past 5 years are more likely to get diabetes

Smoking doesn't affect diabetics

People who reported physical activity in the past 30 days are more likely to get diabetes, But considering the proportions it is normal

Considering the proportions of the other columns it is normal





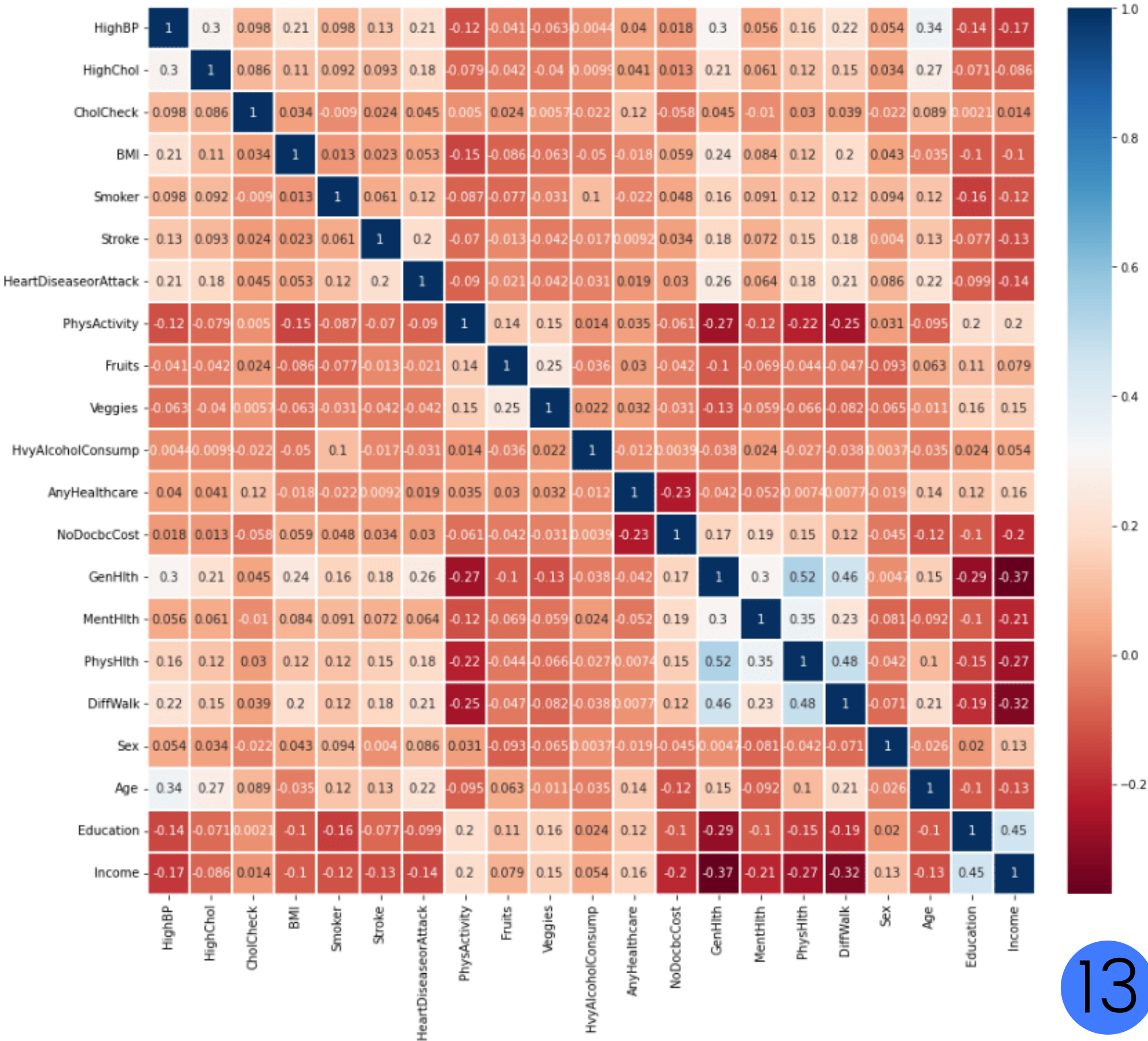
# EXPLORATORY DATA ANALYSIS

Correlation of features

Correlation heatmap shows the relation between columns:

(GenHlth ,PhysHlth ),(PhysHlth, DiffWalk),(GenHlth, DiffWalk )are highly correlated with each other  
=> positive relation

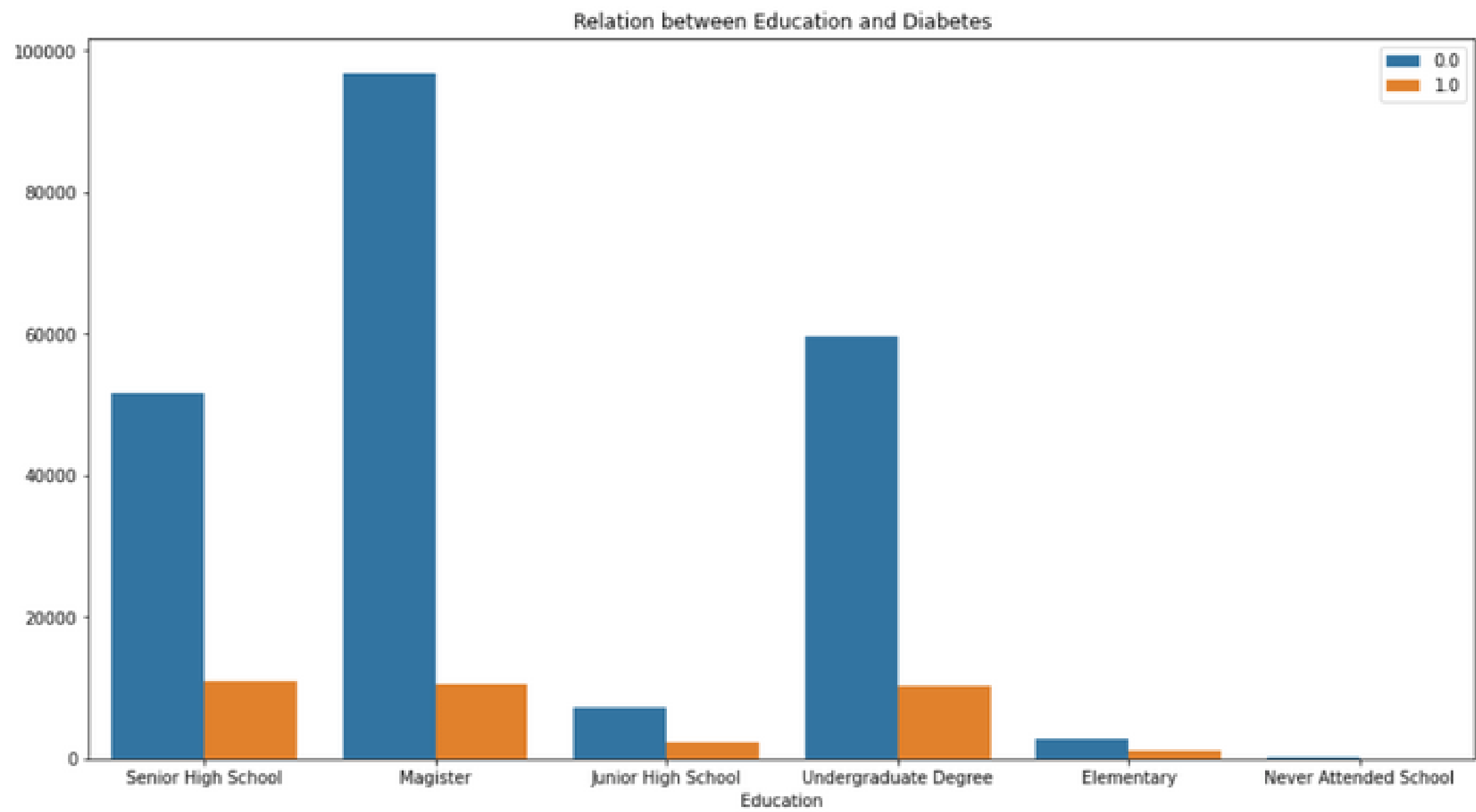
(GenHlth ,Income ) , (DiffWalk, Income) are highly correlated with each other  
=> Negative relation



# EXPLORATORY DATA ANALYSIS

## Relation between Education and diabetes

We can see that most people who have high level of education, healthy people are more than others

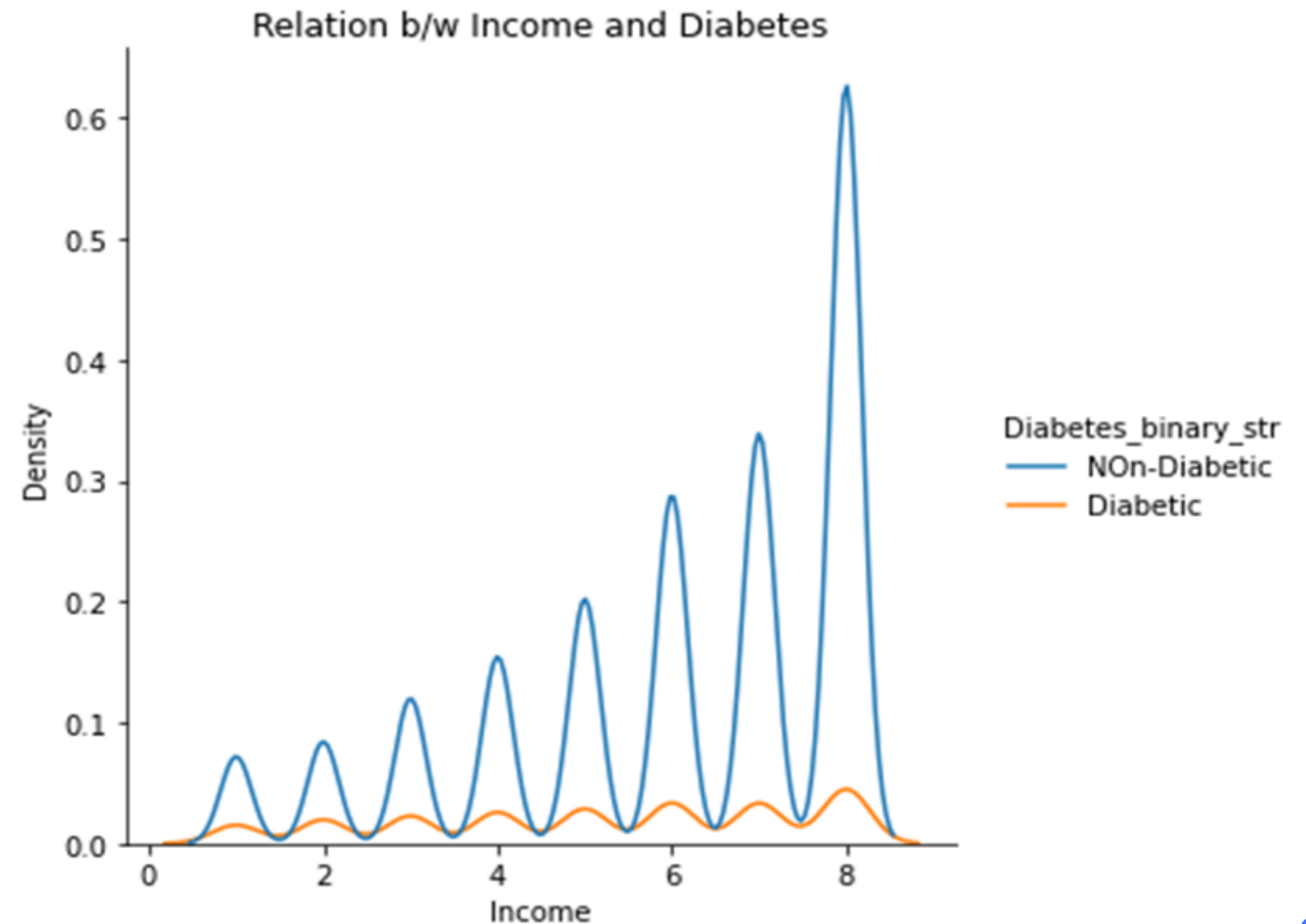




# EXPLORATORY DATA ANALYSIS

## Relation between Income and diabetes

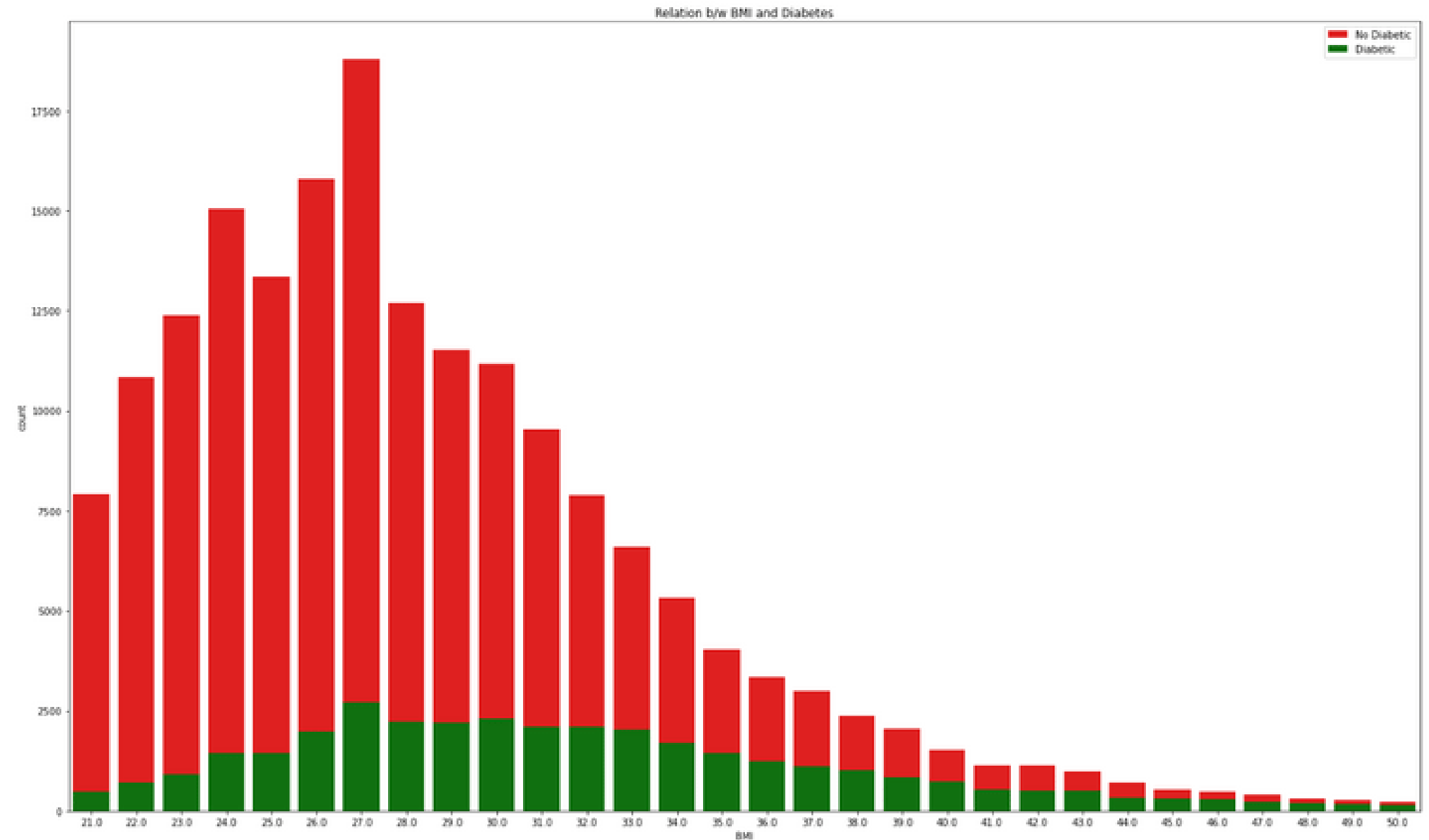
We can see that most of people have high income and in the high level of income, healthy people are more than others



# EXPLORATORY DATA ANALYSIS

## Relation between BMI and diabetes

As we can see people range between 24–33 BMI have more likely to have Diabetic

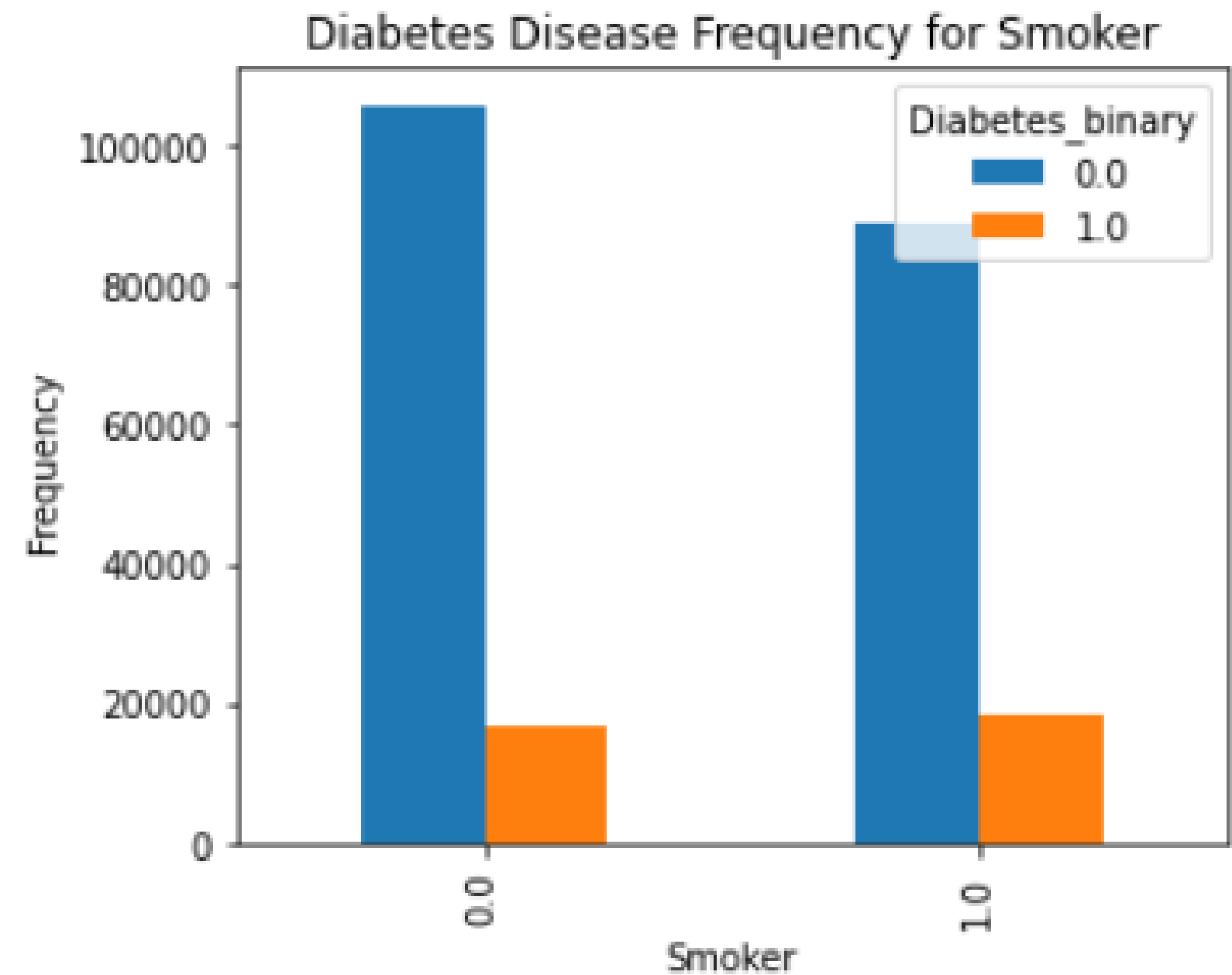


# EXPLORATORY DATA ANALYSIS

## Relation between Smoking and diabetes

According to this data, Only smoking has a minor effect on diabetes

Smoking is injurious to health



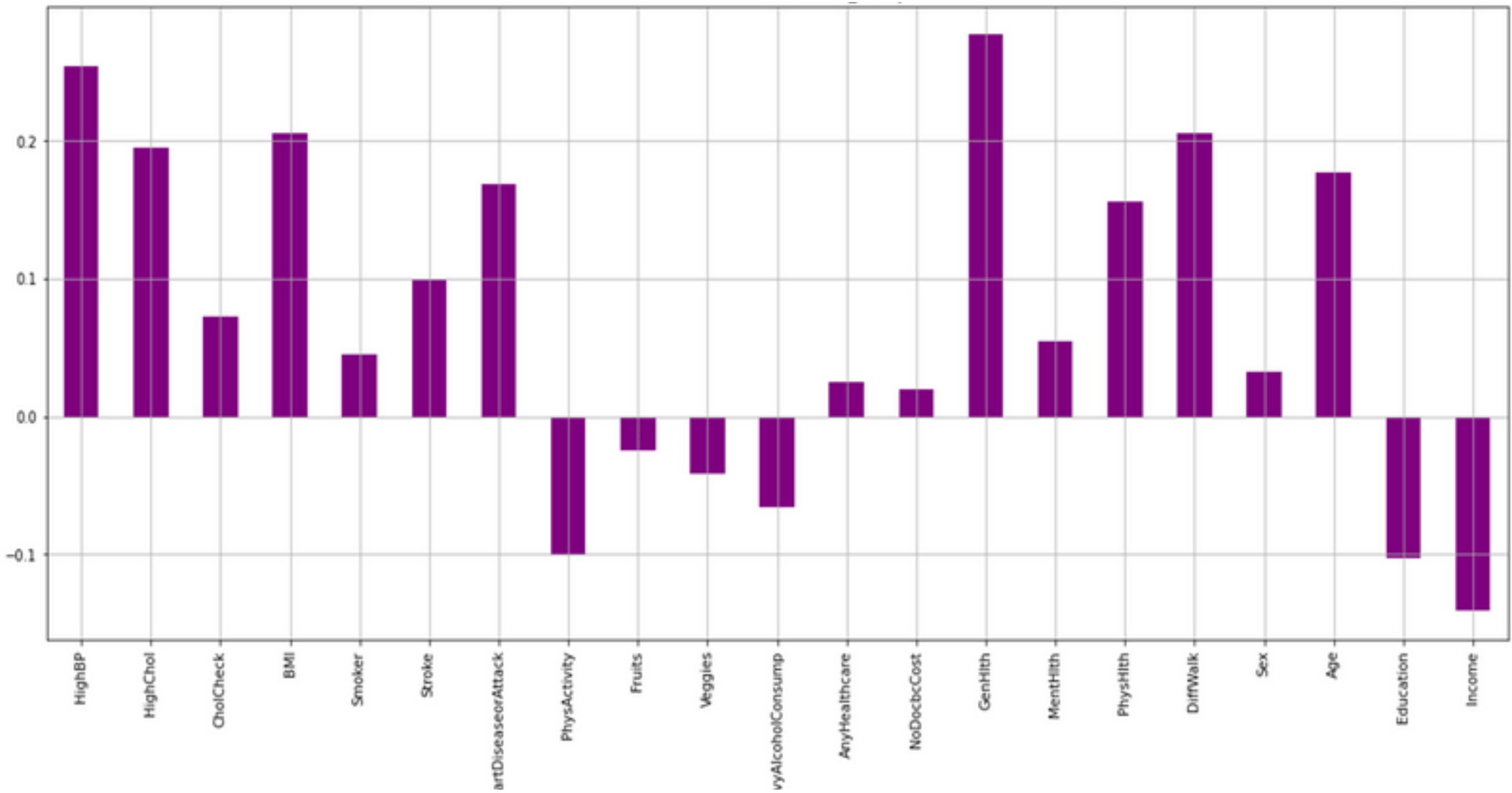
# **EXPLORATORY DATA ANALYSIS**

## **Business solution**

- **Attention to education among the different strata of society**
- **Working to raise the income and standard of living of citizens**
- **Spreading awareness among people of the need to pay attention to public health and physical health**
- **Establishing deterrent regulations for smoking and smokers and spreading awareness of its negative effects**
- **Taking care of the elderly and providing them with all their medical needs**
- **Providing means of treatment and providing permanent examinations and conducting therapeutic analyzes at close intervals for the general public**

# DATA PREPROCESSING

## Feature Extraction & Feature Selection



Fruits, AnyHealthcare, NoDocbcCost, and sex are least correlated with Diabetes binary

HighBP, HighChol, BMI, smoker, stroke, HeartDiseaseorAttack, PhysActivity, Veggies, MentHlth, HvyAlcoholconsump, GenHlth, PhysHlth, Age, Education, Income and DiffWalk have a significant correlation with Diabetes binary

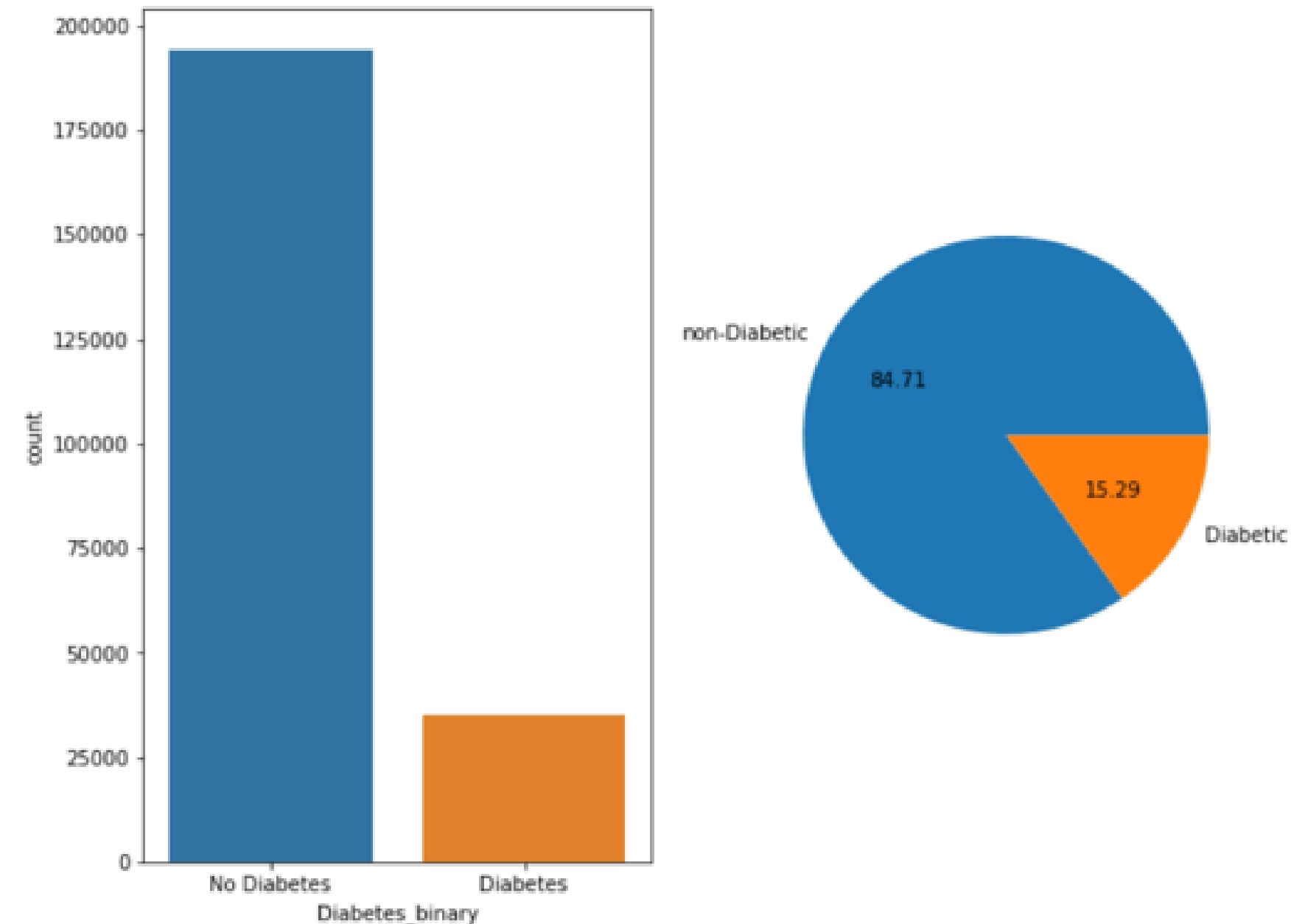
"Fruits" , "Veggies" , "Sex" , "CholCheck" , " AnyHealthcare" will not be with us

# DATA PREPROCESSING

## Handling Imbalanced

❑ **First** : we check the number of Diabetics and Nondiabetics in our target column

- 0 means non-diabetics                      0      194377
- 1 means diabetics                              1      35097





# DATA PREPROCESSING

## Handling Imbalanced

❑ **Second** :We apply under sample technic called “NearMiss”

- 0 means non-diabetics
  - 1 means diabetics
- |   |       |
|---|-------|
| 0 | 35097 |
| 1 | 35097 |

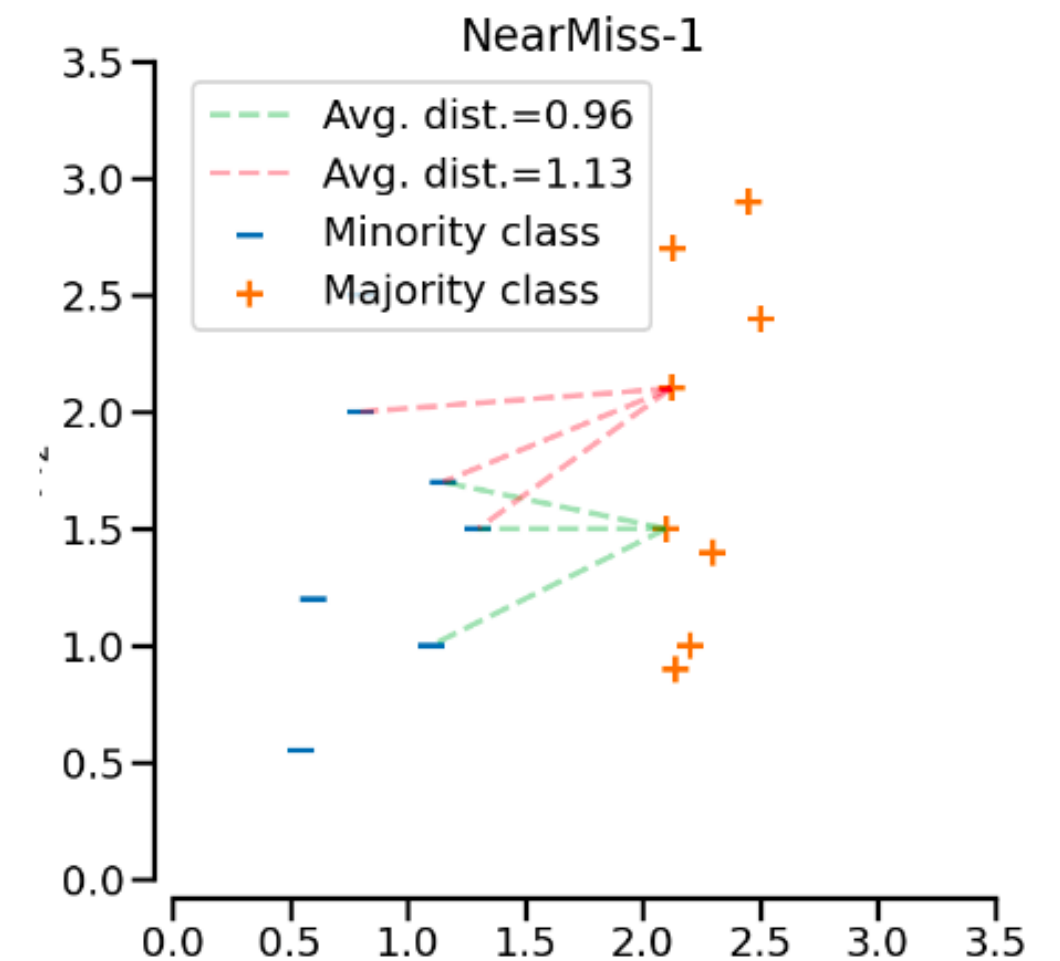
NearMiss has 3 versions and we used version 1

n\_neighbors refer to the size of the neighborhood to consider to compute the average distance to the minority point samples.

NearMiss-1 selects samples from the majority class for which the average distance of the k nearest samples of the minority class is the smallest.

NearMiss-2 selects the samples from the majority class for which the average distance to the farthest samples of the negative class is the smallest.

NearMiss-3 is a 2-step algorithm: first, for each minority sample, their m nearest-neighbors will be kept; then, the majority samples selected are the on for which the average distance to the k nearest neighbors is the largest.



# DATA MODELING

Model	Train Accuracy	Test Accuracy
Logistic Regression	0.8512	0.8472
Decision Tree	0.8657	0.8475
KNN	0.8424	0.8050
Random Forest	0.8713	0.8588
SVM	0.8687	0.8603
XGBoost	0.8770	0.8663

# MODEL EVALUATION

Logistic Regrssion

	precision	recall	f1-score	support
0	0.80	0.93	0.86	10468
1	0.92	0.76	0.83	10591

Decision Tree

	precision	recall	f1-score	support
0	0.78	0.96	0.86	10468
1	0.95	0.74	0.83	10591



# MODEL EVALUATION

K Neighbors

	precision	recall	f1-score	support
0	0.73	0.95	0.83	10468
1	0.93	0.66	0.77	10591

Random Forest

	precision	recall	f1-score	support
0	0.80	0.95	0.87	10468
1	0.94	0.77	0.85	10591



# MODEL EVALUATION

SVM

	precision	recall	f1-score	support
0	0.80	0.96	0.87	10468
1	0.95	0.76	0.85	10591

XGBoost

	precision	recall	f1-score	support
0	0.81	0.95	0.88	10468
1	0.94	0.79	0.86	10591



# CONCLUSION

**Various machine learning algorithms are explored and compared to predict diabetes to further assist the medical healthcare sector. The highest accuracy of the machine learning algorithm model is the XGBoost with 86.63% in predicting diabetes based on health indicators**







**THANK YOU!**