

# Diabetes prediction dataset EDA

## About Authors:

- Saif Ur Rehman - [email](#)
- Zeeshan Shaukat - [email](#)
- Raja Haris - (<mailto:harisraja964@gmail.com>)
- Muhammad Hamza Shakeel - [email](#)
- Ahmad Hassan - [email](#)
- Awais Haleem - [email](#)

## About Dataset:

The Diabetes prediction dataset is a collection of medical and demographic data from patients, along with their diabetes status (positive or negative). The data includes features such as age, gender, body mass index (BMI), hypertension, heart disease, smoking history, HbA1c level, and blood glucose level. This dataset can be used to build machine learning models to predict diabetes in patients based on their medical history and demographic information. This can be useful for healthcare professionals in identifying patients who may be at risk of developing diabetes and in developing personalized treatment plans. Additionally, the dataset can be used by researchers to explore the relationships between various medical and demographic factors and the likelihood of developing diabetes.

## Metadata

- **Gender:** Gender refers to the biological sex of the individual, which can have an impact on their susceptibility to diabetes.
- **Age :** Age is an important factor as diabetes is more commonly diagnosed in older adults. Age ranges from 0-80 in our dataset.
- **Hypertension :** Hypertension is a medical condition in which the blood pressure in the arteries is persistently elevated. It has values a 0.
- **Heart\_disease :** Heart disease is another medical condition that is associated with an increased risk of developing diabetes.
- **Smoking\_history :** Smoking history is also considered a risk factor for diabetes and can exacerbate the complications associated.
- **BMI:** BMI (Body Mass Index) is a measure of body fat based on weight and height. Higher BMI values are linked to a higher risk.
- **HbA1c\_level:** HbA1c (Hemoglobin A1c) level is a measure of a person's average blood sugar level over the past 2-3 months. Higher levels.
- **Blood\_glucose\_level:** Blood glucose level refers to the amount of glucose in the bloodstream at a given time. High blood glucose levels are a key.

- **Diabetes:** Diabetes is the target variable being predicted, with values of 1 indicating the presence of diabetes and 0 indicating the

## Objectives:

The objective of exploratory data analysis (EDA) of the diabetes prediction dataset is to understand the data, identify patterns and trends, and develop hypotheses about the factors that contribute to diabetes. We will use data visualization techniques and statistical tests to explore the data and to develop a deeper understanding of the relationships between the features and the target variable (diabetes). We will also identify any outliers or anomalies in the data and assess its quality.

## Task:

We intend to conduct an Exploratory Data Analysis (EDA) on the given dataset. The EDA will serve as the basis for the necessary Data Wrangling activities to be carried out for the purposes of data cleaning and normalization. During the coding process, we will document our observations. Ultimately, we will produce a summary and draw conclusions from our findings

# 1. Importing Libraries

## 1.1 We will use the following libraries

- **Pandas:** Data manipulation and analysis library.
- **Numpy:** Numerical computing library.
- **Matplotlib:** Data visualization library.
- **Seaborn:** Statistical data visualization library.
- **Plotly:** Interactive Data visualization library.

```
# use matplotlib inline to plot the graphs within the jupyter notebook
%matplotlib inline
```

```
# ignore warnings
import warnings
warnings.filterwarnings('ignore')
```

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

```
# set plt image size globally
plt.rcParams['figure.figsize'] = (15,6)
```

## 2. Loading, Transformation and Exploration of dataset

### 2.1 Loading the dataset from a CSV file.

```
df = pd.read_csv("./hemogloba_data.csv")
```

- let's see the first 5 rows of the dataset to get a feel of how the data looks like

```
df.head()
```

	gender	age	hypertension	heart_disease	smoking_history	bmi	\
0	Female	80.0	0	1	never	25.19	
1	Female	54.0	0	0	No Info	27.32	
2	Male	28.0	0	0	never	27.32	
3	Female	36.0	0	0	current	23.45	
4	Male	76.0	1	1	current	20.14	

	HbA1c_level	blood_glucose_level	diabetes
0	6.6	140	0
1	6.6	80	0
2	5.7	158	0
3	5.0	155	0
4	4.8	155	0

- let's check the number of rows and columns in the dataset

```
print(f"The number of rows are {df.shape[0]}, and columns are {df.shape[1]}")
```

The number of rows are 100000, and columns are 9

- Get a sneak peek of your data

```
df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 100000 entries, 0 to 99999
Data columns (total 9 columns):
#   Column                Non-Null Count  Dtype
---  -
0   gender                100000 non-null object
1   age                  100000 non-null float64
2   hypertension          100000 non-null int64
3   heart_disease         100000 non-null int64
4   smoking_history       100000 non-null object
5   bmi                  100000 non-null float64
6   HbA1c_level           100000 non-null float64
7   blood_glucose_level   100000 non-null int64
```

```
8    diabetes      100000 non-null  int64
dtypes: float64(3), int64(4), object(2)
memory usage: 6.9+ MB
```

## Observations:

- We have 9 columns; 3 columns of float, 4 columns of int and 2 columns of object dtype.
- There is no null values in any of the columns.
- Memory usage is 6.9+ MB

## 2.2 Missing Values

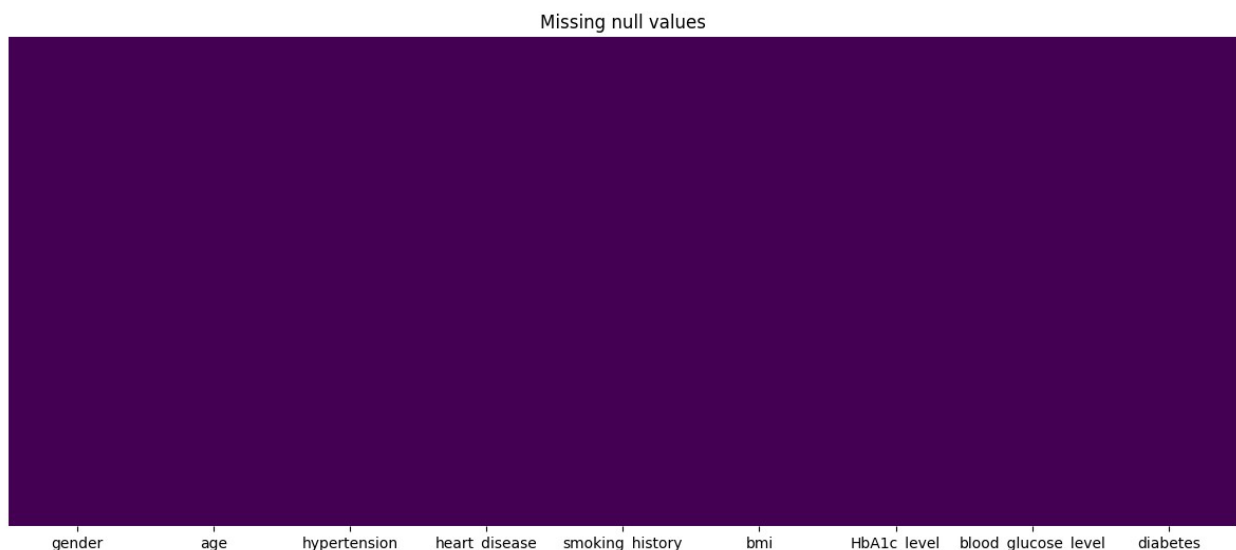
- verify missing values

```
df.isnull().sum()
```

```
gender      0
age         0
hypertension 0
heart_disease 0
smoking_history 0
bmi         0
HbA1c_level 0
blood_glucose_level 0
diabetes    0
dtype: int64
```

let's plot the missing values using heatmap

```
sns.heatmap(df.isnull(),yticklabels = False, cbar = False , cmap =
'viridis')
plt.title("Missing null values")
plt.show()
```



## 2.3 Duplicate values

let's check the duplicate values in the dataset

```
df.duplicated().sum()
```

```
3854
```

```
print(f'We have {df.duplicated().sum()} duplicated rows in the dataset  
\nWhich are {round(df.duplicated().sum() / len(df) * 100,2)}% of the  
dataset')
```

```
We have 3854 duplicated rows in the dataset  
Which are 3.85% of the dataset
```

This is minute observations and we can drop the duplicated rows to maintain the integrity of the dataset.

```
df.drop_duplicates(inplace=True)
```

## 2.3 Summary Statistics of numerical columns

```
df.describe()
```

	age	hypertension	heart_disease	bmi
HbA1c_level \				
count	96146.000000	96146.000000	96146.000000	96146.000000
mean	41.794326	0.077601	0.040803	27.321461
std	5.532609			
std	22.462948	0.267544	0.197833	6.767716
min	0.080000	0.000000	0.000000	10.010000
25%	24.000000	0.000000	0.000000	23.400000
50%	43.000000	0.000000	0.000000	27.320000
75%	59.000000	0.000000	0.000000	29.860000
max	80.000000	1.000000	1.000000	95.690000

	blood_glucose_level	diabetes
count	96146.000000	96146.000000
mean	138.218231	0.088220
std	40.909771	0.283616
min	80.000000	0.000000
25%	100.000000	0.000000

50%	140.000000	0.000000
75%	159.000000	0.000000
max	300.000000	1.000000

## 2.4 unique values and value counts in non-numerical columns

- let's check the nunique values and counts in each non-numerical column

```
# take non_numeric columns
non_numeric_columns =
df.select_dtypes(exclude=np.number).columns.tolist()
for col in non_numeric_columns:
    print(f"{col} has following {df[col].nunique()} unique values: \
n{df[col].unique()}\n")
    print(f"Value counts for {col} are: \n{df[col].value_counts()}\n")
```

gender has following 3 unique values:  
['Female' 'Male' 'Other']

Value counts for gender are:

```
gender
Female    56161
Male      39967
Other         18
Name: count, dtype: int64
```

smoking\_history has following 6 unique values:  
['never' 'No Info' 'current' 'former' 'ever' 'not current']

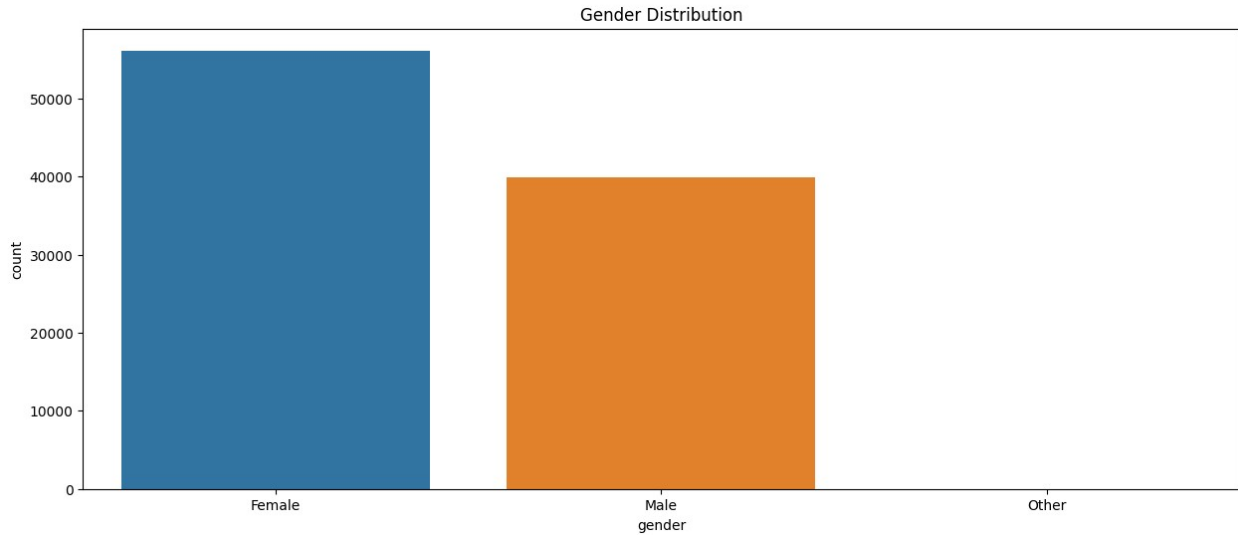
Value counts for smoking\_history are:

```
smoking_history
never      34398
No Info    32887
former      9299
current     9197
not current 6367
ever        3998
Name: count, dtype: int64
```

## 2.4 Distribution of the data

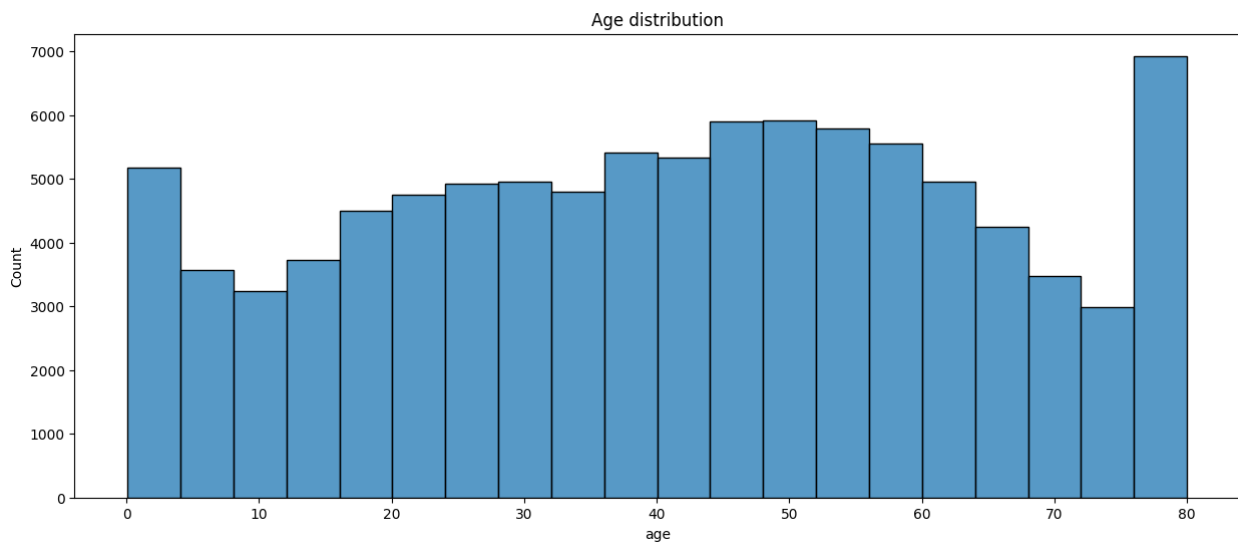
- Gender

```
sns.countplot(df, x='gender')
plt.title("Gender Distribution")
plt.show()
```



- age

```
sns.histplot(df, x='age', bins=20)
plt.title("Age distribution")
plt.show()
```



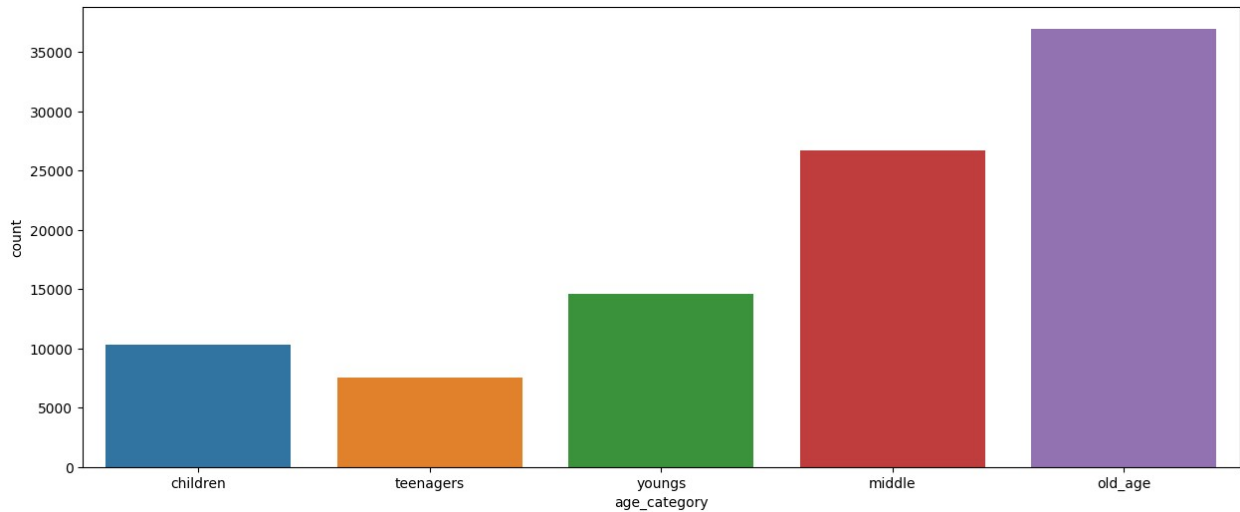
We have almost uniform distribution of age in the dataset. We have all entries from every age group.

- Making new column with age categories from age column

```
bins = [0,10,18,30,50,80]
labels = ['children', 'teenagers', 'youngs', 'middle','old_age']
df['age_category'] = pd.cut(df['age'],bins = bins, labels = labels)

sns.countplot(x='age_category', data=df)

<Axes: xlabel='age_category', ylabel='count'>
```



we have more old age people in our dataset

- let's see which people are more likely to have diabetes

```
df.groupby('age_category')['diabetes'].value_counts()
```

age_category	diabetes	count
children	0	10297
	1	29
teenagers	0	7431
	1	63
youngs	0	14389
	1	209
middle	0	25192
	1	1547
old_age	0	30355
	1	6634

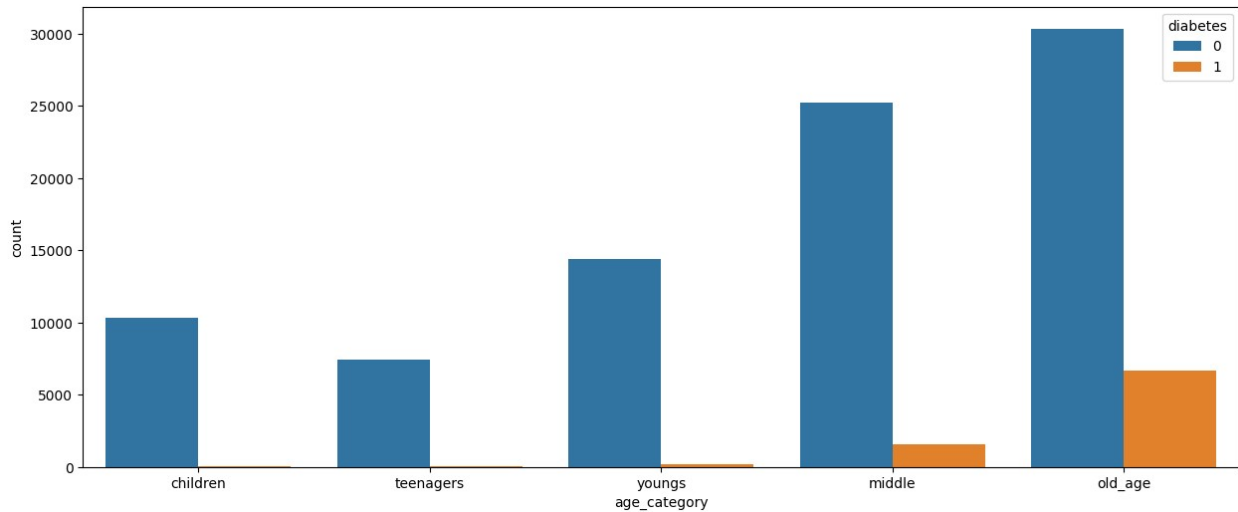
Name: count, dtype: int64

*# make a plot to show diabetes distribution in each age category*

```
sns.countplot(x='age_category', hue='diabetes', data=df)
```

```
<Axes: xlabel='age_category', ylabel='count'>
```





Old age group is more in dataset and also have more diabetes patients.

```
# check Percentage of diabetes in each age category
df.groupby('age_category')
['diabetes'].value_counts(normalize=True).mul(100).round(2).astype(str)
) + '%'
```

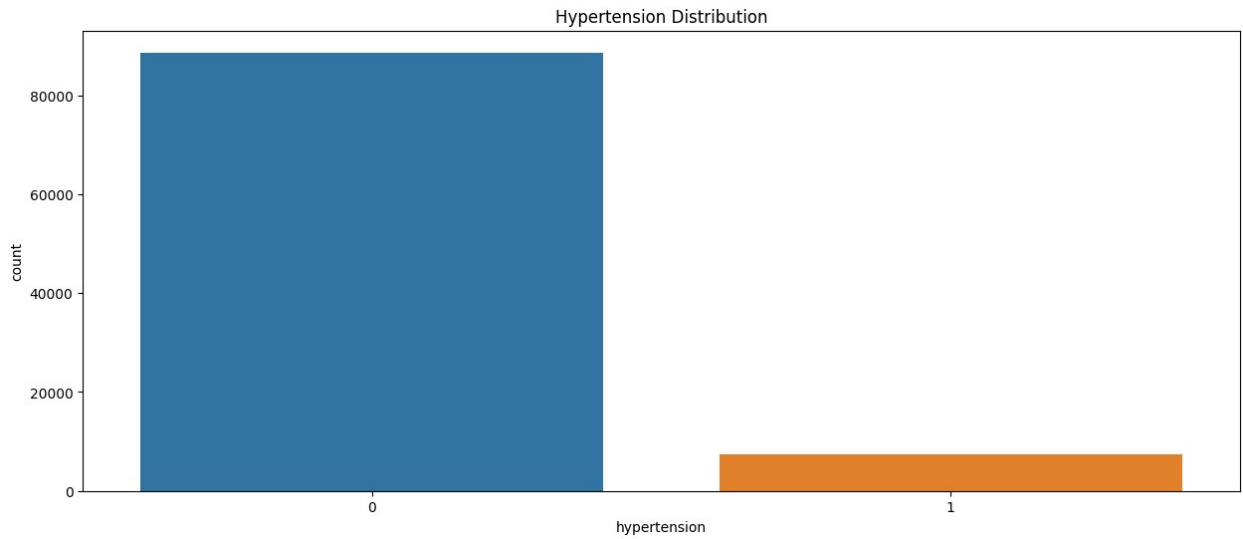
age_category	diabetes	percentage
children	0	99.72%
	1	0.28%
teenagers	0	99.16%
	1	0.84%
youngs	0	98.57%
	1	1.43%
middle	0	94.21%
	1	5.79%
old_age	0	82.06%
	1	17.94%

Name: proportion, dtype: object

Old age group has about 19% of diabetes patients

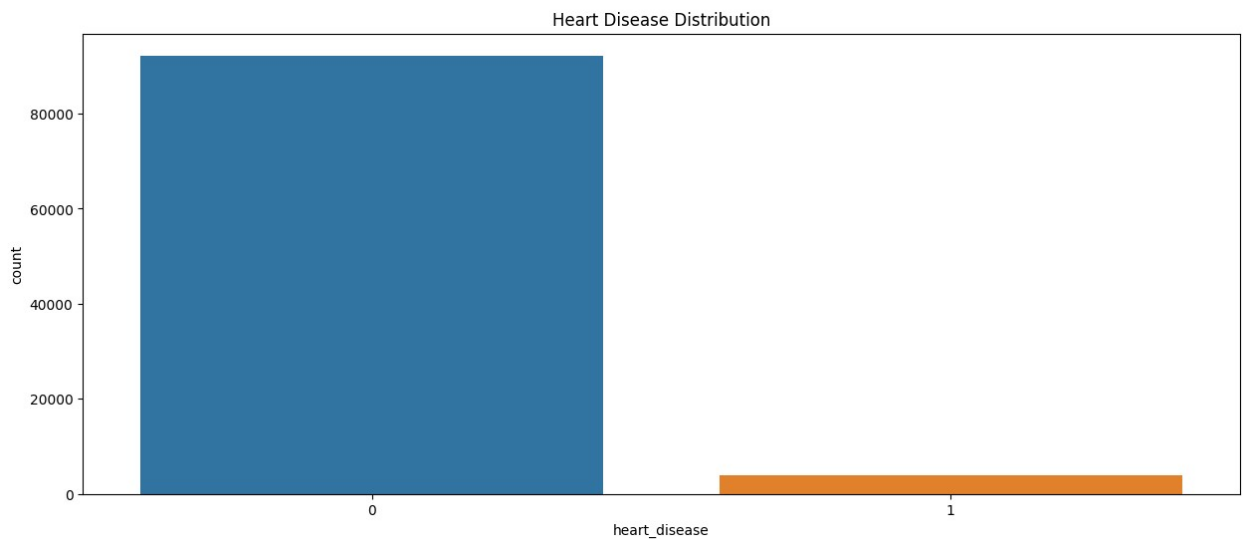
- hypertension

```
sns.countplot(df, x='hypertension')
plt.title("Hypertension Distribution")
plt.show()
```



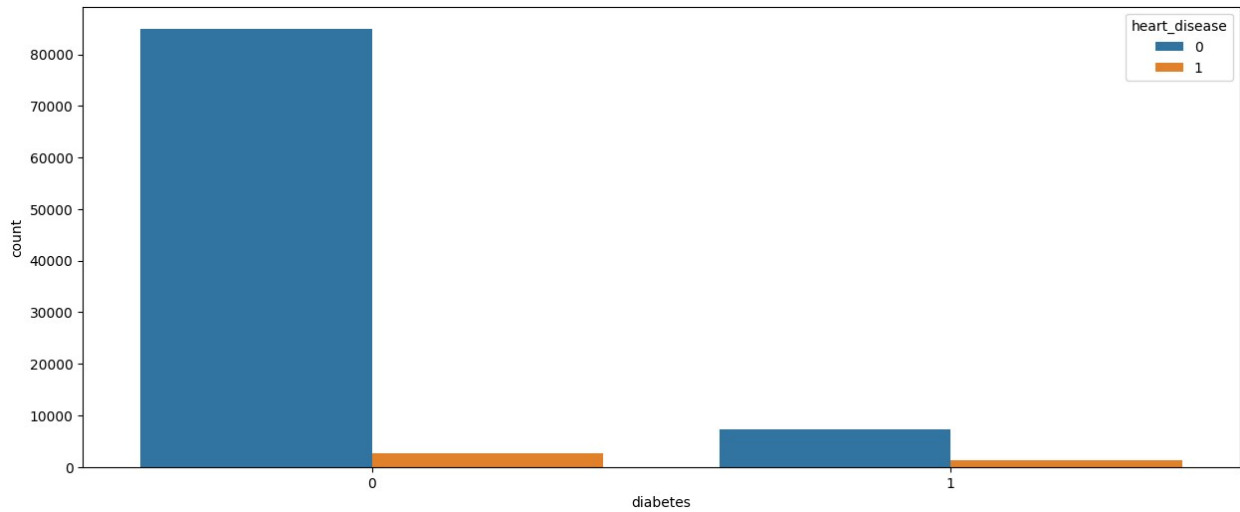
- heart\_disease

```
sns.countplot(df, x='heart_disease')  
plt.title("Heart Disease Distribution")  
plt.show()
```



more people have no heart disease. very few people have heart disease

```
# Is heart disease and diabetes related?  
sns.countplot(x='diabetes', hue='heart_disease', data=df)  
<Axes: xlabel='diabetes', ylabel='count'>
```



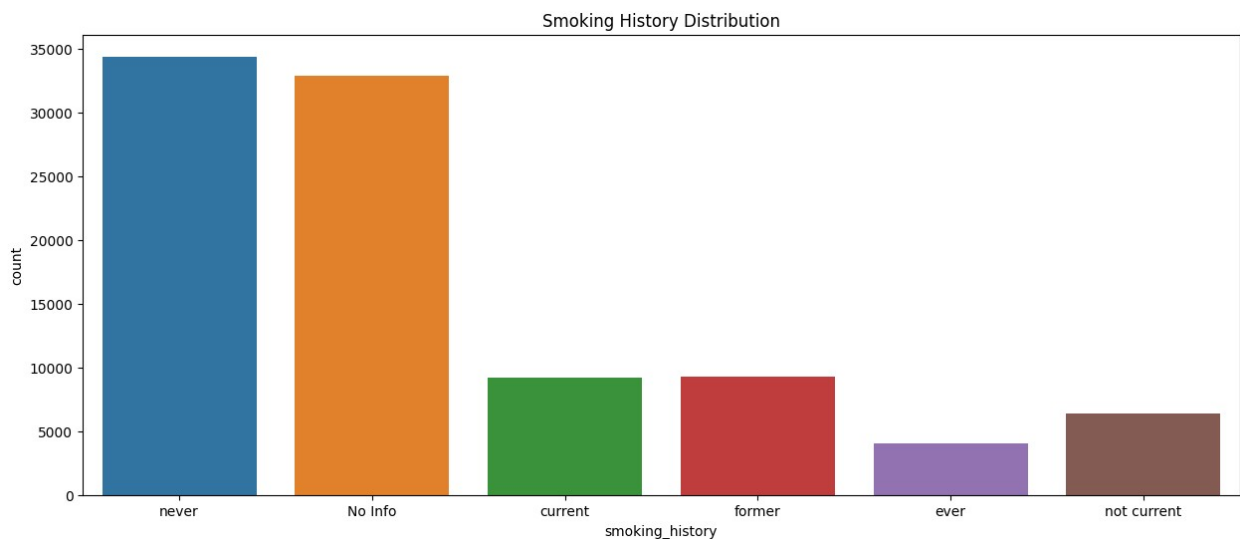
There is no relation between heart disease and diabetes

```
df[['heart_disease', 'diabetes']].corr()
```

	heart_disease	diabetes
heart_disease	1.000000	0.170711
diabetes	0.170711	1.000000

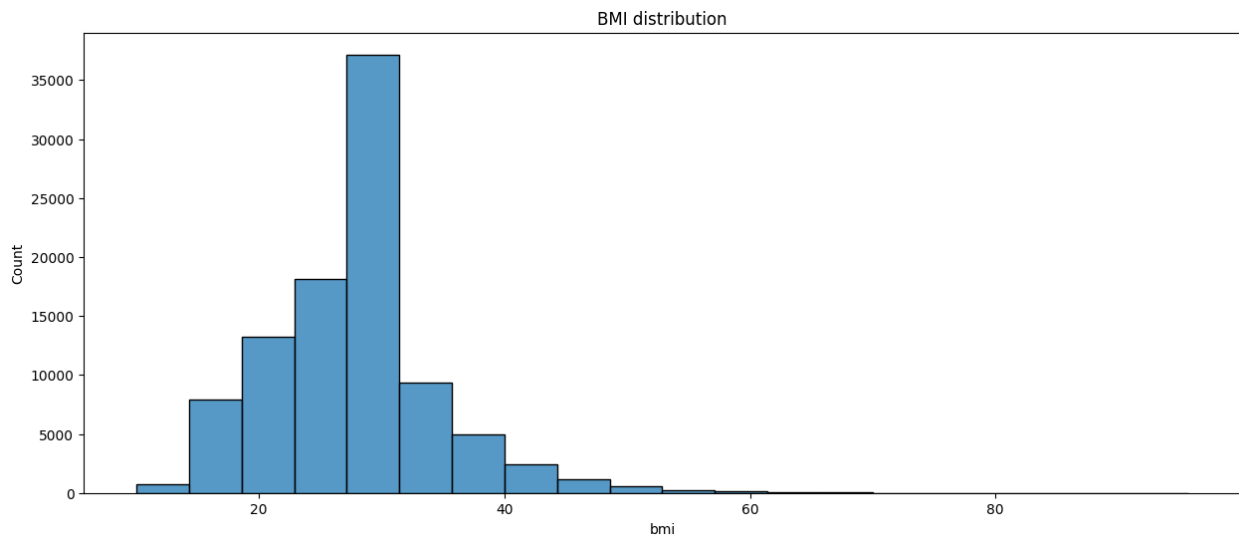
- smoking\_history

```
sns.countplot(df, x='smoking_history')
plt.title("Smoking History Distribution")
plt.show()
```

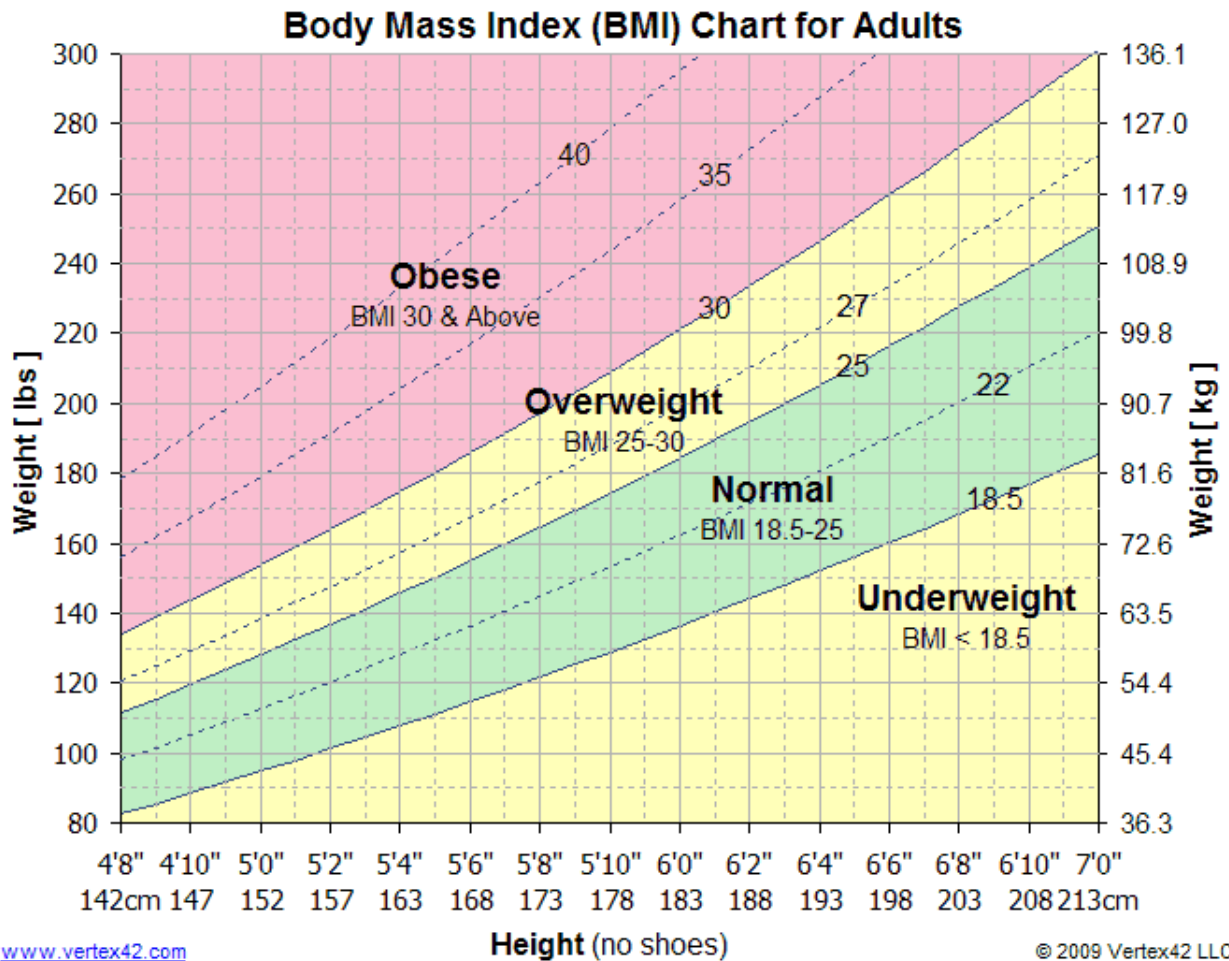


- bmi

```
sns.histplot(df, x='bmi', bins=20)  
plt.title("BMI distribution")  
plt.show()
```



BMI Distribution is slightly right skewed. we have potential outliers in the dataset.

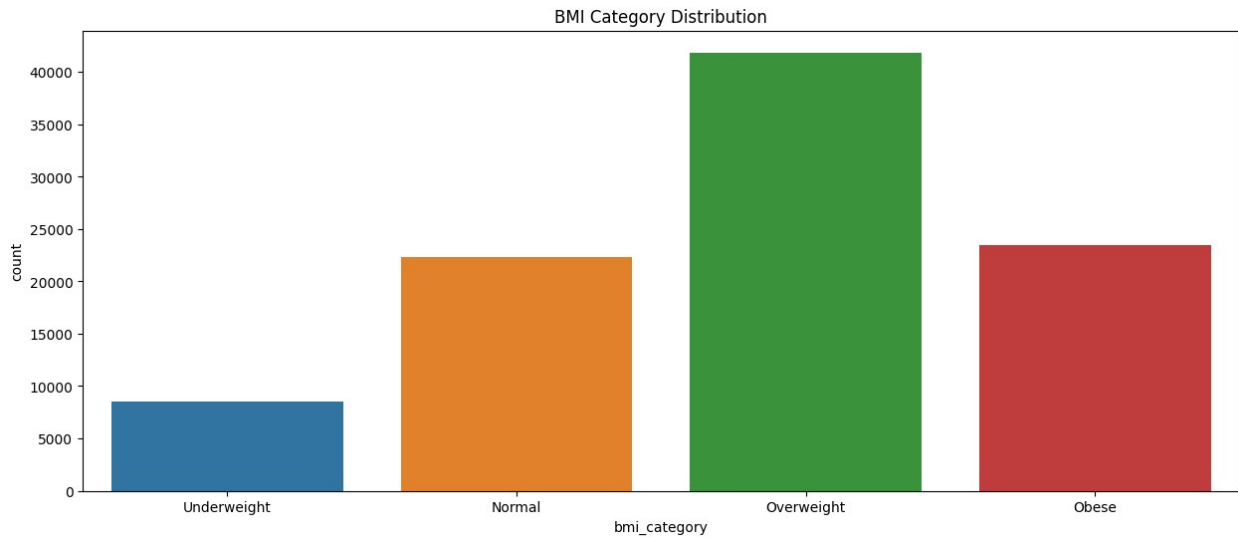


[www.vertex42.com](http://www.vertex42.com)  
bmi\_chart

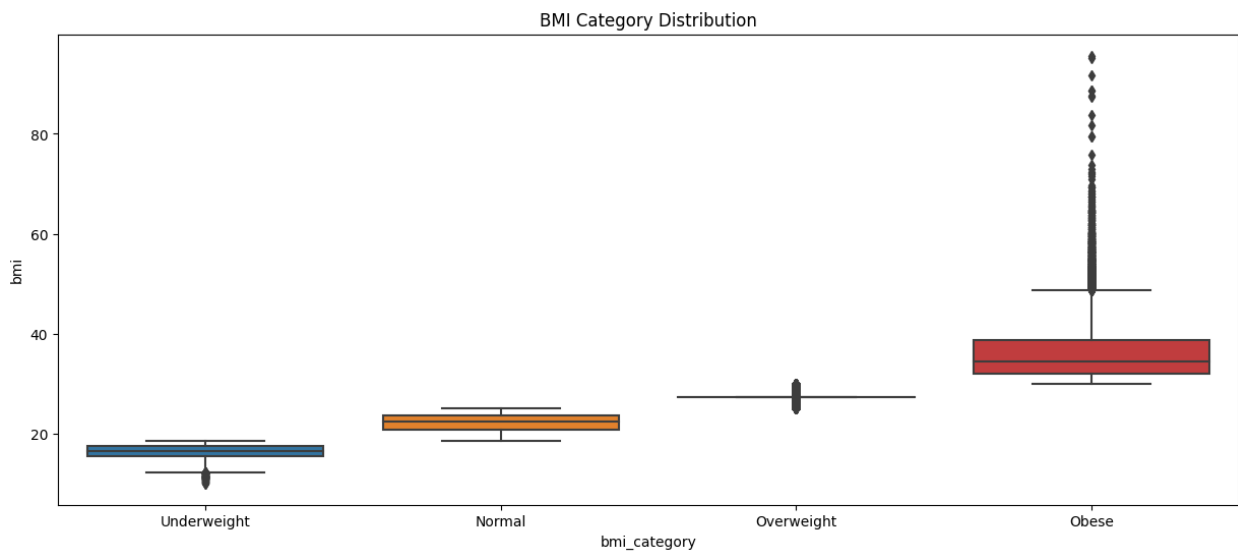
- HbA1c\_level
- Making a new column for BMI category

```
# binn bime into 4 categories
df['bmi_category'] = pd.cut(df['bmi'], bins=[0, 18.5, 25, 30, 100],
labels=['Underweight', 'Normal', 'Overweight', 'Obese'])

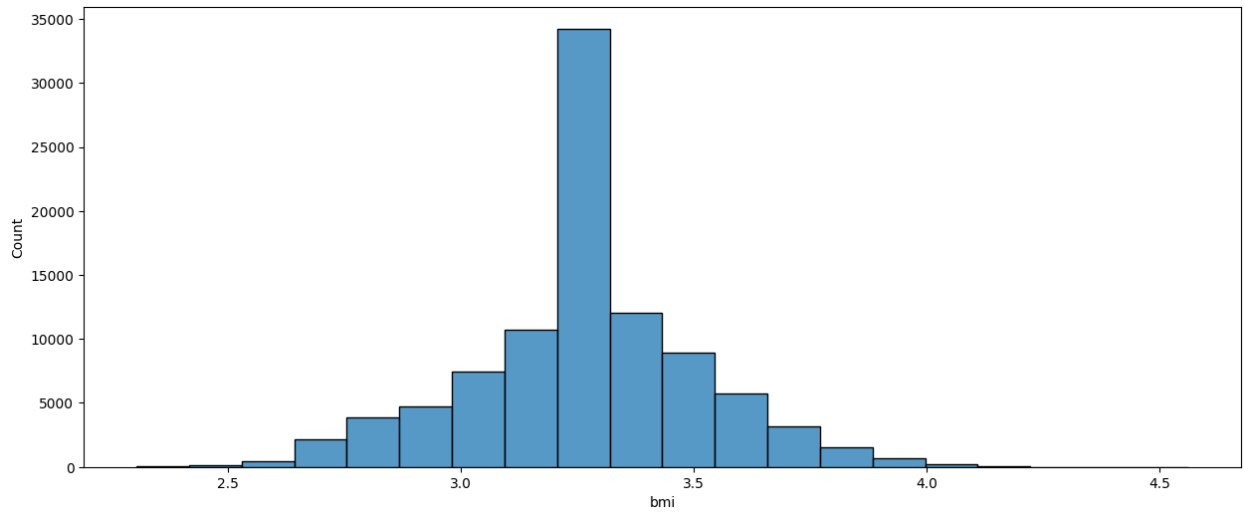
# lets see the distribution of bmi categories
sns.countplot(df, x='bmi_category')
plt.title("BMI Category Distribution")
plt.show()
```



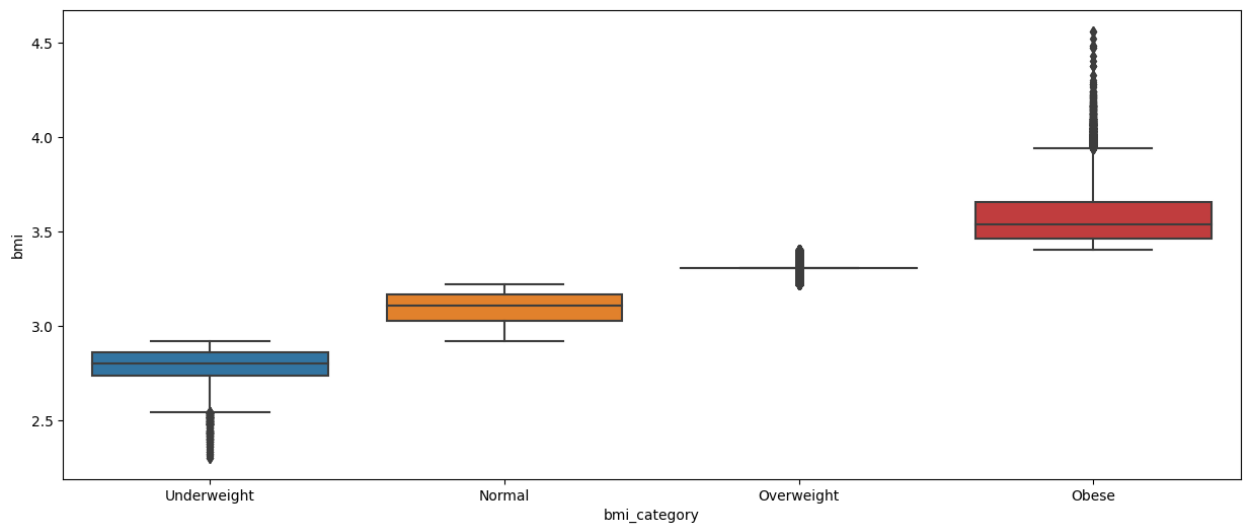
```
# lets see the outliers in bmi_category
sns.boxplot(df, x='bmi_category', y='bmi')
plt.title("BMI Category Distribution")
Text(0.5, 1.0, 'BMI Category Distribution')
```



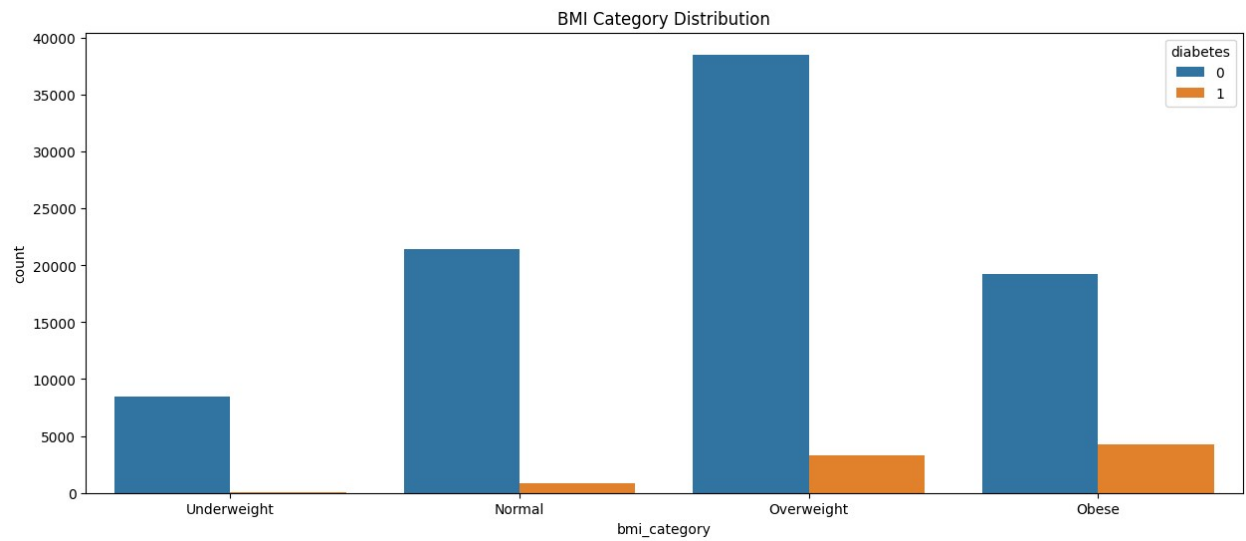
```
# tranforming bmi in log scale and plotting the distribution
sns.histplot(np.log(df['bmi']), bins=20)
<Axes: xlabel='bmi', ylabel='Count'>
```



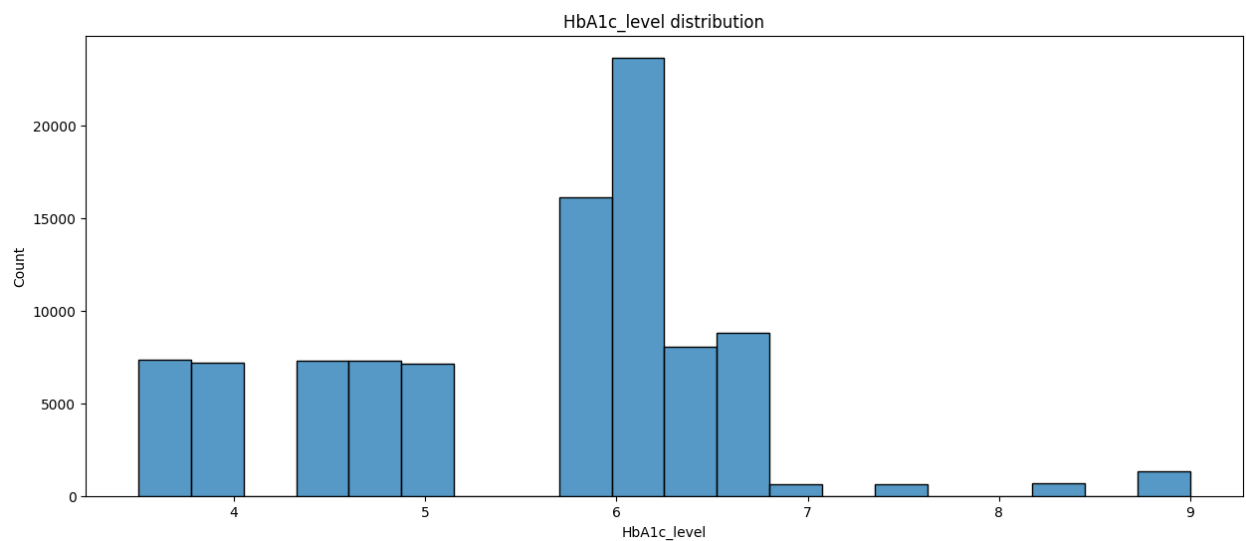
```
sns.boxplot(df, x='bmi_category', y=np.log(df['bmi']))  
<Axes: xlabel='bmi_category', ylabel='bmi'>
```



```
# let's see diabetes in every bmi category  
sns.countplot(df, x='bmi_category', hue='diabetes')  
plt.title("BMI Category Distribution")  
plt.show()
```



```
sns.histplot(df, x='HbA1c_level', bins=20)  
plt.title("HbA1c_level distribution")  
plt.show()
```





# HbA1c(%)

# ELO

Optimal  
←5.7%

Elevated  
5.7–6.4

High  
6.4→



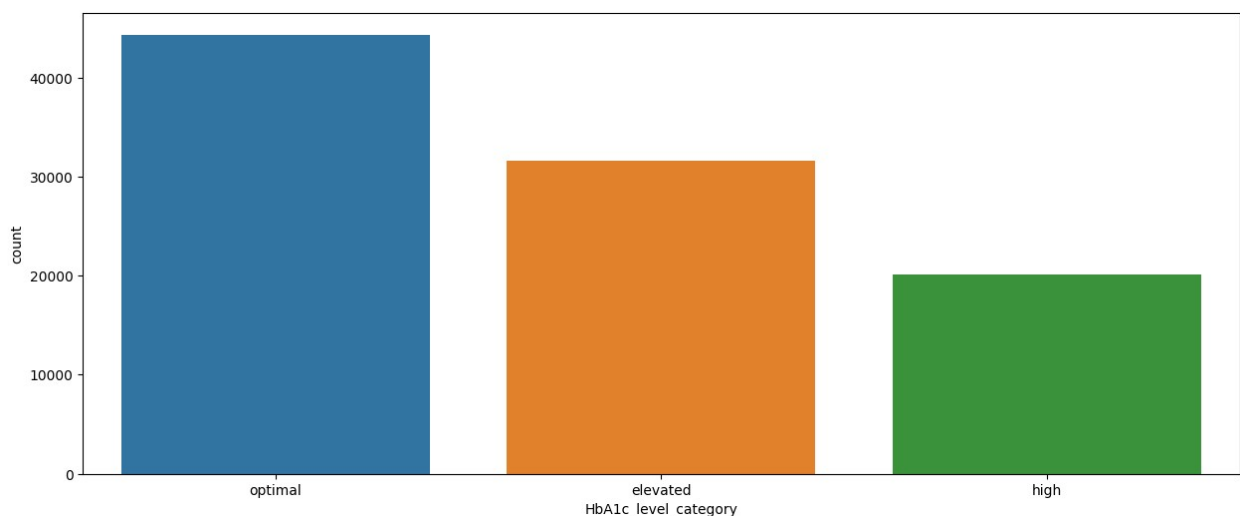
WWW.ELO.HEALTH

HB1c\_level\_chart

```
# making HBa1c level in 3 categories
bins = [0, 5.7, 6.4, 100]
labels = ['optimal', 'elevated', 'high']
df['HbA1c_level_category'] = pd.cut(df['HbA1c_level'], bins=bins,
labels=labels)

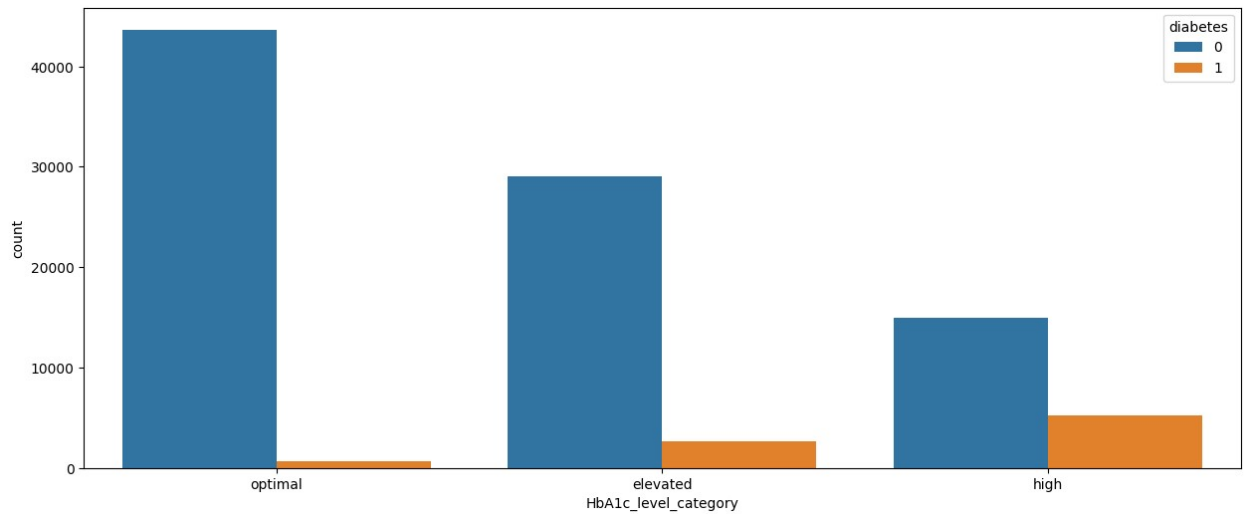
sns.countplot(df, x='HbA1c_level_category')

<Axes: xlabel='HbA1c_level_category', ylabel='count'>
```

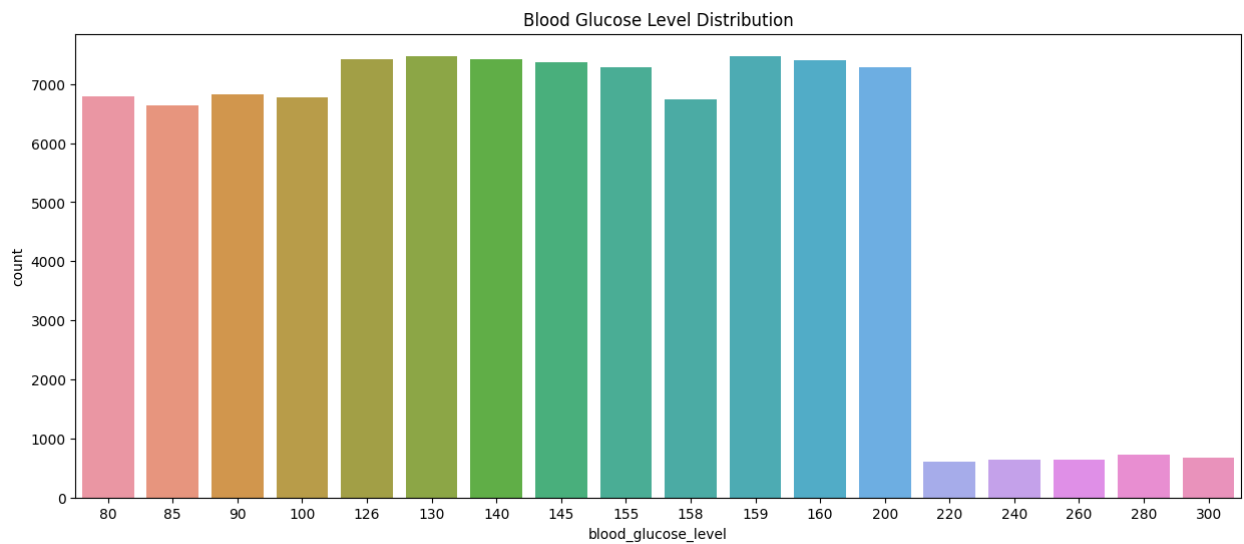


```
sns.countplot(df, x='HbA1c_level_category', hue='diabetes')
```

```
<Axes: xlabel='HbA1c_level_category', ylabel='count'>
```



```
sns.countplot(df, x='blood_glucose_level')  
plt.title("Blood Glucose Level Distribution")  
plt.show()
```



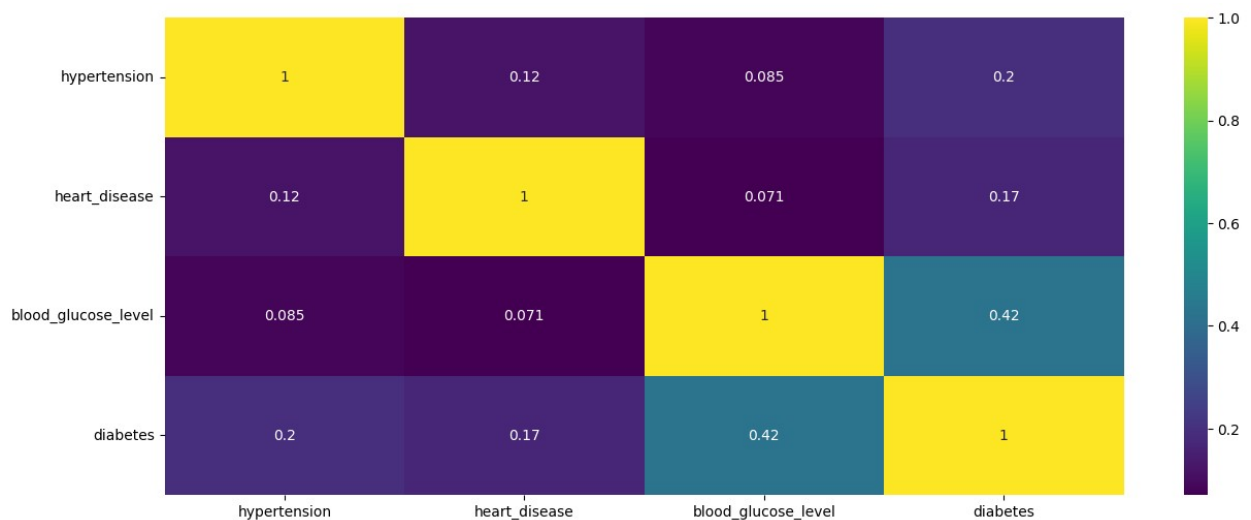
Blood Glucose Chart			
Mg/DL	Fasting	After Eating	2-3 Hours After Eating
Normal	80-100	170-200	120-140
Impaired Glucose	101-125	190-230	140-160
Diabetic	126+	220-300	200+

blood\_glocose\_chart

We have patients with blood glucose level above 200.

```
# bllood Glucose lever, blood desies, hypertenstion and diabetes
distribution looks same, lets see the correlation
sns.heatmap(df[["hypertension", "heart_disease",
"blood_glucose_level", "diabetes"]].corr(), annot=True,
cmap='viridis')
```

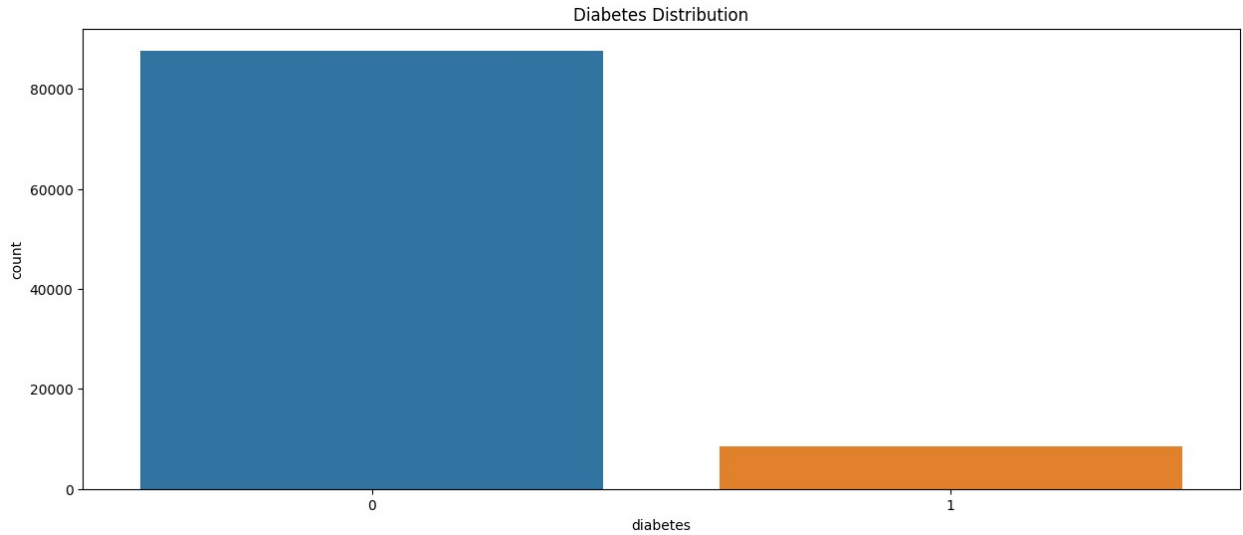
<Axes: >



Diabetes and blood\_glucose\_level are positively correlated. Other than that, there is positive correlation between other variables but it's not too much strong.

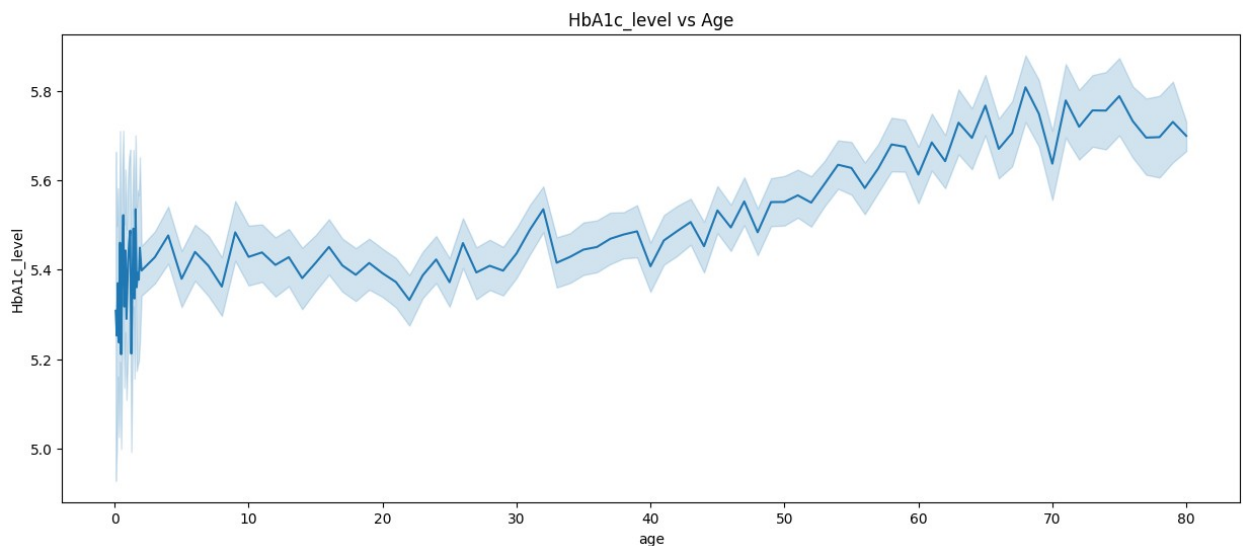
- diabetes

```
sns.countplot(df, x='diabetes')
plt.title("Diabetes Distribution")
plt.show()
```



## Trend between age and HbA1c\_level

```
sns.lineplot(df, x='age', y='HbA1c_level')
plt.title("HbA1c_level vs Age")
plt.show()
```



- Draw sunburst on df columns gender, diabetes\_categorical, smoking\_history

```
fig = px.sunburst(
    df,
    path=['gender',
          'age_catogry', 'hypertension_categorical', 'heart_disease_categorical']
    ,
    values= 'diabetes',
```

```

        color='gender',
    )

# Show the Sunburst chart
fig.show()

{"config":{"plotlyServerURL":"https://plot.ly"},"data":
[{"branchvalues":"total","customdata":[["Female"],["Other"],["Male"],
["Female"],["Other"],["Other"],["Female"],["Female"],["Other"],[null],
["Male"],["Male"],[null],["Male"],["Male"],[null],[null],[null],
[null],["Female"],["Male"],[null],["Female"],["Female"],["Other"],
["Male"],["Female"],["Other"],["Other"],["Male"],["Female"],["Other"],
["Female"],["Other"],["Male"],[null],["Female"],[null],["Male"],
[null],["Female"],[null],[null],[null],[null],["Female"],["Male"],
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["Male"],[null],["Other"],[null],["Male"],["Female"],["Female"],
["Male"],["Female"],["Female"],["Female"],["Female"],["Male"],
["Other"],[null],["Male"],["Male"],["Female"],[null],[null],[null],
[null],["Female"],["Male"],[null],["Female"],[null],["Male"],["Male"],
[null],["Male"],["Female"],["Male"],["Female"],["Female"],[null],
["Male"],["Male"],["Other"],["Other"]],"domain":{"x":[0,1],"y":
[0,1]},"hvertemplate":"labels=%{label}<br>diabetes=%
{value}<br>parent=%{parent}<br>id=%{id}<br>gender=%
{customdata[0]}<extra></extra>","ids":["Female/children/0/0","Other/
old_age/0","Male/old_age/0","Female/old_age/0","Other/middle/
0","Other/youngs/0","Female/youngs/0","Female/teenagers/0","Other/
children/0","Female/children/1","Male/children/0","Male/old_age/
1/1","Other/middle/1/1","Male","Male/youngs/1/1","Female/youngs/
1/1","Male/teenagers/1/1","Other/children/1/1","Male/children/
1/1","Female/children/0","Male/children/1","Other/children/
1","Female/teenagers/1","Female","Other/old_age","Male/
old_age","Female/old_age","Other/middle","Other/youngs","Male/
youngs","Female/youngs","Other/teenagers","Female/teenagers","Other/
children","Male/children","Other/old_age/1","Female/old_age/
1","Other/middle/1","Male/middle/1","Other/youngs/1","Female/youngs/
1","Other/teenagers/1","Female/children/1/1","Other/old_age/
0/1","Other/youngs/1/1","Female/old_age/0/1","Male/children/
1/0","Male/old_age/0/0","Male/old_age/0/1","Other/youngs/
1/0","Female/middle/1/0","Male/middle/1/0","Other/middle/1/0","Other/
middle/0/0","Male/old_age/1/0","Male/middle/0/0","Female/middle/
0/0","Female/children/0/1","Male/children/0/1","Other/children/
0/1","Other/youngs/0/0","Male/youngs/0/0","Male/middle/0/1","Other/
children/0/0","Female/middle/0/1","Male/teenagers/0/0","Female/
teenagers/0/1","Female/youngs/0/1","Male/youngs/0/1","Male/teenagers/
0/1","Other/teenagers/0/0","Other/children/1/0","Male/
middle","Female/old_age/0/0","Female/children","Male/
teenagers","Female/middle","Female/teenagers/0/0","Female/middle/
1","Female/youngs/0/0","Male/old_age/1","Other/old_age/0/0","Female/

```

```
children/1/0","Male/children/0/0","Male/youngs/1/0","Female/
teenagers/1/0","Other/middle/0/1","Female/teenagers/1/1","Other/
teenagers/1/1","Other/youngs/0/1","Female/middle/1/1","Male/middle/
1/1","Other/teenagers/0/1","Female/old_age/1/1","Other/old_age/
1/1","Male/youngs/1","Male/teenagers/0","Other/old_age/1/0","Male/
youngs/0","Female/middle/0","Male/middle/0","Female/old_age/
1/0","Female/youngs/1/0","Other/teenagers/1/0","Male/teenagers/
1","Male/teenagers/1/0","Other/teenagers/0","Other"},"labels":
["0","0","0","0","0","0","0","0","0","1","0","1","1","Male","1","1","1
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## Insights

- The observation on basis of gender is as follows:

Female: 4447

- Old\_age : 3464 Diabetic Patients
- Middle : 812 Diabetic Patients
- Youngs : 119 Diabetic Patients
- teenagers : 35 Diabetic Patients

Male: 4035

- Old\_age : 3170 Diabetic Patients
- Middle : 735 Diabetic Patients
- Youngs : 90 Diabetic Patients
- teenagers : 28 Diabetic Patients

Others: 0

Q1. Comparatively who has more Diabetes?

A. Based on given data females has more diabetes than male.

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*# Show the Sunburst chart*

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## Q2. Comparatively who has more heart\_disease?

A. Based on given data male has more heart issues than female.

- Male: 2367
- Female: 1556
- Others: 0

```

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```

*# Show the Sunburst chart*  
fig.show()

```

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### Q3. Comparatively who has more hypertension?

A. Based on given data female has more heart issues than male.

- Female: 4179
- Male: 3282
- Others: 0

```
fig = px.sunburst(
    df,

    path=['hypertension_categorical', 'heart_disease_categorical', 'gender',
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    values= 'diabetes',
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)
```

```
# Show the Sunburst chart
```

```
fig.show()
```

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```

```

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```

Q4. People with Hypertension and Heart-disease on the basis of gender, who got more diabetes?

A. Based on Data people with Hypertension and Hear\_disease, Male has more Diabetes.

- Male : 194
- Female : 164

Q5. People with no Hypertension and no Heart-disease on the basis of gender, who got more diabetes?

A. Based on Data people with Hypertension and Hear\_disease, female has more Diabetes.

- Male 2518
- Female 2969

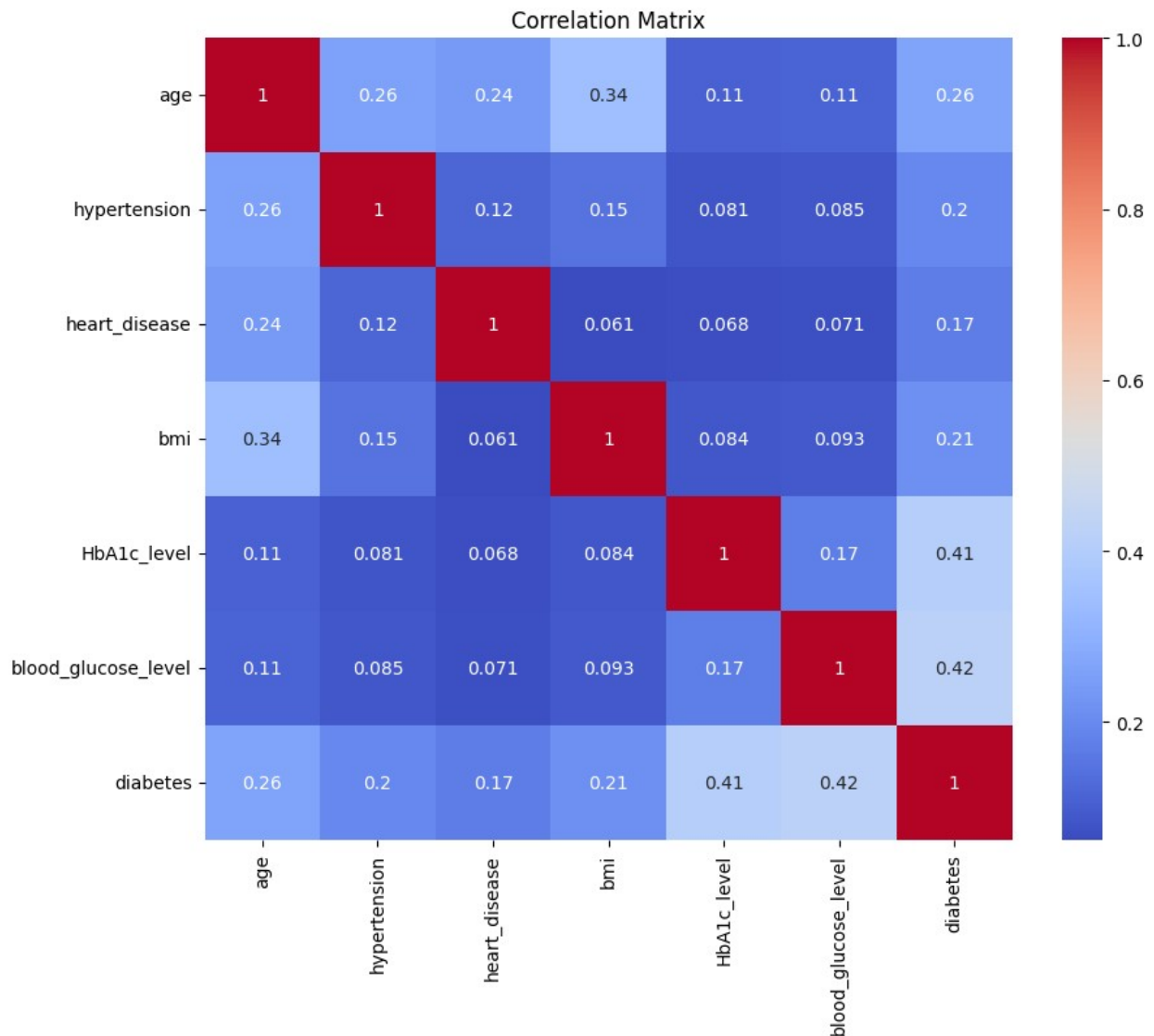
```

# Select only numeric columns
numeric_columns = df.select_dtypes(include='number')

# Calculate the correlation matrix
correlation_matrix = numeric_columns.corr()

# Create a heatmap to visualize the correlations
plt.figure(figsize=(10, 8))
sns.heatmap(correlation_matrix, annot=True, cmap="coolwarm")
plt.title("Correlation Matrix")
plt.show()

```



### Observing Age with BMI, HbA1c\_level, and blood\_glucose\_level

- On the basis of data, by increasing age bmi also increases.
- On the basis of data, Positive low correlation between age and HbA1c\_level.
- On the basis of data, Positive low correlation between age and blood\_glucose\_level.

```
df.drop(df[df['smoking_history'] == 'No Info'].index, inplace=True)

fig = px.sunburst(
    df[df['smoking_history'] != 'No info'],

    path=['smoking_history', 'hypertension_categorical', 'heart_disease_categorical', 'gender', 'age_category'],
    values='diabetes',
    color='gender',
)
```

```
# Show the Sunburst chart
fig.show()
```

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Q6. On the basis of gender who had more diabetes that were smoking, had heart disease and hypertension?

A. Based on Data people who were smoking, had heart disease and hypertension, male were having higher rate of Diabetes.

- Male: 28
- Female: 15

Q7. On the basis of gender who had more diabetes that never smoked?

A. Based on Data people who never smoked, female were having higher rate of Diabetes.

- Female: 1288
  - Male: 868
-

## Summary

### Q1. Comparatively who has more Diabetes?

A. Based on given data females has more diabetes than male.

### Q2. Comparatively who has more heart\_disease?

A. Based on given data male has more heart issues than female.

- Male: 2367
- Female: 1556
- Others: 0

### Q3. Comparatively who has more hypertension?

A. Based on given data female has more heart issues than male.

- Female: 4179
- Male: 3282
- Others: 0

### Q4. People with Hypertension and Heart-disease on the basis of gender, who got more diabetes?

A. Based on Data people with Hypertension and Hear\_disease, Male has more Diabetes.

- Male : 194
- Female : 164

### Q5. People with no Hypertension and no Heart-disease on the basis of gender, who got more diabetes?

A. Based on Data people with Hypertension and Hear\_disease, female has more Diabetes.

- Male 2518
- Female 2969

### Q6. On the basis of gender who had more diabetes that were smoking, had heart disease and hypertension?

A. Based on Data people who were smoking, had heart disease and hypertension, male were having higher rate of Diabetes.

- Male: 28
- Female: 15

### Q7. On the basis of gender who had more diabetes that never smoked?

A. Based on Data people who never smoked, female were having higher rate of Diabetes.



- Female: 1288
- Male: 868

### Conclusion:

Based on the given data, here are the conclusions for each question:

- Comparatively, females have more diabetes than males.
- Comparatively, males have more heart disease than females.
- Comparatively, females have more hypertension than males.
- Among people with both hypertension and heart disease, males have more diabetes than females.
- Among people with neither hypertension nor heart disease, females have more diabetes than males.
- Among individuals who were smoking, had heart disease, and hypertension, males have a higher rate of diabetes compared to females.
- Among individuals who never smoked, females have a higher rate of diabetes compared to males.