

EC9630 Machine Learning - laboratory 01

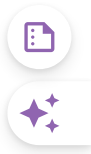
Task: STASTICAL PATTEN CLASSIFICATION

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RegNo: 2020/E/082

Date: 28th April 2024

Time:



## Question 02

(a). The Diabetes Health Indicators Dataset contains healthcare statistics and lifestyle survey information about people in general along with their diagnosis of diabetes. The 35 features consist of some demographics, lab test results, and answers to survey questions for each patient. The target variable for classification is whether a patient has diabetes, is pre-diabetic, or healthy. Dataset Characteristics Tabular, Multivariate

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Subject Area: Health and Medicine

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Associated Tasks: Classification

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Feature Type: Categorical, Integer

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Instances: 253680

---

Features: 21

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(b). Categorical and Integer

(c). Features: The dataset includes various features like demographics, health history, and personal information that are used to predict the target variable, which is the diabetes status of the individual.

Labels: The target variable in the dataset is the diabetes diagnosis, categorized into three classes: Diabetes, Pre-diabetes, and Healthy. This variable is used for classification purposes to predict the health status of individuals based on the provided features.

(d). 21 Features.

```
import pandas as pd
```

```
# Load the dataset
```

```
df = pd.read_csv("/content/drive/MyDrive/CSV FILE/diabetes_012_health_indicators_BRFSS2013.csv")  
df.describe()
```



	Diabetes_012	HighBP	HighChol	CholCheck	BMI	
<b>count</b>	253680.000000	253680.000000	253680.000000	253680.000000	253680.000000	253680.000000
<b>mean</b>	0.296921	0.429001	0.424121	0.962670	28.382364	
<b>std</b>	0.698160	0.494934	0.494210	0.189571	6.608694	
<b>min</b>	0.000000	0.000000	0.000000	0.000000	12.000000	
<b>25%</b>	0.000000	0.000000	0.000000	1.000000	24.000000	
<b>50%</b>	0.000000	0.000000	0.000000	1.000000	27.000000	
<b>75%</b>	0.000000	1.000000	1.000000	1.000000	31.000000	
<b>max</b>	2.000000	1.000000	1.000000	1.000000	98.000000	

8 rows × 22 columns

```
# the Variable type informationS
print(df.info())
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 253680 entries, 0 to 253679
Data columns (total 22 columns):
#   Column                                Non-Null Count  Dtype
---  -
0   Diabetes_012                          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)
memory usage: 42.6 MB
None
```



```
# Identify features (all columns except the label column)
features = df.drop(columns=['Diabetes_012'])
```

```
labels = df['Diabetes_012']
```

```
# Display the list of features
print("Features:")
print(features.columns.tolist())
print("\nLabel:")
print(labels.name)
```

```
Features:
['HighBP', 'HighChol', 'CholCheck', 'BMI', 'Smoker', 'Stroke', 'HeartDiseaseorAttack
```

```
Label:
Diabetes_012
```

```
import matplotlib.pyplot as plt
import seaborn as sns
```

```
# Print the column names
print("Column names:")
print(df.columns)
```

```
Column names:
Index(['Diabetes_012', 'HighBP', 'HighChol', 'CholCheck', 'BMI', 'Smoker',
      'Stroke', 'HeartDiseaseorAttack', 'PhysActivity', 'Fruits', 'Veggies',
      'HvyAlcoholConsump', 'AnyHealthcare', 'NoDocbcCost', 'GenHlth',
      'MentHlth', 'PhysHlth', 'DiffWalk', 'Sex', 'Age', 'Education',
      'Income'],
      dtype='object')
```

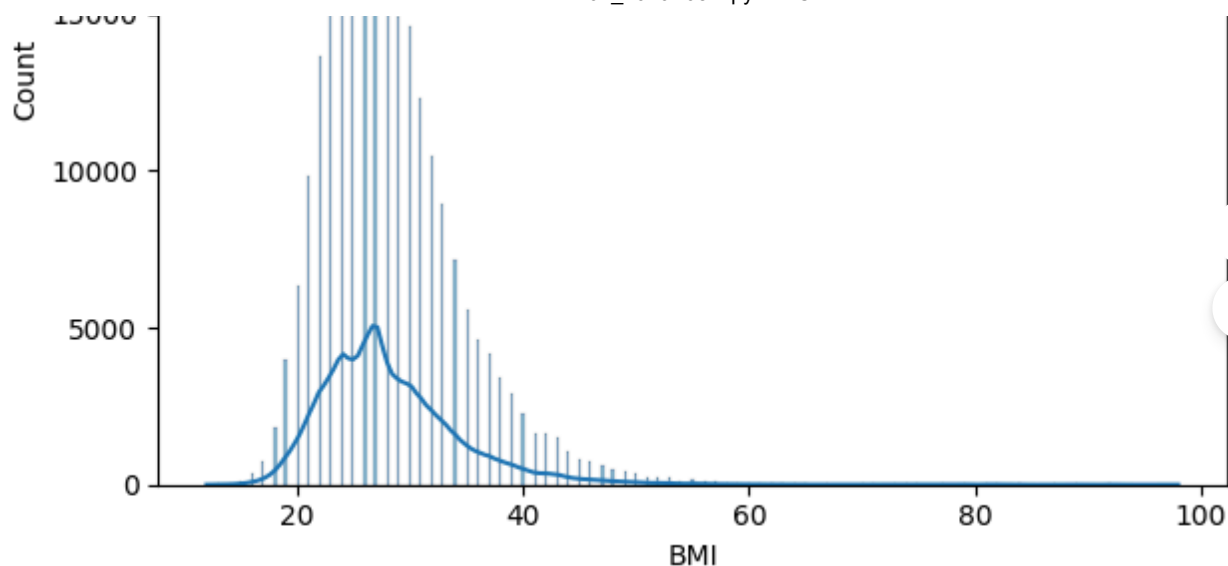


```
# missing values
print("\nMissing values per column:")
print(df.isnull().sum())

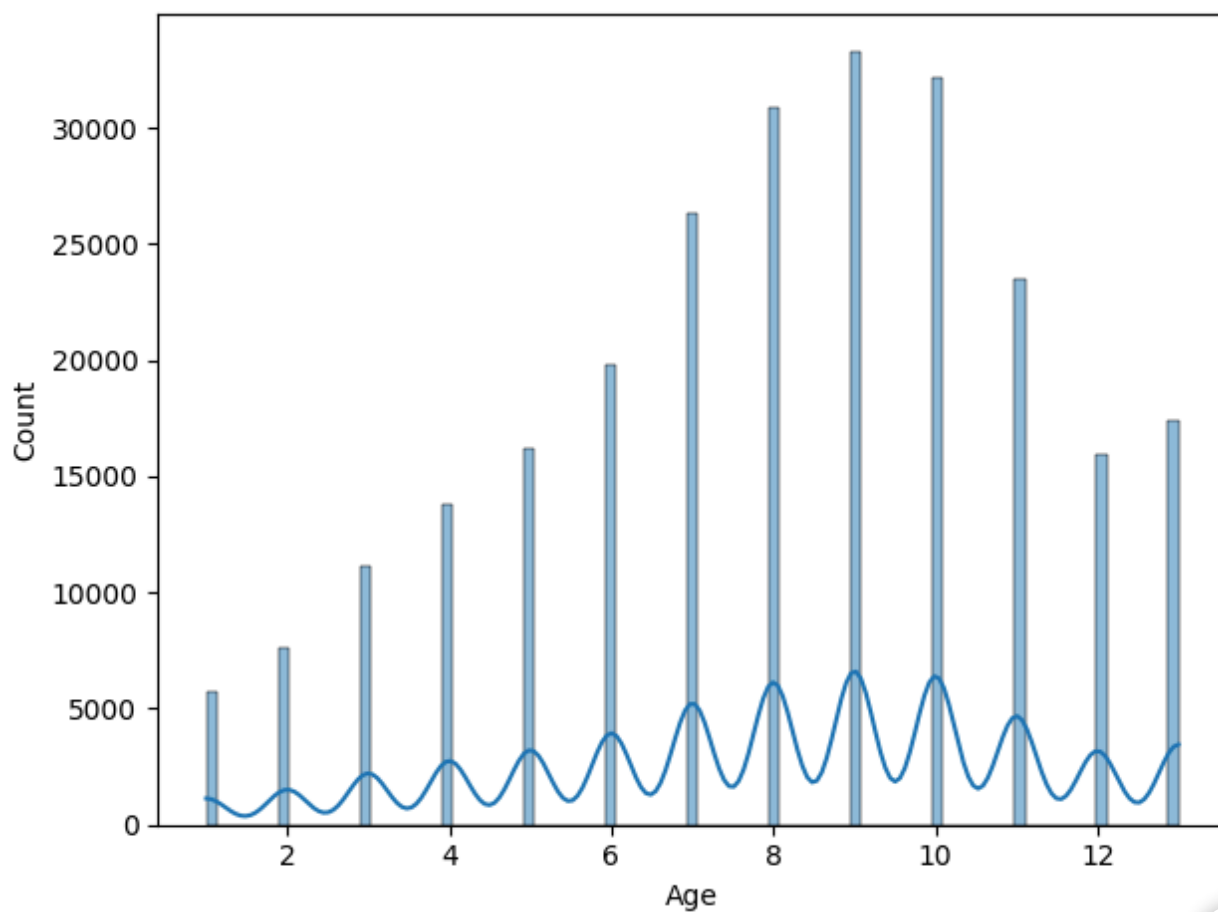
# Distributions-Numerical feature
print("\nNumerical feature distributions:")
for col in ['BMI', 'Age', 'Fruits', 'Veggies', 'PhysActivity']:
    print(f"\n{col}:")
    print(df[col].describe())
    plt.figure()
    sns.histplot(data=df, x=col, kde=True)
    plt.tight_layout()
    plt.show()

# Distributions-Categorical feature
print("\nCategorical feature distributions:")
for col in ['Sex', 'HighBP', 'HighChol', 'Smoker']:
    print(f"\n{col}:")
    print(df[col].value_counts())
    plt.figure()
    sns.countplot(data=df, x=col)
    plt.tight_layout()
    plt.show()
```



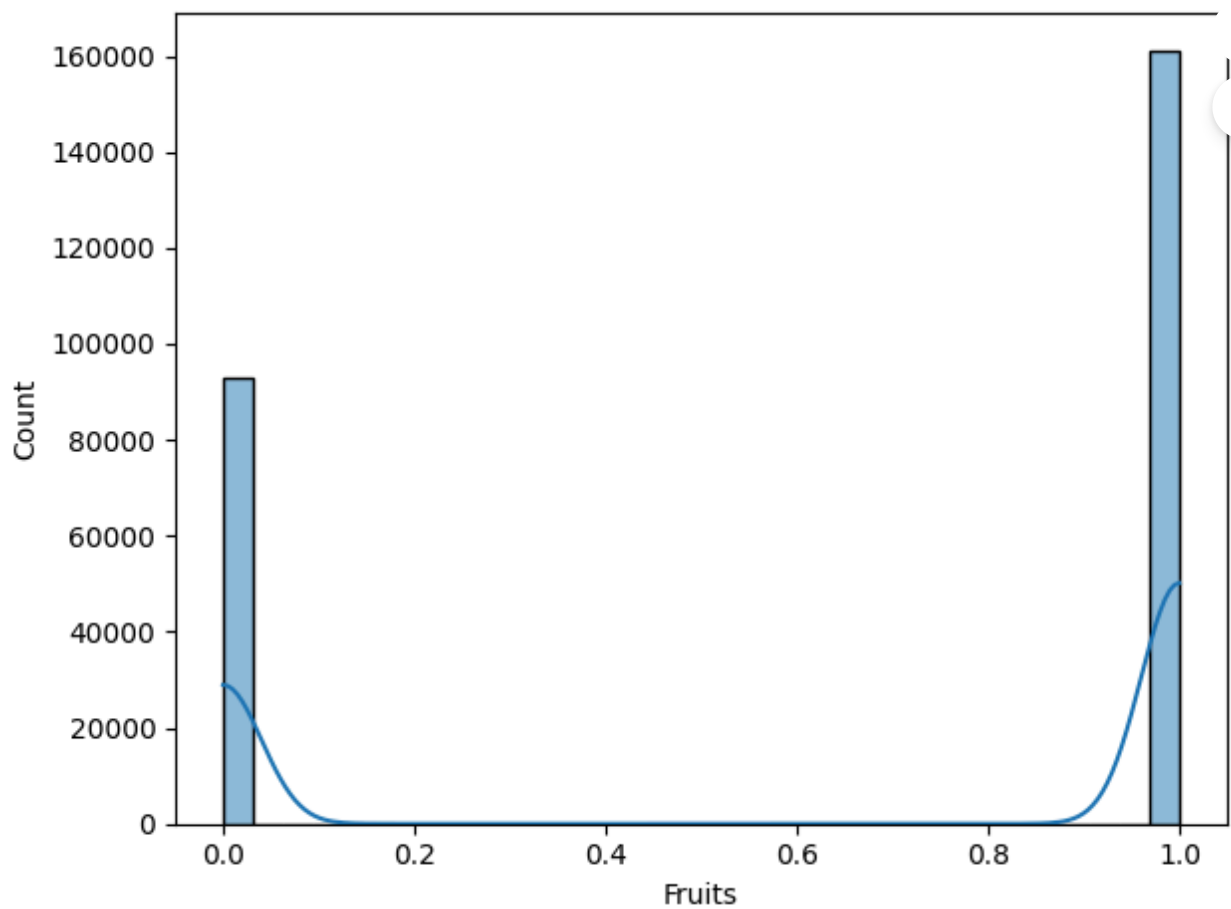


```
Age:
count    253680.000000
mean       8.032119
std        3.054220
min        1.000000
25%        6.000000
50%        8.000000
75%       10.000000
max       13.000000
Name: Age, dtype: float64
```



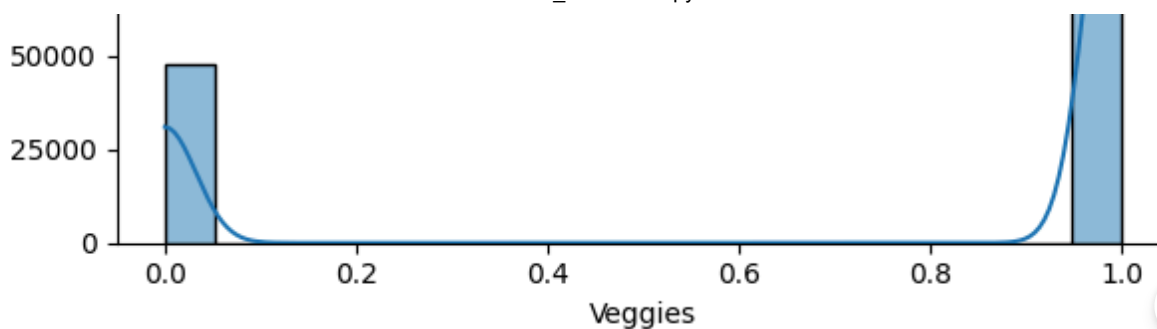
```
Fruits:
count    253680.000000
mean       0.634256
std        0.481639
min        0.000000
```

```
25%      0.000000
50%      1.000000
75%      1.000000
max       1.000000
Name: Fruits, dtype: float64
```



```
Veggies:
count    253680.000000
mean      0.811420
std       0.391175
min       0.000000
25%       1.000000
50%       1.000000
75%       1.000000
max       1.000000
Name: Veggies, dtype: float64
```

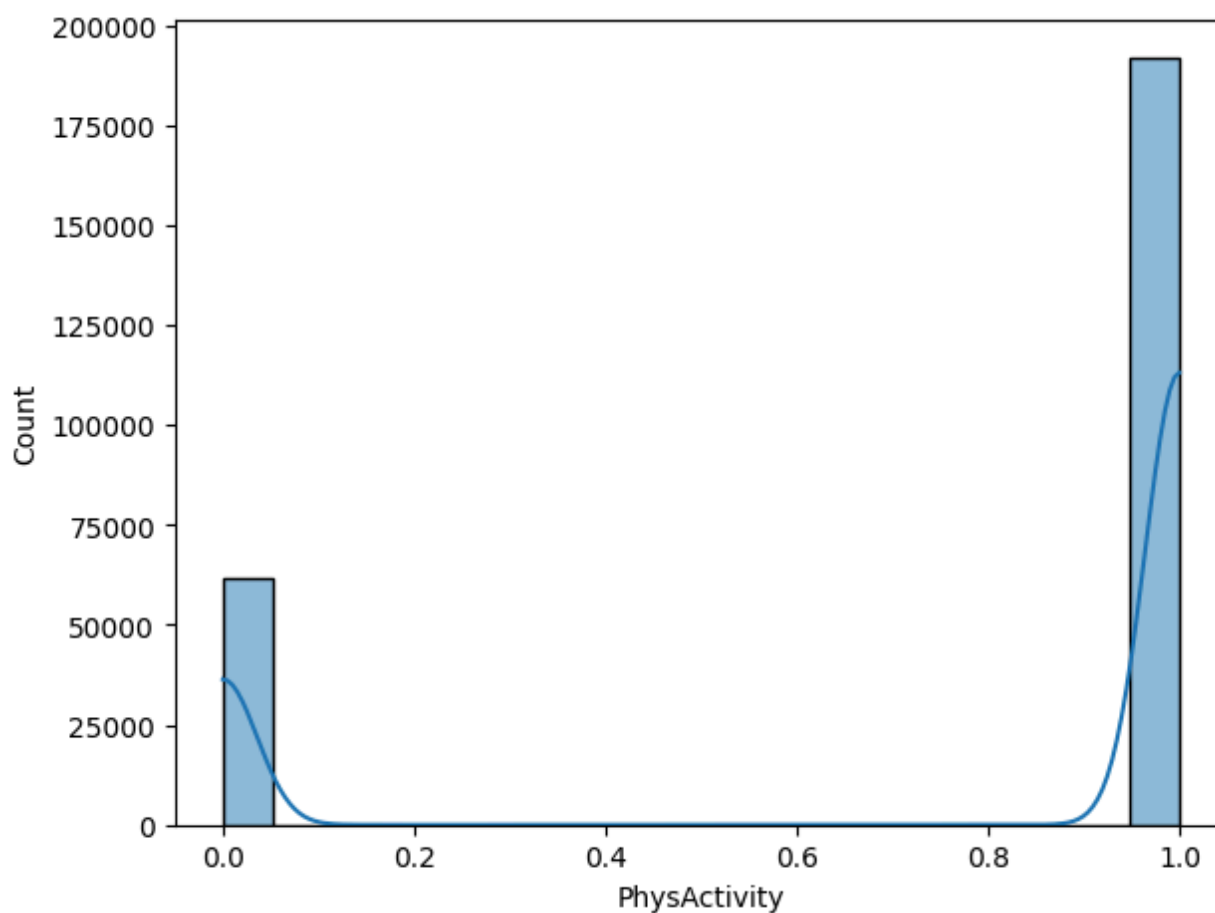




PhysActivity:

```
count    253680.000000
mean      0.756544
std       0.429169
min       0.000000
25%       1.000000
50%       1.000000
75%       1.000000
max       1.000000
```

Name: PhysActivity, dtype: float64



Categorical feature distributions:

Sex:

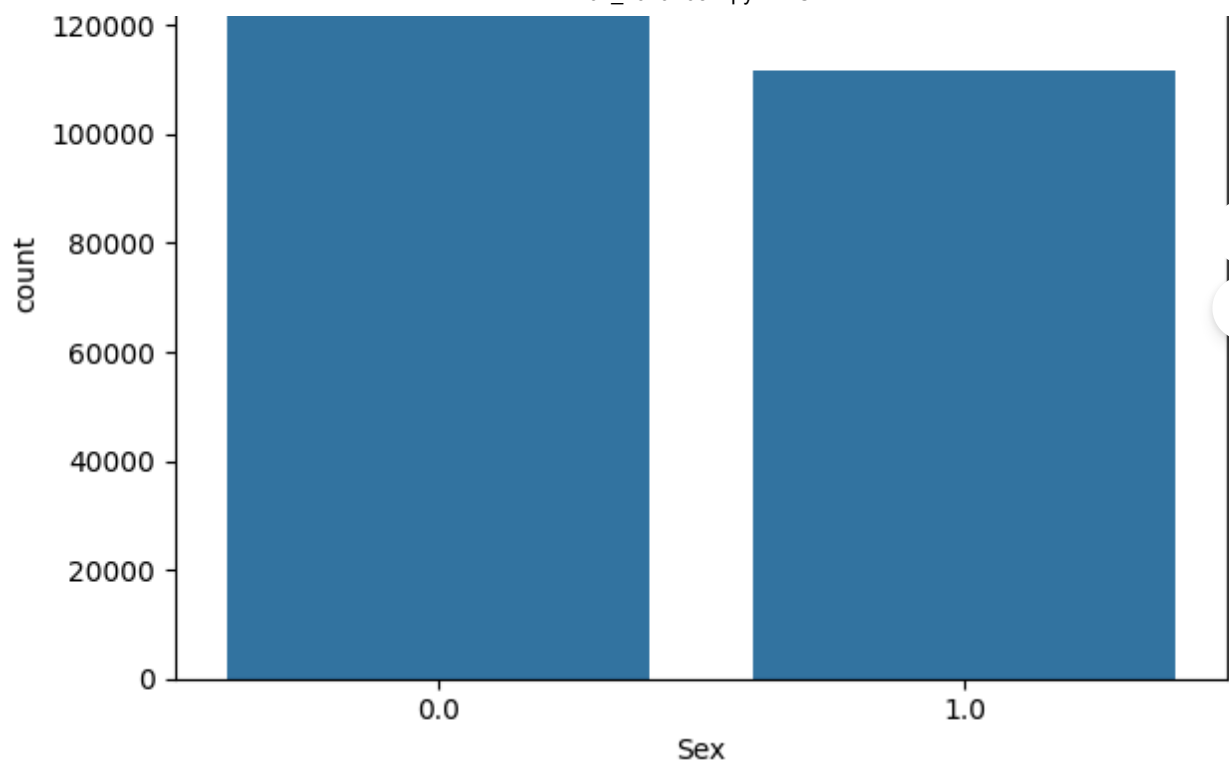
Sex

0.0 141974

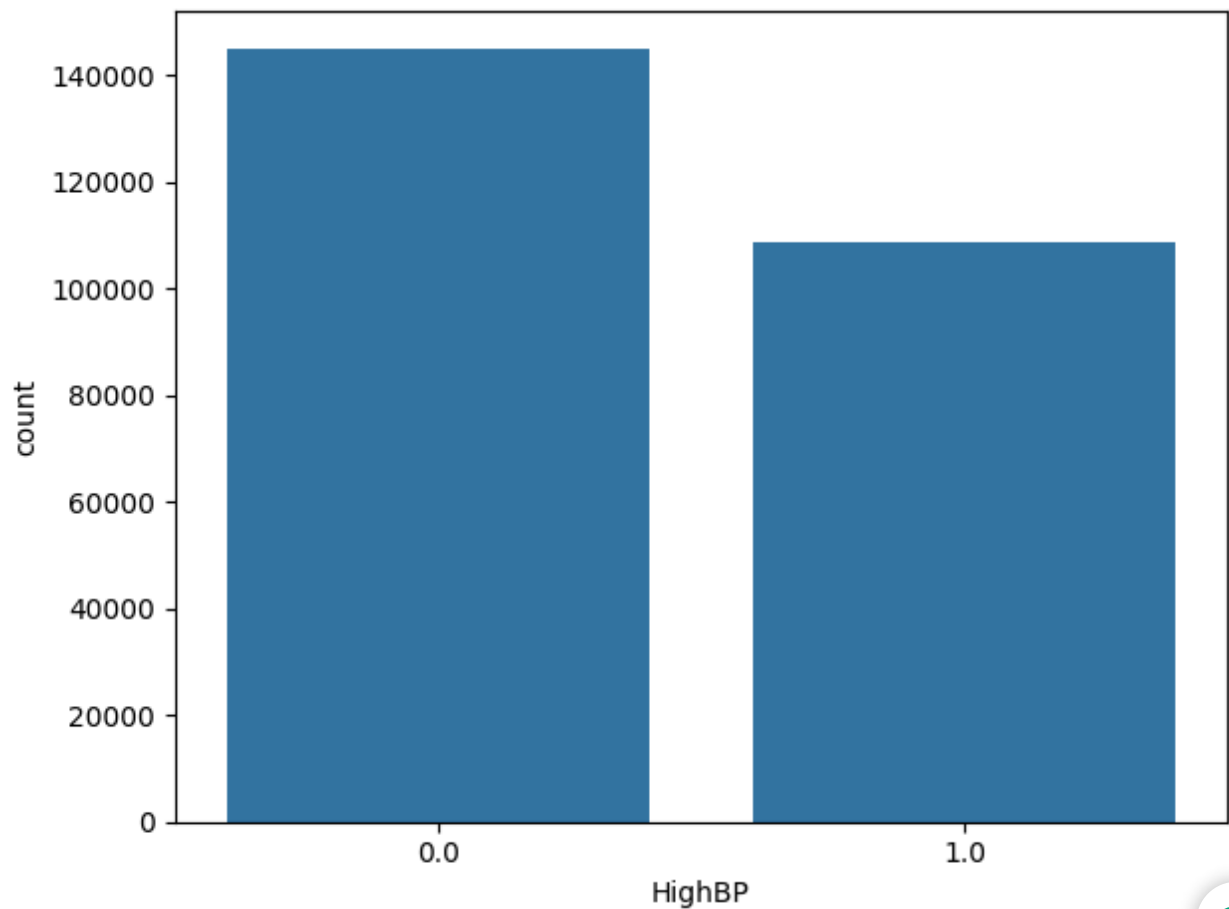
1.0 111706

Name: count, dtype: int64





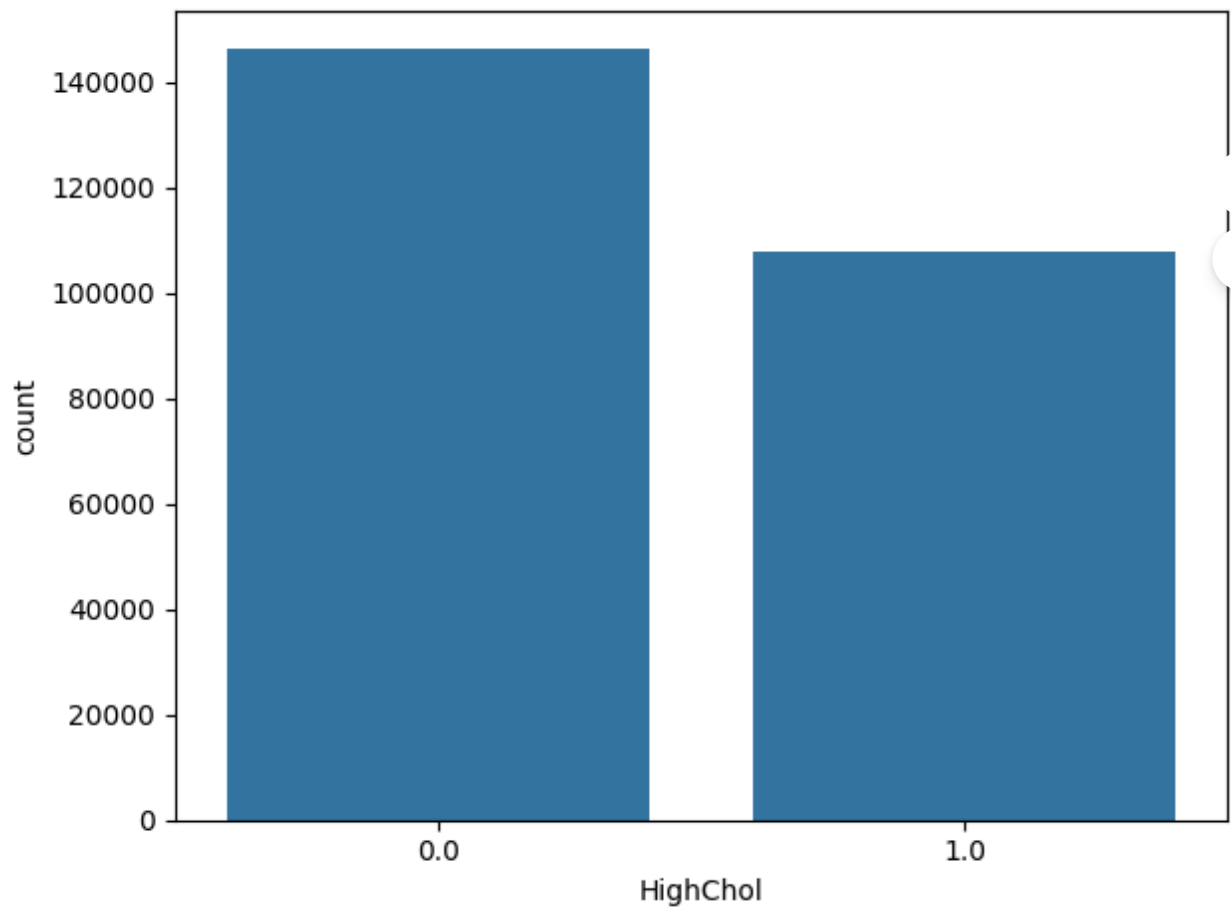
```
HighBP:
HighBP
0.0    144851
1.0    108829
Name: count, dtype: int64
```



```
HighChol:
HighChol
0.0    146089
1.0    107591
Name: count, dtype: int64
```



```
name: count, dtype: int64
```



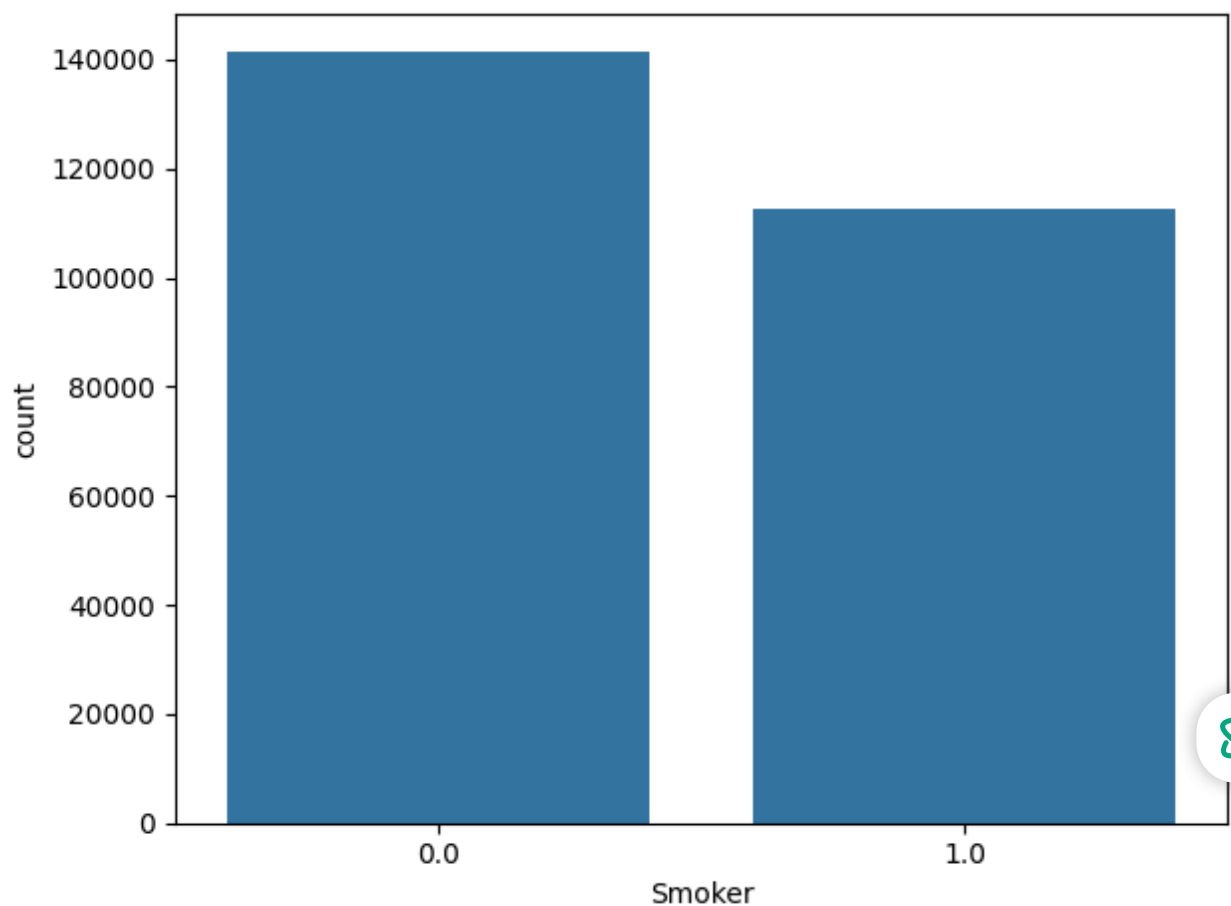
```
Smoker:
```

```
Smoker
```

```
0.0    141257
```

```
1.0    112423
```

```
Name: count, dtype: int64
```





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

# Display the first few rows of the dataset
print("First few rows of the dataset:")
print(df.head())

# Check for missing values in each column
print("\nMissing values per column:")
print(df.isnull().sum())

# Generate descriptive statistics for numerical columns
print("\nDescriptive statistics:")
print(df.describe())

# Create a correlation matrix for numerical variables
print("\nCorrelation matrix:")
print(df.corr())

# Create a scatter plot matrix for numerical variables
plt.figure(figsize=(12, 10))
sns.pairplot(df, diag_kind="kde")
plt.tight_layout()
plt.show()

# Handling missing values
df = df.dropna() # Dropping rows with missing values
# Alternatively, fill missing values with the mean: df = df.fillna(df.mean())

# Encoding categorical variables using one-hot encoding
categorical_cols = ['Sex', 'HighBP', 'HighChol', 'Smoker']
df = pd.get_dummies(df, columns=categorical_cols, drop_first=True)

# Splitting features and labels
X = df.drop('Diabetes_012', axis=1)
y = df['Diabetes_012']

# Display the shapes of features and labels
print("\nShapes of features and labels:")
print("Features shape:", X.shape)
print("Labels shape:", y.shape)
```



First few rows of the dataset:

	Diabetes_012	HighBP	HighChol	CholCheck	BMI	Smoker	Stroke	\
0	0.0	1.0	1.0	1.0	40.0	1.0	0.0	
1	0.0	0.0	0.0	0.0	25.0	1.0	0.0	
2	0.0	1.0	1.0	1.0	28.0	0.0	0.0	
3	0.0	1.0	0.0	1.0	27.0	0.0	0.0	
4	0.0	1.0	1.0	1.0	24.0	0.0	0.0	

	HeartDiseaseorAttack	PhysActivity	Fruits	...	AnyHealthcare	\
0	0.0	0.0	0.0	...	1.0	
1	0.0	1.0	0.0	...	0.0	
2	0.0	0.0	1.0	...	1.0	
3	0.0	1.0	1.0	...	1.0	
4	0.0	1.0	1.0	...	1.0	

	NoDocbcCost	GenHlth	MentHlth	PhysHlth	DiffWalk	Sex	Age	Education	\
0	0.0	5.0	18.0	15.0	1.0	0.0	9.0	4.0	
1	1.0	3.0	0.0	0.0	0.0	0.0	7.0	6.0	
2	1.0	5.0	30.0	30.0	1.0	0.0	9.0	4.0	
3	0.0	2.0	0.0	0.0	0.0	0.0	11.0	3.0	
4	0.0	2.0	3.0	0.0	0.0	0.0	11.0	5.0	

	Income
0	3.0
1	1.0
2	8.0
3	6.0
4	4.0

[5 rows x 22 columns]

Missing values per column:

Diabetes_012	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

dtype: int64

Descriptive statistics:

	Diabetes_012	HighBP	HighChol	CholCheck	\
count	253680.000000	253680.000000	253680.000000	253680.000000	
mean	0.296921	0.429001	0.424121	0.962670	

std	0.698160	0.494934	0.494210	0.189571
min	0.000000	0.000000	0.000000	0.000000
25%	0.000000	0.000000	0.000000	1.000000
50%	0.000000	0.000000	0.000000	1.000000
75%	0.000000	1.000000	1.000000	1.000000
max	2.000000	1.000000	1.000000	1.000000

	BMI	Smoker	Stroke	HeartDiseaseorAttack
count	253680.000000	253680.000000	253680.000000	253680.000000
mean	28.382364	0.443169	0.040571	0.094186
std	6.608694	0.496761	0.197294	0.292087
min	12.000000	0.000000	0.000000	0.000000
25%	24.000000	0.000000	0.000000	0.000000
50%	27.000000	0.000000	0.000000	0.000000
75%	31.000000	1.000000	0.000000	0.000000
max	98.000000	1.000000	1.000000	1.000000

	PhysActivity	Fruits	...	AnyHealthcare	NoDocbcCost
count	253680.000000	253680.000000	...	253680.000000	253680.000000
mean	0.756544	0.634256	...	0.951053	0.084177
std	0.429169	0.481639	...	0.215759	0.277654
min	0.000000	0.000000	...	0.000000	0.000000
25%	1.000000	0.000000	...	1.000000	0.000000
50%	1.000000	1.000000	...	1.000000	0.000000
75%	1.000000	1.000000	...	1.000000	0.000000
max	1.000000	1.000000	...	1.000000	1.000000

	GenHlth	MentHlth	PhysHlth	DiffWalk
count	253680.000000	253680.000000	253680.000000	253680.000000
mean	2.511392	3.184772	4.242081	0.168224
std	1.068477	7.412847	8.717951	0.374066
min	1.000000	0.000000	0.000000	0.000000
25%	2.000000	0.000000	0.000000	0.000000
50%	2.000000	0.000000	0.000000	0.000000
75%	3.000000	2.000000	3.000000	0.000000
max	5.000000	30.000000	30.000000	1.000000

	Sex	Age	Education	Income
count	253680.000000	253680.000000	253680.000000	253680.000000
mean	0.440342	8.032119	5.050434	6.053875
std	0.496429	3.054220	0.985774	2.071148
min	0.000000	1.000000	1.000000	1.000000
25%	0.000000	6.000000	4.000000	5.000000
50%	0.000000	8.000000	5.000000	7.000000
75%	1.000000	10.000000	6.000000	8.000000
max	1.000000	13.000000	6.000000	8.000000

[8 rows x 22 columns]

Correlation matrix:

	Diabetes_012	HighBP	HighChol	CholCheck	BMI
Diabetes_012	1.000000	0.271596	0.209085	0.067546	0.224379
HighBP	0.271596	1.000000	0.298199	0.098508	0.213748
HighChol	0.209085	0.298199	1.000000	0.085642	0.106722
CholCheck	0.067546	0.098508	0.085642	1.000000	0.034495
BMI	0.224379	0.213748	0.106722	0.034495	1.000000
Smoker	0.062914	0.096991	0.091299	-0.009929	0.013804
Stroke	0.107179	0.129575	0.092620	0.024158	0.020153
HeartDiseaseorAttack	0.180272	0.209361	0.180765	0.044206	0.052904
PhysActivity	-0.121947	-0.125267	-0.078046	0.004190	-0.147294

Fruits	-0.042192	-0.040555	-0.040859	0.023849	-0.087518
Veggies	-0.058972	-0.061266	-0.039874	0.006121	-0.062275
HvyAlcoholConsump	-0.057882	-0.003972	-0.011543	-0.023730	-0.048736
AnyHealthcare	0.015410	0.038425	0.042230	0.117626	-0.018471
NoDocbcCost	0.035436	0.017358	0.013310	-0.058255	0.058206
GenHlth	0.302587	0.300530	0.208426	0.046589	0.239185
MentHlth	0.073507	0.056456	0.062069	-0.008366	0.085310
PhysHlth	0.176287	0.161212	0.121751	0.031775	0.121141
DiffWalk	0.224239	0.223618	0.144672	0.040585	0.197078
Sex	0.031040	0.052207	0.031205	-0.022115	0.042950
Age	0.185026	0.344452	0.272318	0.090321	-0.036618
Education	-0.130517	-0.141358	-0.070802	0.001510	-0.103932
Income	-0.171483	-0.171235	-0.085459	0.014259	-0.100069

	Smoker	Stroke	HeartDiseaseorAttack	PhysActivity	\
Diabetes_012	0.062914	0.107179	0.180272	-0.121947	
HighBP	0.096991	0.129575	0.209361	-0.125267	
HighChol	0.091299	0.092620	0.180765	-0.078046	
CholCheck	-0.009929	0.024158	0.044206	0.004190	
BMI	0.013804	0.020153	0.052904	-0.147294	
Smoker	1.000000	0.061173	0.114441	-0.087401	
Stroke	0.061173	1.000000	0.203002	-0.069151	
HeartDiseaseorAttack	0.114441	0.203002	1.000000	-0.087299	
PhysActivity	-0.087401	-0.069151	-0.087299	1.000000	
Fruits	-0.077666	-0.013389	-0.019790	0.142756	
Veggies	-0.030678	-0.041124	-0.039167	0.153150	
HvyAlcoholConsump	0.101619	-0.016950	-0.028991	0.012392	
AnyHealthcare	-0.023251	0.008776	0.018734	0.035505	
NoDocbcCost	0.048946	0.034804	0.031000	-0.061638	
GenHlth	0.163143	0.177942	0.258383	-0.266186	
MentHlth	0.092196	0.070172	0.064621	-0.125587	
PhysHlth	0.116460	0.148944	0.181698	-0.219230	
DiffWalk	0.122463	0.176567	0.212709	-0.253174	
Sex	0.093662	0.002978	0.086096	0.032482	
Age	0.120641	0.126974	0.221618	-0.092511	
Education	-0.161955	-0.076009	-0.099600	0.199658	
Income	-0.123937	-0.128599	-0.141011	0.198539	

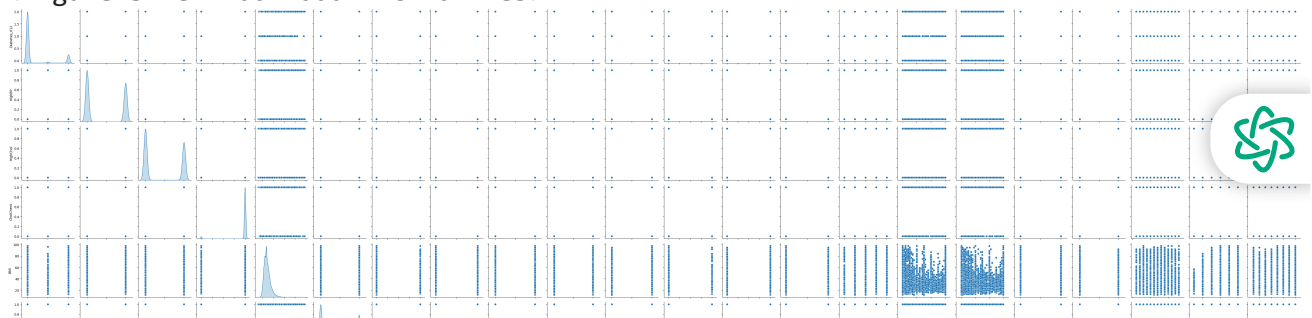
	Fruits	...	AnyHealthcare	NoDocbcCost	GenHlth	\
Diabetes_012	-0.042192	...	0.015410	0.035436	0.302587	
HighBP	-0.040555	...	0.038425	0.017358	0.300530	
HighChol	-0.040859	...	0.042230	0.013310	0.208426	
CholCheck	0.023849	...	0.117626	-0.058255	0.046589	
BMI	-0.087518	...	-0.018471	0.058206	0.239185	
Smoker	-0.077666	...	-0.023251	0.048946	0.163143	
Stroke	-0.013389	...	0.008776	0.034804	0.177942	
HeartDiseaseorAttack	-0.019790	...	0.018734	0.031000	0.258383	
PhysActivity	0.142756	...	0.035505	-0.061638	-0.266186	
Fruits	1.000000	...	0.031544	-0.044243	-0.103854	
Veggies	0.254342	...	0.029584	-0.032232	-0.123066	
HvyAlcoholConsump	-0.035288	...	-0.010488	0.004684	-0.036724	
AnyHealthcare	0.031544	...	1.000000	-0.232532	-0.040817	
NoDocbcCost	-0.044243	...	-0.232532	1.000000	0.166397	
GenHlth	-0.103854	...	-0.040817	0.166397	1.000000	
MentHlth	-0.068217	...	-0.052707	0.192107	0.301674	
PhysHlth	-0.044633	...	-0.008276	0.148998	0.524364	
DiffWalk	-0.048352	...	0.007074	0.118447	0.456920	
Sex	-0.091175	...	-0.019405	-0.044931	-0.006091	
Age	0.064547	...	0.138046	-0.119777	0.152450	
Education	0.110187	...	0.122514	-0.100701	-0.284912	
Income	0.079929	...	0.157999	-0.203182	-0.370014	

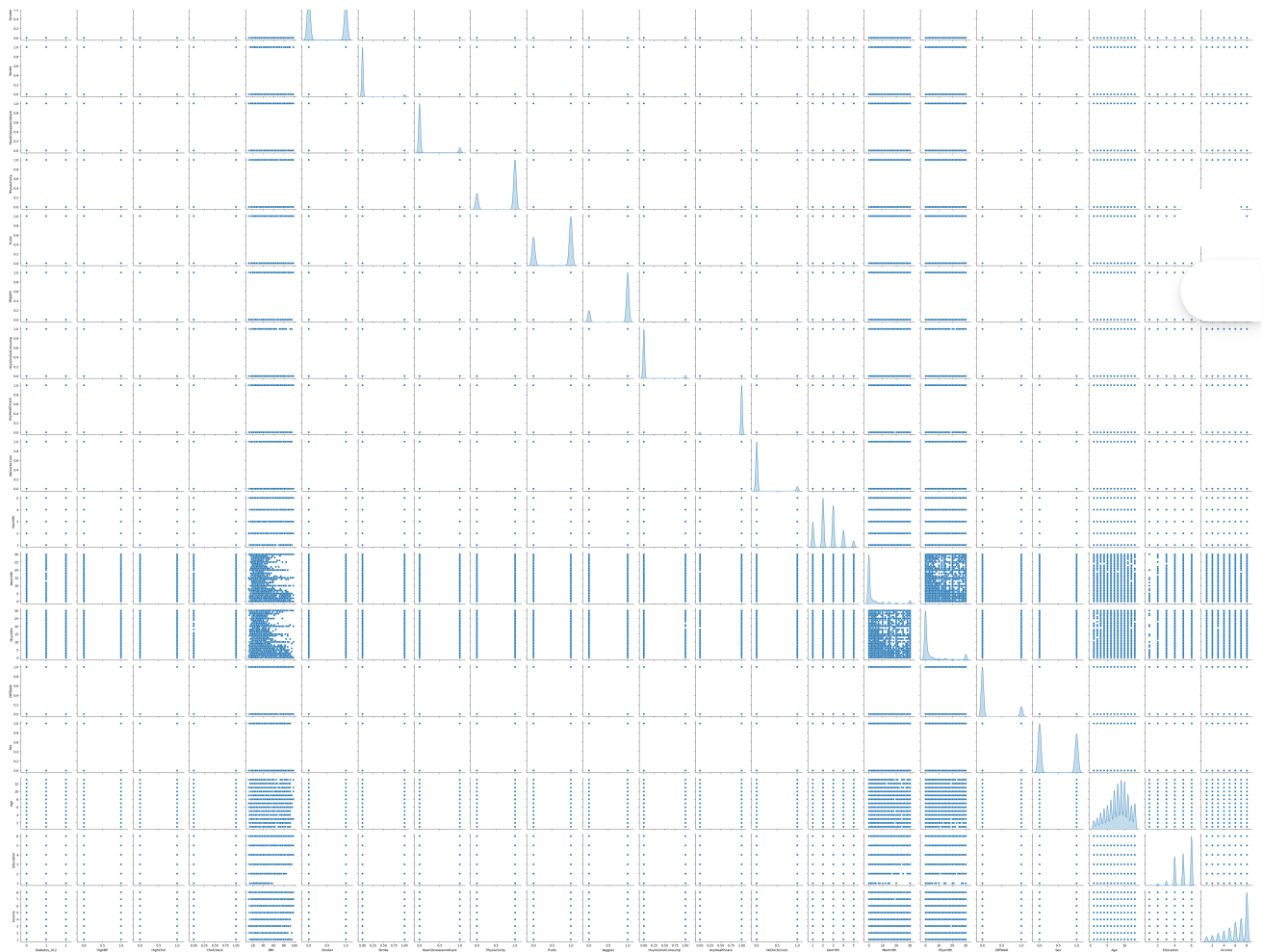
	MentHlth	PhysHlth	DiffWalk	Sex	Age
Diabetes_012	0.073507	0.176287	0.224239	0.031040	0.185026
HighBP	0.056456	0.161212	0.223618	0.052207	0.344452
HighChol	0.062069	0.121751	0.144672	0.031205	0.272318
CholCheck	-0.008366	0.031775	0.040585	-0.022115	0.090321
BMI	0.085310	0.121141	0.197078	0.042950	-0.036618
Smoker	0.092196	0.116460	0.122463	0.093662	0.120641
Stroke	0.070172	0.148944	0.176567	0.002978	0.126974
HeartDiseaseorAttack	0.064621	0.181698	0.212709	0.086096	0.221618
PhysActivity	-0.125587	-0.219230	-0.253174	0.032482	-0.092511
Fruits	-0.068217	-0.044633	-0.048352	-0.091175	0.064547
Veggies	-0.058884	-0.064290	-0.080506	-0.064765	-0.009771
HvyAlcoholConsump	0.024716	-0.026415	-0.037668	0.005740	-0.034578
AnyHealthcare	-0.052707	-0.008276	0.007074	-0.019405	0.138046
NoDocbcCost	0.192107	0.148998	0.118447	-0.044931	-0.119777
GenHlth	0.301674	0.524364	0.456920	-0.006091	0.152450
MentHlth	1.000000	0.353619	0.233688	-0.080705	-0.092068
PhysHlth	0.353619	1.000000	0.478417	-0.043137	0.099130
DiffWalk	0.233688	0.478417	1.000000	-0.070299	0.204450
Sex	-0.080705	-0.043137	-0.070299	1.000000	-0.027340
Age	-0.092068	0.099130	0.204450	-0.027340	1.000000
Education	-0.101830	-0.155093	-0.192642	0.019480	-0.101901
Income	-0.209806	-0.266799	-0.320124	0.127141	-0.127775

	Education	Income
Diabetes_012	-0.130517	-0.171483
HighBP	-0.141358	-0.171235
HighChol	-0.070802	-0.085459
CholCheck	0.001510	0.014259
BMI	-0.103932	-0.100069
Smoker	-0.161955	-0.123937
Stroke	-0.076009	-0.128599
HeartDiseaseorAttack	-0.099600	-0.141011
PhysActivity	0.199658	0.198539
Fruits	0.110187	0.079929
Veggies	0.154329	0.151087
HvyAlcoholConsump	0.023997	0.053619
AnyHealthcare	0.122514	0.157999
NoDocbcCost	-0.100701	-0.203182
GenHlth	-0.284912	-0.370014
MentHlth	-0.101830	-0.209806
PhysHlth	-0.155093	-0.266799
DiffWalk	-0.192642	-0.320124
Sex	0.019480	0.127141
Age	-0.101901	-0.127775
Education	1.000000	0.449106
Income	0.449106	1.000000

[22 rows x 22 columns]

<Figure size 1200x1000 with 0 Axes>





Shapes of features and labels:

Features shape: (253680, 21)

Labels shape: (253680,)





```
# Task 4: Separating features and labels
X = df.drop('Diabetes_012', axis=1) # Features
y = df['Diabetes_012'] # Labels
```

```
# Print the shapes of features and labels
print("Shape of features (X):", X.shape)
print("Shape of labels (y):", y.shape)
```

```
Shape of features (X): (253680, 21)
Shape of labels (y): (253680,)
```

```
# Task 5: Calculating Information Gain
from sklearn.metrics import mutual_info_score
```

```
# Separate features and labels
X = df.drop('Diabetes_012', axis=1)
y = df['Diabetes_012']
```

```
# Calculate information gain for each feature
information_gain = []
for feature in X.columns:
    info_gain = mutual_info_score(X[feature], y)
    information_gain.append(info_gain)
```

```
# Create a DataFrame to display feature information gain
feature_info_gain = pd.DataFrame({'Feature': X.columns, 'Information Gain': information_
feature_info_gain = feature_info_gain.sort_values(by='Information Gain', ascending=False)
print(feature_info_gain)
```

	Feature	Information Gain
10	GenHlth	0.047067
18	HighBP_1.0	0.037435
1	BMI	0.030067
14	Age	0.022087
19	HighChol_1.0	0.022049
13	DiffWalk	0.021242
16	Income	0.015080
12	PhysHlth	0.014923
3	HeartDiseaseorAttack	0.013050
15	Education	0.008582
4	PhysActivity	0.006936
2	Stroke	0.004573
0	CholCheck	0.003242
11	MentHlth	0.003085
7	HvyAlcoholConsump	0.002043
20	Smoker_1.0	0.001980
6	Veggies	0.001671
5	Fruits	0.000884
9	NoDocbcCost	0.000722
17	Sex_1.0	0.000492
8	AnyHealthcare	0.000143

## # Task 6: Splitting Data into Train, Validation, and Test Sets

```

from sklearn.model_selection import train_test_split

# Separate features and labels
X = df.drop('Diabetes_012', axis=1)
y = df['Diabetes_012']

# Split the data into train, validation, and test sets
X_train, X_temp, y_train, y_temp = train_test_split(X, y, test_size=0.3, random_state=42)
X_val, X_test, y_val, y_test = train_test_split(X_temp, y_temp, test_size=0.25, random_state=42)

# Print the shapes of the data splits
print("Shape of X_train:", X_train.shape)
print("Shape of y_train:", y_train.shape)
print("Shape of X_val:", X_val.shape)
print("Shape of y_val:", y_val.shape)
print("Shape of X_test:", X_test.shape)
print("Shape of y_test:", y_test.shape)

```

```

Shape of X_train: (177576, 21)
Shape of y_train: (177576,)
Shape of X_val: (57078, 21)
Shape of y_val: (57078,)
Shape of X_test: (19026, 21)
Shape of y_test: (19026,)

```

## # Task 7: Training a Decision Tree Model

```

from sklearn.tree import DecisionTreeClassifier
from sklearn.metrics import accuracy_score

```

## # Train the decision tree model

```

dtc = DecisionTreeClassifier(random_state=42)
dtc.fit(X_train, y_train)

```

## # Make predictions on the test set

```

y_pred = dtc.predict(X_test)

```

## # Calculate the accuracy of the model

```

accuracy = accuracy_score(y_test, y_pred)
print(f"Accuracy: {accuracy:.2f}")

```

```

Accuracy: 0.77

```

## In Task 7,

1. Imports the DecisionTreeClassifier from scikit-learn's tree module and the accuracy\_score function from the metrics module.
2. Creates an instance of the DecisionTreeClassifier with random\_state=42 for reproducibility.
3. Trains the decision tree model on the training data X\_train and y\_train using the fit method.



4. Makes predictions on the test data `X_test` using the `predict` method and stores the predictions in `y_pred`.
5. Calculates the accuracy of the model by comparing the predicted labels `y_pred` with the true labels `y_test` using the `accuracy_score` function.
6. Prints the accuracy score.

Hyperparameters of the decision tree model and their significance:

1. `criterion` (default='gini'): This parameter specifies the function to measure the quality of a split. The two options are 'gini' for the Gini impurity and 'entropy' for the information gain (entropy). According to the scikit-learn documentation, the Gini impurity is a measure of node impurity, and it is the default criterion for classification tasks.
2. `max_depth` (default=None): This parameter sets the maximum depth of the tree. A higher value allows the tree to grow deeper, which can lead to overfitting on the training data. Setting it to None (default) means that the tree can grow without any depth restriction.
3. `min_samples_split` (default=2): This parameter specifies the minimum number of samples required to split an internal node. A higher value can prevent overfitting by not allowing splits in nodes with too few samples.
4. `min_samples_leaf` (default=1): This parameter sets the minimum number of samples required to be at a leaf node. Increasing this value can prevent overfitting by pruning the tree and reducing the number of leaf nodes.
5. `max_features` (default=None): This parameter limits the number of features to consider when looking for the best split. Setting it to None (default) means that all features are considered for splitting.
6. `random_state` (default=None): This parameter controls the randomness of the decision tree's behavior when using random subsets of features for splitting. Setting a fixed value ensures reproducibility of results.

```
# Task 8: Evaluating Training Accuracy
from sklearn.tree import DecisionTreeClassifier
from sklearn.metrics import accuracy_score

# Train the decision tree model
dtc = DecisionTreeClassifier(random_state=42)
dtc.fit(X_train, y_train)

# Make predictions on the train set
y_train_pred = dtc.predict(X_train)

# Calculate the training accuracy
train_accuracy = accuracy_score(y_train, y_train_pred)
print(f"Training Accuracy: {train_accuracy:.2f}")
```

Training Accuracy: 0.99