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# **Diabetes Prediction Using Machine Learning**

Diabetes is a medical disorder that impacts how well our body uses food as fuel. Most food we eat daily is converted to sugar, commonly known as glucose, and then discharged into the bloodstream. Our pancreas releases insulin when the blood sugar levels rise.

Diabetes can cause blood sugar levels to rise if it is not continuously and carefully managed, which raises the chance of severe side effects like heart attack and stroke. We, therefore, choose to forecast using Python machine learning.

#### Steps

- 1. Installing the Libraries
- 2. Importing the Dataset
- 3. Filling the Missing Values
- 4. Exploratory Data Analysis
- 5. Feature Engineering
- 6. Implementing Machine Learning Models
- 7. Predicting Unseen Data
- 8. Concluding the Report

## Installing the Libraries

We first have to import the most popular Python libraries, which we will use for implementing machine learning algorithms in the first step of building the project, including Pandas, Seaborn, Matplotlib, and others.





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We will use Python because it is the most adaptable and powerful programming language for data analysis purposes. In the world of software development, we also use Python.

#### Code

# Import libraries

import numpy as np # for linear algebra

import pandas as pd # for data processing, CSV file I/O (e.g. pd.read\_csv)

import seaborn as sns # for data visualization

import matplotlib.pyplot as plt # to plot data visualization charts

from collections import Counter

import os

# Modeling Libraries

from sklearn.metrics import confusion\_matrix, accuracy\_score, precision\_score

from sklearn.preprocessing import QuantileTransformer

from sklearn.linear\_model import LogisticRegression

from sklearn.neighbors import KNeighborsClassifier

from sklearn.tree import DecisionTreeClassifier

from sklearn.ensemble import RandomForestClassifier, AdaBoostClassifier, GradientBoostingClassifier

from sklearn.model\_selection import GridSearchCV, cross\_val\_score, StratifiedKFold, learning\_curve, train\_test\_split

from sklearn.svm import SVC

The Sklearn toolkit is incredibly practical and helpful and has practical applications. It offers a vast selection of ML models and algorithms.

## Importing the Dataset

We are using the Diabetes Dataset from Kaggle for this study. The National Institute of Diabetes and Digestive and Kidney Diseases is the original source of this database.

#### Code

```
# Importing the dataset from Kaggle

data = pd.read_csv("../input/pima-indians-diabetes-database/diabetes.csv")

# First step is getting familiar with the structure of the dataset

data.info()
```

### Output

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 768 entries, 0 to 767
Data columns (total 9 columns):
    Column
                              Non-Null Count Dtype
    -----
                              _____
---
0
    Pregnancies
                              768 non-null
                                              int64
1
    Glucose
                              768 non-null
                                              int64
    BloodPressure
                              768 non-null
                                              int64
    SkinThickness
 3
                              768 non-null
                                              int64
    Insulin
4
                              768 non-null
                                              int64
 5
    BMI
                              768 non-null
                                              float64
                                              float64
    DiabetesPedigreeFunction 768 non-null
 6
7
                              768 non-null
                                              int64
    Age
                              768 non-null
                                              int64
    Outcome
dtypes: float64(2), int64(7)
memory usage: 54.1 KB
```

As we can see, all of the columns are integers, except for the BMI and DiabetesPedigreeFunction. The target variable is the labels with values of 1 and 0. A person's diabetes status is indicated by a one or a zero.

## Code

# Showing the top 5 rows of the dataset data.head()

## Output

Pregnancies   Glucose   BloodPressure   SkinThickness   Insulin   BMI   DiabetesPedigreeFunction   Age   Outcome
--

0	6	148	72	35	0	33.6	0.627	50	1
1	1	85	66	29	0	26.6	0.351	31	0
2	8	183	64	0	0	23.3	0.672	32	1
3	1	89	66	23	94	28.1	0.167	21	0
4	0	137	40	35	168	43.1	2.288	33	1

# Filling the Missing Values

The next step is cleaning the dataset, which is a crucial step in data analysis. When modelling and making predictions, missing data can result in incorrect results.

#### Code

# Exploring the missing values in the diabetes dataset data.isnull().sum()

### Output

Pregnancies	0
Glucose	0
BloodPressure	0
SkinThickness	0
Insulin	0
BMI	0
DiabetesPedigreeFunction	0
Age	0
Outcome	0
dtype: int64	

We found no missing values in the dataset, yet independent features like skin thickness, insulin, blood pressure, ;and glucose each have some 0 values, which is practically impossible. A particular column's mean or median scores must be used to replace unwanted 0 values.

## Code

- # Replacing 0 values with the mean of that column
- # Replacing 0 values of Glucose

data['Glucose'] = data['Glucose'].replace(0, data['Glucose'].median())

# Filling 0 values of Blood Pressure

data['BloodPressure'] = data['BloodPressure'].replace(0, data['BloodPressure'].median())

```
# Replacing 0 values in BMI

data['BMI'] = data['BMI'].replace(0, data['BMI'].mean())

# Replacing the missing values of Insulin and SkinThickness

data['SkinThickness'] = data['SkinThickness'].replace(0, data['SkinThickness'].mean())

data['Insulin'] = data['Insulin'].replace(0, data['Insulin'].mean())
```

## Output

data.head()

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	ВМІ	DiabetesPedigreeFunction	Age	Outcome
0	6	148	72	35.000000	79.799479	33.6	0.627	50	1
1	1	85	66	29.000000	79.799479	26.6	0.351	31	0
2	8	183	64	20.536458	79.799479	23.3	0.672	32	1
3	1	89	66	23.000000	94.000000	28.1	0.167	21	0
4	0	137	40	35.000000	168.000000	43.1	2.288	33	1

Let's now examine the data statistics.

### Code

# Reviewing the dataset statistics data.describe()

## Output

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	вмі	DiabetesPedigreeFunction	Age	Οι
count	768.000000	768.000000	768.000000	768.000000	768.000000	768.000000	768.000000	768.000000	76
mean	3.845052	121.656250	72.386719	26.606479	118.660163	32.450805	0.471876	33.240885	0.3
std	3.369578	30.438286	12.096642	9.631241	93.080358	6.875374	0.331329	11.760232	0.4
min	0.000000	44.000000	24.000000	7.000000	14.000000	18.200000	0.078000	21.000000	0.0
25%	1.000000	99.750000	64.000000	20.536458	79.799479	27.500000	0.243750	24.000000	0.0
50%	3.000000	117.000000	72.000000	23.000000	79.799479	32.000000	0.372500	29.000000	0.0
75%	6.000000	140.250000	80.000000	32.000000	127.250000	36.600000	0.626250	41.000000	1.(
max	17.000000	199.000000	122.000000	99.000000	846.000000	67.100000	2.420000	81.000000	1.(
4				'					•

Now our dataset is free of missing and unwanted values.

# **Exploratory Data Analysis**

We will demonstrate analytics using the Seaborn GUI in this tutorial.

### Correlation

Correlation is the relationship between two or more variables. Finding the important features and cleaning the dataset before we begin modelling also helps make the model efficient.

### Code

```
# Correlation plot of the independent variables

plt.figure(figsize = (10, 8))

sns.heatmap(data.corr(), annot = True, fmt = ".3f", cmap = "YlGnBu")

plt.title("Correlation heatmap")
```

## Output



Observations show that characteristics like pregnancy, glucose, BMI, and age are more closely associated with outcomes. I demonstrated a detailed illustration of these aspects in the following phases.

## Pregnancy

#### Code

```
# Exploring Pregnancy and target variables together

plt.figure(figsize = (10, 8))
```

```
# Plotting density function graph of the pregnancies and the target variable

kde = sns.kdeplot(data["Pregnancies"][data["Outcome"] == 1], color = "Red", shade = True)

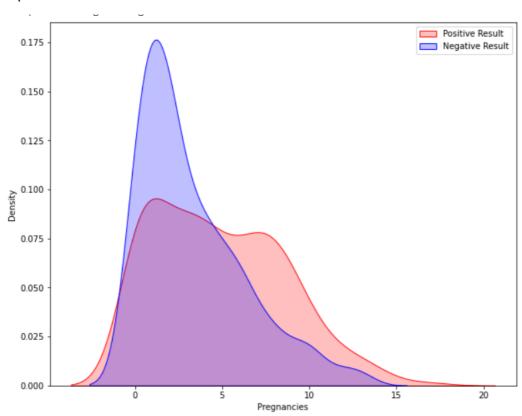
kde = sns.kdeplot(data["Pregnancies"][data["Outcome"] == 0], ax = kde, color = "Blue", shade = True)

kde.set_xlabel("Pregnancies")

kde.set_ylabel("Density")

kde.legend(["Positive Result", "Negative Result"])
```

### Output

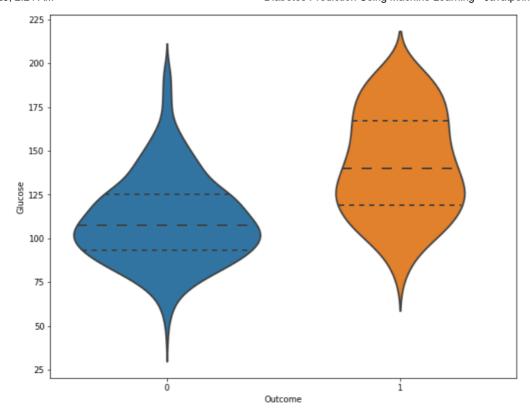


According to the data, women having diabetes have given birth to healthy infants. However, the risk for future complications can be decreased by managing diabetes. The risk of pregnancy issues, such as hypertension, depression, preterm birth, birth abnormalities, and pregnancy loss, is increased if women have uncontrolled diabetes.

## Glucose

```
# Exploring the Glucose and the Target variables together
plt.figure(figsize = (10, 8))
sns.violinplot(data = data, x = "Outcome", y = "Glucose",
split = True, inner = "quart", linewidth = 2)
```

#### Output



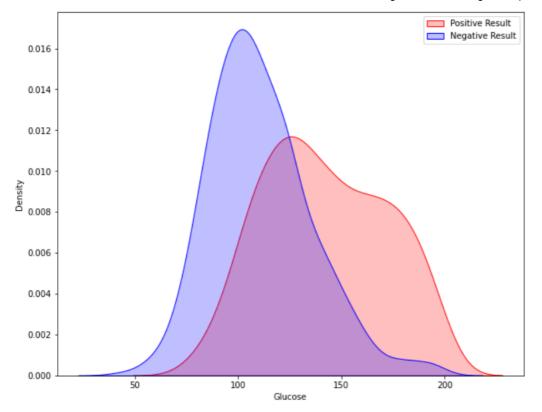
The likelihood of developing diabetes gradually climbs with glucose levels.

### Code

```
# Exploring the density function plot of the Glucose levels

plt.figure(figsize = (10, 8))
kde = sns.kdeplot(data["Glucose"][data["Outcome"] == 1], color = "Red", shade = True)
kde = sns.kdeplot(data["Glucose"][data["Outcome"] == 0], ax = kde, color = "Blue", shade= True)
kde.set_xlabel("Glucose")
kde.set_ylabel("Glucose")
kde.legend(["Positive Result","Negative Result"])
```

### Output



## Implementing Machine Learning Models

We will test many machine learning models and compare their accuracy in this part. After that, we will tune the hyperparameters on models with good precision.

We will use sklearn.preprocessing to convert the data into quantiles before dividing the dataset.

## Code

# Transforming the data into quartiles

quartile = QuantileTransformer()

X = quartile.fit\_transform(data)

dataset = quartile.transform(X)

dataset = pd.DataFrame(X)

 $dataset. columns = \hbox{['Pregnancies', 'Glucose', 'BloodPressure', 'SkinThickness', 'Insulin', 'BMI', 'DiabetesPedigreeFunction', 'Age', 'Outcome']}$ 

# Showing the top 5 rows of the transformed dataset

dataset.head()

## Output

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	ВМІ	DiabetesPedigreeFunction	Age	Outcome
0	0.747718	0.810300	0.494133	0.801825	0.380052	0.591265	0.750978	0.889831	1.0
1	0.232725	0.091265	0.290091	0.644720	0.380052	0.213168	0.475880	0.558670	0.0
2	0.863755	0.956975	0.233377	0.308996	0.380052	0.077575	0.782269	0.585398	1.0
3	0.232725	0.124511	0.290091	0.505867	0.662973	0.284224	0.106258	0.000000	0.0
4	0.000000	0.721643	0.005215	0.801825	0.834420	0.926988	0.997392	0.606258	1.0

## **Data Splitting**

We will now divide the data into a training and testing dataset. We will use the training and testing datasets to train and evaluate different models. We will also perform cross-validation for multiple models before predicting the testing data.