

# Document File Containing the Working of Model

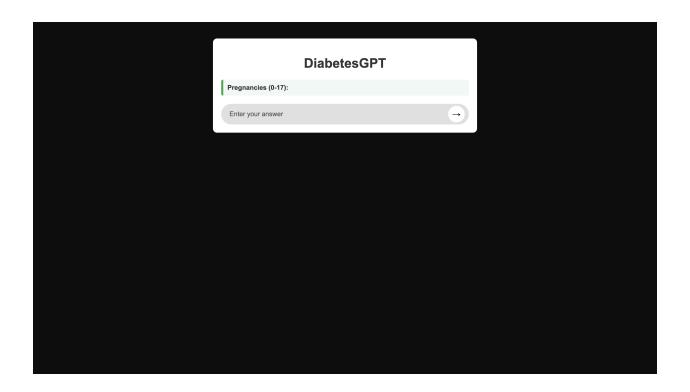
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BATCH: 11

### Working of the Flask App:



- $\rightarrow$  This is the home page of the flask app , i.e. , <u>app.py</u> which is linked with my Machine Learning Model , i.e. , Diabetic-Model.pkl.
- → Now the user enters real time data in the flask app for each features.
- First we enter that data that should predict Diabetes in the final predication.

Pregnancies: 2

Glucose: 120

**Blood Pressure: 80** 

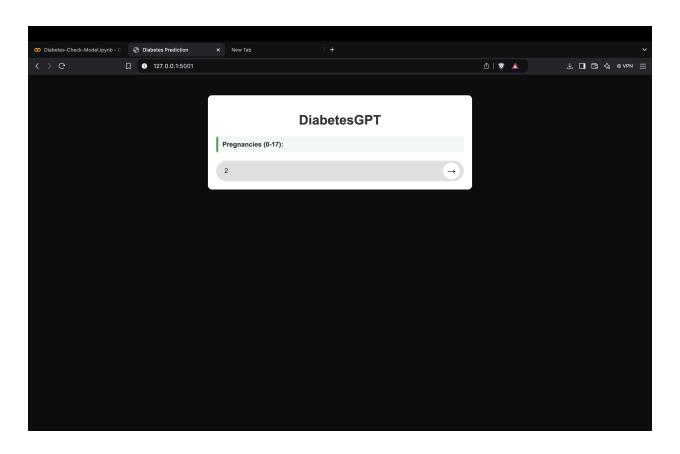
Skin Thickness: 28.5

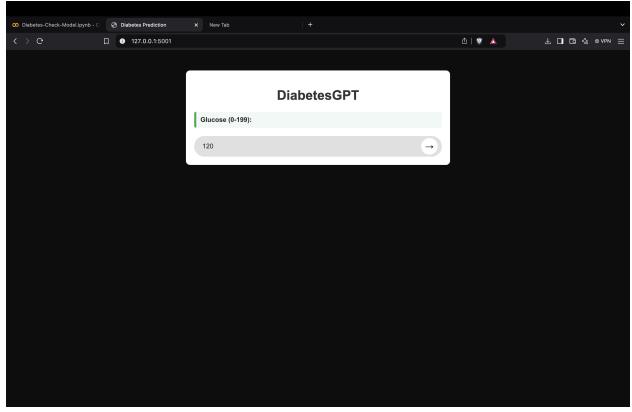
Insulin: 80

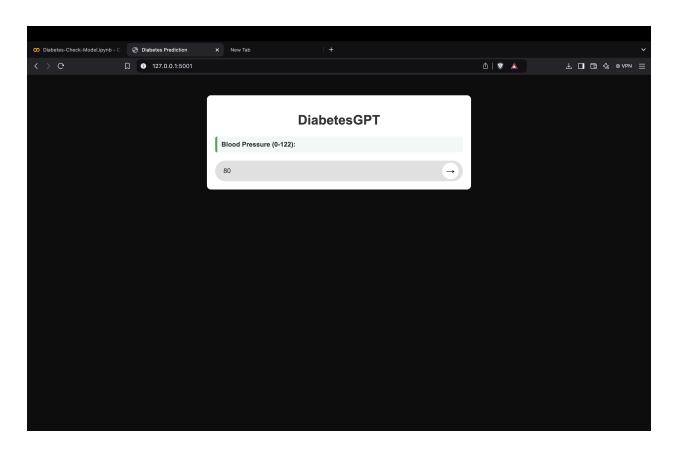
**BMI: 28** 

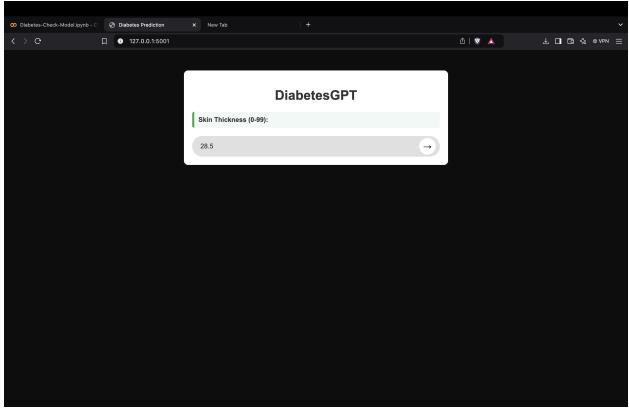
**Diabetic Pedigree Function: 0.45** 

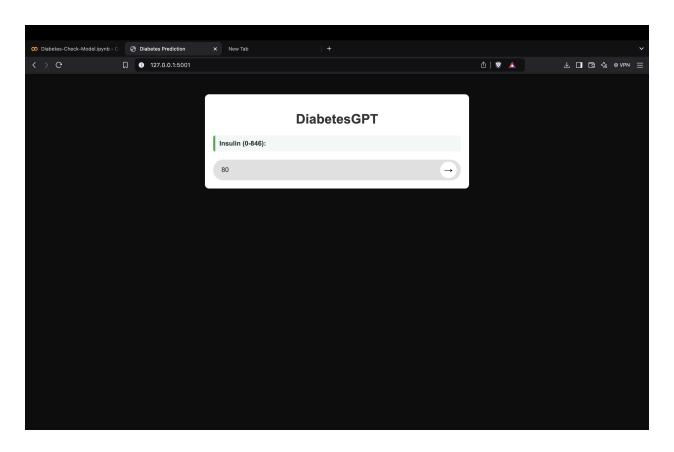
Age: 45

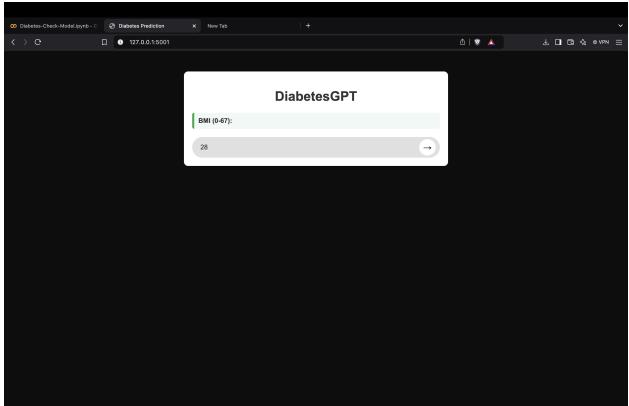


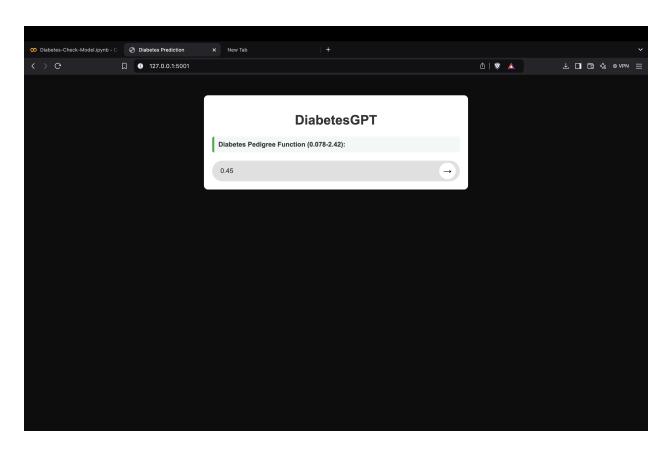


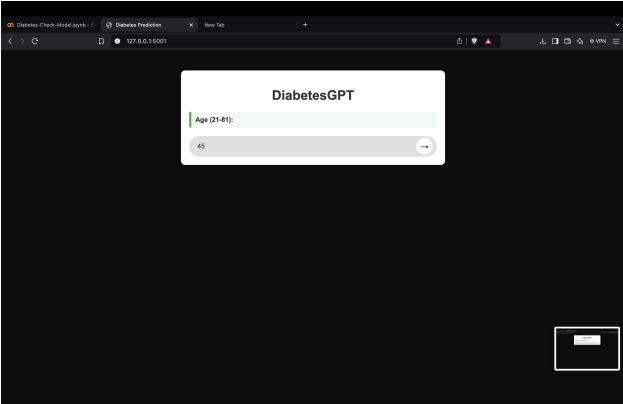


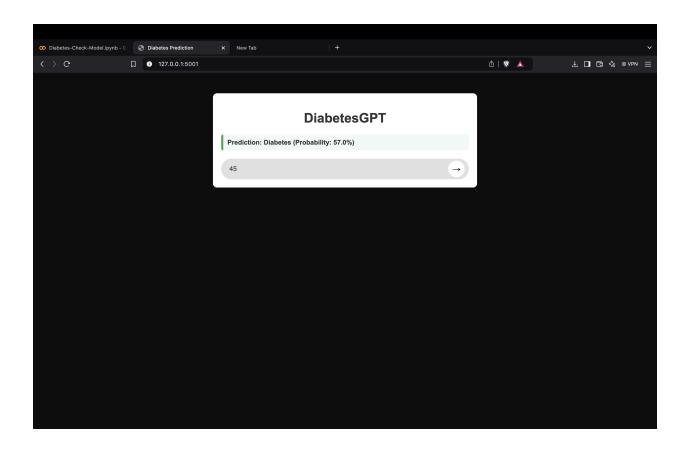












Hence it predicts: 57.0 %, i.e., chances of this person getting diabetic!

At the backend , we also do console.log to see that user entered data is successfully getting extracted for the model prediction or not :

```
Description

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```

## Now we look into our model working:

1. Install required libraries:

```
# Install required libraries for imbalanced learning and XGBoost
!pip install -q imbalanced-learn xgboost
```

- This cell installs two libraries:
  - imbalanced-learn: Used for handling imbalanced datasets, particularly with techniques like SMOTE (Synthetic Minority Over-sampling Technique).
  - xgboost: A library for the XGBoost model, which is often used in classification tasks due to its high performance with structured data.

The q flag is used to suppress unnecessary output during the installation.

```
# Import necessary libraries
import pandas as pd
import numpy as np
from sklearn.model_selection import KFold, cross_val_score, tra:
from sklearn.linear_model import LogisticRegression
from sklearn.ensemble import RandomForestClassifier
from sklearn.svm import SVC
from sklearn.neighbors import KNeighborsClassifier
from xgboost import XGBClassifier
from sklearn.neural network import MLPClassifier
from sklearn.metrics import accuracy_score, precision_score, red
from imblearn.over_sampling import SMOTE
from sklearn.preprocessing import StandardScaler
import warnings
import joblib
from google.colab import files
```

- Imports various libraries and modules:
  - pandas and numpy: Essential libraries for data handling and numerical operations.
  - sklearn.model\_selection: Provides tools for splitting data (train\_test\_split), k-fold cross-validation (KFold), and evaluating models (cross\_val\_score).
  - sklearn.linear\_model, ensemble, svm, neighbors, neural\_network: Includes classifiers like Logistic Regression, Random Forest, SVM, K-Nearest Neighbors, and Neural Network, all used later in model comparisons.
  - XGBClassifier: XGBoost classifier from xgboost, a high-performing model commonly used in machine learning competitions.
  - sklearn.metrics: For evaluating model performance using metrics like accuracy, precision, recall, F1 score, and ROC AUC score.
  - SMOTE: Oversampling technique from imbalanced-learn to balance the class distribution.
  - StandardScaler: Used to scale features for improved model performance.

- warnings: To control warning messages.
- joblib: For saving trained models.
- o google.colab.files: Allows downloading files when using Google Colab.

#### 3. Suppress the specific UserWarning:

```
# Suppress the specific UserWarning from sklearn warnings.filterwarnings("ignore", category=UserWarning, module=
```

Suppresses Userwarning from Sklearn.base, ensuring these warnings do not clutter the output. This can be helpful when some warnings aren't relevant or may confuse the user.

#### 4. Load Dataset:

```
# Load Dataset
url = 'https://raw.githubusercontent.com/plotly/datasets/master,
try:
    data = pd.read_csv(url)
except Exception as e:
    print("Error loading data:", e)

# Data Preprocessing
# Handle missing values and separate features/target
for column in ['Glucose', 'BloodPressure', 'SkinThickness', 'Institute data.loc[data[column] == 0, column] = data[column].median()

X = data.drop('Outcome', axis=1)
y = data['Outcome']

# Balance classes using SMOTE
smote = SMOTE(random_state=42)
X, y = smote.fit_resample(X, y)
```

```
# Scale features
scaler = StandardScaler()
X = scaler.fit_transform(X)
```

- Data Loading: Loads the dataset from a URL.
  - pd.read\_csv(url): Reads the CSV file from the given URL. If loading fails, an error message is printed.

#### Data Preprocessing:

- **Missing Value Handling:** For selected columns, any instance of (which is likely erroneous for attributes like Glucose and BMI) is replaced with the median of the column.
- **Feature/Target Split**: Splits the data into x (features) and y (target variable outcome).

#### • Class Balancing:

• Uses **SMOTE** to balance classes in **y**, which can help improve model performance on imbalanced datasets.

#### Feature Scaling:

- Scales features to standardize them (mean=0, variance=1), which helps models like logistic regression and neural networks.
- 5. Split Dataset for Training and Testing Purpose:

```
# Split the dataset into training and testing sets
X_train, X_test, y_train, y_test = train_test_split(X, y, test_s)
# Check the shapes of the training and testing sets
X_train.shape, X_test.shape, y_train.shape, y_test.shape
```

• **Train-Test Split**: Splits data into training and testing sets (80-20 split) with train\_test\_split.

- stratify=y ensures both sets have the same proportion of classes, preserving the class distribution.
- **Shape Check**: Outputs the shapes of the training and testing sets, verifying the split was done correctly.
- 6. Comparison Of Various Models for Comparison:

```
# Define Models for Comparison
models = {
    'Logistic Regression': LogisticRegression(max iter=200),
    'Random Forest': RandomForestClassifier(),
    'Support Vector Machine': SVC(probability=True),
    'K-Nearest Neighbors': KNeighborsClassifier(),
    'XGBoost': XGBClassifier(use label encoder=False, eval metri
    'Neural Network': MLPClassifier(max_iter=300)
}
# Model Training and Evaluation with K-Fold Cross Validation
kf = KFold(n_splits=5, shuffle=True, random_state=42)
results = {}
for model_name, model in models.items():
    with np.errstate(divide='ignore', invalid='ignore'):
        accuracy = cross_val_score(model, X_train, y_train, cv=l
        precision = cross val score(model, X train, y train, cv:
        results[model_name] = {'Accuracy': accuracy, 'Precision'
```

#### Model Definitions:

• Initializes several models in a dictionary (models) to compare their performance on the data.

#### K-Fold Cross-Validation:

• KFold is set up to split data into 5 subsets for cross-validation.

#### Model Evaluation:

- For each model, <a href="mailto:cross\_val\_score">cross\_val\_score</a> calculates average <a href="mailto:accuracy">accuracy</a> and <a href="mailto:precision">precision</a> across folds.
- Results are stored in results, associating each model with its performance metrics.

#### 7. Displaying Precision and Accuracy Results For Each Model:

```
# Displaying results for each model
for model_name, metrics in results.items():
    print(f"{model_name}:")
    print(f" Accuracy: {metrics['Accuracy']:.2f}")
    print(f" Precision: {metrics['Precision']:.2f}")
```

#### Displaying Results:

- Iterates over the results dictionary.
- For each model, it prints the Accuracy and Precision scores, formatted to two decimal places.
- This helps in identifying the models that perform best on both metrics.

#### 8. Train and Save the best performing model:

```
# Train and save the best-performing model
best_model = LogisticRegression(max_iter=200)
best_model.fit(X_train, y_train)

# Save the trained model
joblib.dump(best_model, "diabetes_model.pkl")
```

#### Selecting and Training Best Model:

Chooses LogisticRegression as the best model based on previous results.
 This choice may be based on an assumption, or a selection process from prior cells could guide it.

• The model is trained using x\_train and y\_train.

#### Saving the Model:

- o joblib.dump saves the trained model as diabetes\_model.pkl.
- This serialized model file can be loaded later for making predictions without retraining.
- 9. Test the prediction probability of the model:

```
# Test the Best Model on Sample Input Data
# Here, we select the best model based on accuracy and precision
best model = models[best model name]
best_model.fit(X_train, y_train) # Train on the full training (
try:
    # Collect user inputs for each feature
    user input = []
    user_input.append(float(input("Pregnancies (0-17): ")))
    user_input.append(float(input("Glucose (0-199): ")))
    user_input.append(float(input("Blood Pressure (0-122): ")))
    user_input.append(float(input("Skin Thickness (0-99): ")))
    user_input.append(float(input("Insulin (0-846): ")))
    user input.append(float(input("BMI (0-67): ")))
    user_input.append(float(input("Diabetes Pedigree Function ()
    user_input.append(float(input("Age (21-81): ")))
    # Convert to numpy array and scale the input
    user_input_scaled = scaler.transform(np.array([user_input]))
    # Make prediction
    with np.errstate(divide='ignore', invalid='ignore'):
        prediction = best_model.predict(user_input_scaled)
        prediction_proba = best_model.predict_proba(user_input_s)
```

```
# Output prediction and probability
print(f"\nSample Prediction (0 = No Diabetes, 1 = Diabetes)
print(f"Probability of Diabetes: {prediction_proba[0][1]:.21
except ValueError:
    print("Invalid input. Please enter numeric values.")
```

- best\_model = models[best\_model\_name]:
  - Selects the best-performing model based on previously calculated metrics (accuracy and precision).
  - best\_model\_name is likely the name of the model with the highest evaluation scores.
- best\_model.fit(X\_train, y\_train):
  - Retrains the best model on the entire training set to ensure it has the most complete information before making predictions.
  - This final training step is done after K-Fold validation to fine-tune the model.
- Initializes an empty list user\_input.
- Each line prompts the user to enter a specific feature value for prediction:
  - Pregnancies, Glucose, Blood Pressure, Skin Thickness, Insulin, BMI,
     Diabetes Pedigree Function, Age
  - Each feature input is converted to float to ensure numeric values, and all inputs are appended to the user\_input list.
- Expected ranges are provided in parentheses to guide valid input values.
- np.array([user\_input]) converts the list of user inputs to a numpy array, compatible with the scaler.
- scaler.transform(...) applies scaling to the input, standardizing it according to the scaling parameters derived from the training data.
- user\_input\_scaled now contains the scaled values ready for prediction.

- **Suppressing Warnings**: The np.errstate(divide='ignore', invalid='ignore') context manager suppresses warnings (e.g., division by zero or invalid operations) during prediction.
- best\_model.predict(user\_input\_scaled):
  - Predicts the class (0 or 1) for diabetes.
- best\_model.predict\_proba(user\_input\_scaled):
  - Calculates the probability for each class. <a href="mailto:proba[0][1]">proba[0][1]</a> accesses the probability of the positive class (diabetes).
- Outputs the class prediction and the probability of diabetes.
- prediction[0]: Prints the result (0 for non-diabetic, 1 for diabetic).
- prediction\_proba[0][1]:.2f: Prints the probability with two decimal places for readability.
- Catches any ValueError (e.g., if the user inputs a non-numeric value).
- Provides a helpful error message to prompt users for correct input.