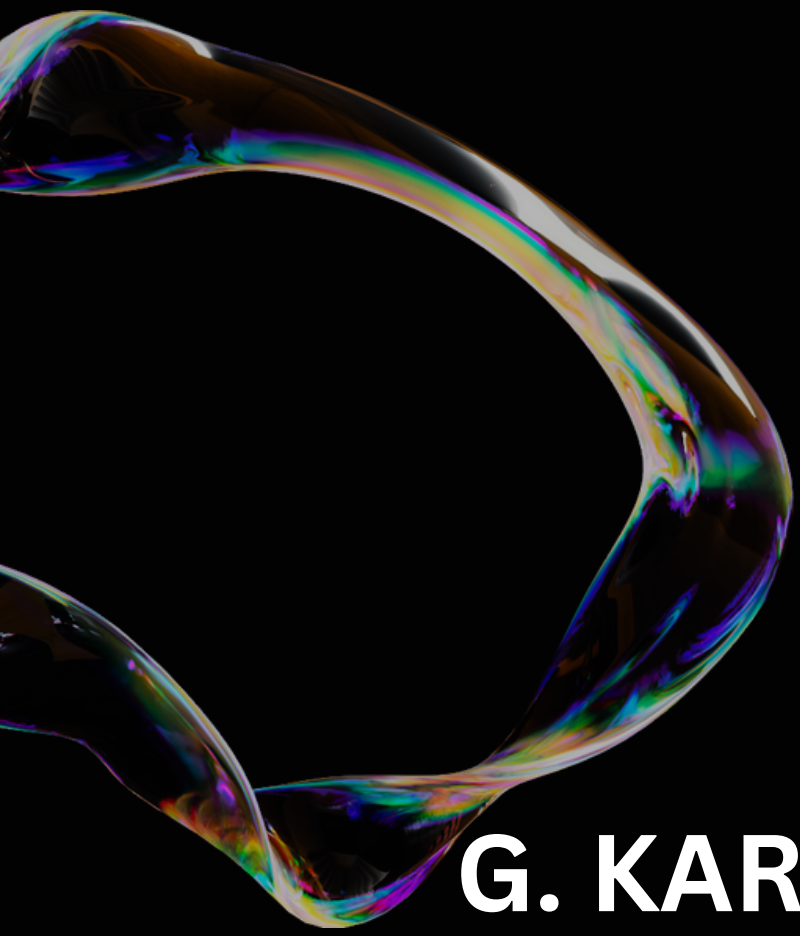
AI BASED DIABETES PREDICTION SYSTEM



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```
IMPORT DATASET
DF = PD.READ_CSV("../INPUT/PIMA-INDIANS-DIABETES-DATABASE/DIABETES.CSV")
# GET FAMILIER WITH DATASET STRUCTURE
DF.INFO()
```

```
lass 'pandas.core.frame.DataFrame'>
ngeIndex: 768 entries, 0 to 767
ta columns (total 9 columns):
    Column                    Non-Null Count  Dtype
-   ------                    --------------  -----
    Pregnancies               768 non-null    int64
    Glucose                   768 non-null    int64
    BloodPressure             768 non-null    int64
    SkinThickness             768 non-null    int64
    Insulin                   768 non-null    int64
    BMI                       768 non-null    float64
    DiabetesPedigreeFunction  768 non-null    float64
    Age                       768 non-null    int64
    Outcome                   768 non-null    int64
ypes: float64(2), int64(7)
mory usage: 54.1 KB
```

```
DF['GLUCOSE'] = DF['GLUCOSE'].REPLACE(0,
DF['GLUCOSE'].MEAN())
# CORRECTING MISSING VALUES IN BLOOD
PRESSURE
DF['BLOODPRESSURE'] =
DF['BLOODPRESSURE'].REPLACE(0,
DF['BLOODPRESSURE'].MEAN()) # THERE
ARE 35 RECORDS WITH 0 BLOODPRESSURE
IN DATASET
# CORRECTING MISSING VALUES IN BMI
DF['BMI'] = DF['BMI'].REPLACE(0,
DF['BMI'].MEDIAN())
```

```
# Data Transformation
q  = QuantileTransformer()
X = q.fit_transform
transformedDF = q.transform(X)
transformedDF = pd.DataFrame(X)
transformedDF.columns =
['Pregnancies', 'Glucose',
'BloodPressure', 'SkinThickness',
'Insulin', 'BMI',
'DiabetesPedigreeFunction', 'Age',
'Outcome']
# Show top 5 rows
transformedDF.head
```

```
import pandas as pd
from sklearn.model_selection
import train_test_split
from sklearn.preprocessing
import StandardScaler,
LabelEncoder

# Load the dataset
data =
pd.read_csv('your_dataset.csv')

# Data Exploration (Optional)
# You can explore
```

Data Cleaning

```
# Handle missing values and remove
duplicates if necessary
```

```
# Data Preprocessing
# For example, encoding categorical variables
label_encoder = LabelEncoder()
data['categorical_column'] = label_encoder

X = data.drop('target_column', axis=1)  # Features
y = data['target_column']  # Target variable
```