

**ST JOSEPH COLLEGE OF ENGINEERING**

**TITLE : AI-BASED DIABETICS  
PREDICTION MODEL  
PHASE-5**

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# TITLE : AI-BASED DIABETICS PREDICTION MODEL

## C O D E

### Step 1 – Importing Modules

Now, let's import the necessary **Python libraries** into our notebook.

**Keras API** already includes Python's TensorFlow deep learning package, which is critical in the diabetes prediction challenge

CODE:

```
import numpy as np
import pandas as pd
import tensorflow as tf
from keras.layers import Dense,Dropout
from sklearn.model_selection import
train_test_split
import matplotlib as mlp
import matplotlib.pyplot as plt
%matplotlib inline
from sklearn.preprocessing import StandardScaler
```

# CODE

## Step 2 – Loading the Dataset

We are now ready to begin importing the dataset. In the next piece of code, we import the dataset and use the `head()` method to get the top five data points.

CODE:

```
data=pd.read_csv("pima-indians-  
diabetes.csv")  
data.head()
```

	6	148	72	35	0	33.6	0.627	50	1
0	1	85	66	29	0	26.6	0.351	31	0
1	8	183	64	0	0	23.3	0.672	32	1
2	1	89	66	23	94	28.1	0.167	21	0
3	0	137	40	35	168	43.1	2.288	33	1
4	5	116	74	0	0	25.6	0.201	30	0

the diabetics and their related documented data are taken from the exact code provided



# C O D E

## Step 3 – Renaming the Columns

You've probably realized that the columns are meaningless

CODE:

head() in Pandas

```
data = data.rename(index=str, columns={"6":"preg"})
data = data.rename(index=str,
columns={"148":"gluco"})data =
data.rename(index=str, columns={"72":"bp"})
data = data.rename(index=str,
columns={"35":"stinmm"})data =
data.rename(index=str, columns={"0":"insulin"})
data = data.rename(index=str,
columns={"33.6":"mass"})data
=data.rename(index=str, columns={"0.627":"dpf"})
data = data.rename(index=str, columns={"50":"age"})
data = data.rename(index=str,
columns={"1":"target"})
data.head()
```

### Step 3:

	preg	gluco	bp	stinmm	insulin	mass	dpf	age	target
0	1	85	66	29	0	26.6	0.351	31	0
1	8	183	64	0	0	23.3	0.672	32	1
2	1	89	66	23	94	28.1	0.167	21	0
3	0	137	40	35	168	43.1	2.288	33	1
4	5	116	74	0	0	25.6	0.201	30	0

the previous schedule of this table is modified by adding some names to the rows and columns to easily recognise what happens exactly and whats the data given to it

## Step 4 – Separating Inputs and Outputs

C  
O  
D  
E

Code:

```
X = data.iloc[:, :-1]
```

```
Y = data.iloc[:, 8]
```

The **X** and **Y** values look somewhat like this:

We separated our dataset into input and target datasets, which implies that the first eight

columns will serve as input features for our model and the last column will serve as the target class.



```
print(X)
```

	preg	gluco	bp	stinmm	insulin	mass	dpf	age
0	1	85	66	29	0	26.6	0.351	31
1	8	183	64	0	0	23.3	0.672	32
2	1	89	66	23	94	28.1	0.167	21
3	0	137	40	35	168	43.1	2.288	33
4	5	116	74	0	0	25.6	0.201	30
..	..	..	..	..	..	..	..	..
762	10	101	76	48	180	32.9	0.171	63
763	2	122	70	27	0	36.8	0.340	27
764	5	121	72	23	112	26.2	0.245	30
765	1	126	60	0	0	30.1	0.349	47
766	1	93	70	31	0	30.4	0.315	23

[767 rows x 8 columns]

```
print(Y)
```

0	0
1	1
2	0
3	1
4	0
..	..
762	0
763	0
764	0
765	1
766	0

Name: target, Length: 767, dtype: int64

the collected data is now allocated with **x,y** valuse in the represented table which allocates the source of the **csv file**

## Step 5 – Train-Test Split of the Data

The next step involves the training and testing **split into data** and then standardizing the data to make computations simpler later on

C

code:

O

```
X_train_full, X_test, y_train_full, y_test =  
train_test_split(X, Y, random_state=42)
```

D

```
X_train, X_valid, y_train, y_valid =  
train_test_split(X_train_full, y_train_full,  
random_state from sklearn.preprocessing
```

E

```
import StandardScaler  
scaler = StandardScaler()
```

```
X_train = scaler.fit_transform(X_train)X_valid  
= scaler.transform(X_valid)  
X_test = scaler.transform(X_test)
```



# C O D E

## **Step 6** – Building the Model

We start off by using a random seed to generate a pseudo-random number and setting it to the tf graph. Then, we will be using a sequential model, and also some dropout layers in the model to avoid overfitting of the data.

**code:**

```
np.random.seed(42)
tf.random.set_seed(42)
model=Sequential()
model.add(Dense(15,input_dim=8, activation='relu'))
model.add(Dense(10,activation='relu'))
model.add(Dense(8,activation='relu'))
model.add(Dropout(0.25))
model.add(Dense(1, activation='sigmoid'))
```

# C O D E

## **Step 7** – Training and Testing of the Model

Now, let's move forward to train our model and then fit the model on the testing dataset.

**code:**

```
model.compile(loss="binary_crossentropy",  
optimizer="SGD", metrics=['accuracy'])  
model_history = model.fit(X_train, y_train,  
epochs=200, validation_data=(X_valid,  
y_valid))
```

You will realize that will train the model for 200 epochs and use binary-cross entropy loss function and SGD optimizer.

step:8

## The conclusion:

The **AI-based diabetes prediction system** represents a promising advancement in the field of healthcare technology. By leveraging sophisticated algorithms and machine learning techniques, it has demonstrated **its potential to accurately forecast the likelihood of diabetes onset in individuals**. This system holds the potential to revolutionize early intervention and preventative care strategies, ultimately improving the quality of life for those at risk of developing diabetes. However, it is essential to continue refining the model, validating its predictions through extensive clinical trials, and ensuring its seamless **integration into existing healthcare workflows**. With further development and implementation, this AI system has the potential to significantly impact public health outcomes and contribute to a more proactive approach in managing diabetes.





# THANK YOU !!

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