

(LETS GROW TOGETHER)

An PROJECT

on

"DIABETES CLASSIFICATION"

Submitted in partial fulfillment for the INTERNSHIP

BATCH 23

IN

MACHINE LEARNING DOMAIN

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TECHNOHACKS EDUTECH INTERNSHIP PROGRAM
2023

DIABETES CLASSIFICATION:

use a dataset containing medical data of pateints to predict if a person has diabetes or not.

Certainly! To build a model to predict whether a person has diabetes or not using a dataset containing medical data, you can follow these general steps. In this example, I'll use the popular diabetes dataset from scikit-learn.

CODE:

```
# Import necessary libraries
import numpy as np
import pandas as pd
from sklearn.model selection import train test split
from sklearn.preprocessing import StandardScaler
from sklearn.linear model import LogisticRegression
from sklearn.metrics import accuracy score, classification report,
confusion matrix
# Load the diabetes dataset
from sklearn.datasets import load diabetes
diabetes = load diabetes()
data = pd.DataFrame(data=np.c [diabetes['data'], diabetes['target']],
columns=diabetes['feature names'] + ['target'])
# Assuming the target variable is binary (1 for diabetes, 0 for no diabetes)
data['target'] = data['target'].apply(lambda x: 1 if x > 150 else 0)
# Split the data into features (X) and target variable (y)
X = data.drop('target', axis=1)
y = data['target']
```

```
# Split the data into training and testing sets
X train, X test, y train, y test = train test split(X, y, test size=0.2,
random state=42)
# Standardize the features
scaler = StandardScaler()
X train scaled = scaler.fit transform(X train)
X test scaled = scaler.transform(X test)
# Build and train the Logistic Regression classifier
logreg classifier = LogisticRegression(random state=42)
logreg classifier.fit(X train scaled, y train)
# Make predictions on the test set
y pred = logreg classifier.predict(X test scaled)
# Evaluate the model
accuracy = accuracy score(y test, y pred)
conf matrix = confusion matrix(y test, y pred)
class report = classification report(y test, y pred)
print(f'Accuracy: {accuracy}')
print(f'Confusion Matrix:\n{conf matrix}')
print(f'Classification Report:\n{class report}')
```

Note: This example assumes that the target variable in the diabetes dataset is continuous. In this case, I've transformed it into a binary variable based on a threshold value (150). You may need to adapt this threshold or use a different approach depending on your specific dataset.

Also, replace the dataset and preprocessing steps with your actual medical dataset. Ensure that your dataset has appropriate features and a binary target variable indicating the presence or absence of diabetes.

OUTPUT:

```
Accuracy: 0.7640449438202247
Confusion Matrix:
[[42 7]
[14 26]]
Classification Report:
        precision recall f1-score support
           0.75
      0
                  0.86
                         0.80
                                 49
           0.79
                  0.65
                         0.71
                                 40
      1
  accuracy
                                 89
                         0.76
                            0.76
               0.77
                      0.75
                                     89
 macro avg
weighted avg
                                      89
               0.77
                       0.76 0.76
```