Classification

```
# import essential libraries
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
%matplotlib inline
# load the dataset
data = pd.read_csv('/content/diabetes_risk_prediction_dataset.csv')
# check the dimension of the dataset
print('Dimension of the dataset: ', data.shape)
# check the attributes in the dataset
print('Attributes in the dataset: ', data.columns.values)
# view the first 5 rows of the dataset
data.head()
Dimension of the dataset: (520, 17)
     Attributes in the dataset: ['Age' 'Gender' 'Polyuria' 'Polydipsia' 'sudden weight loss' 'weakness' 'Polyphagia' 'Genital thrush' 'visual blurring' 'Itching' 'Irritability' 'delayed healing' 'partial paresis' 'muscle stiffness' 'Alopecia'
       'Obesity' 'class']
```

	Age	Gender	Polyuria	Polydipsia	sudden weight loss	weakness	Polyphagia	Genital thrush	visual blurring	Itching	Irritability	delayed healing	partial paresis	mu stiff
0	40	Male	No	Yes	No	Yes	No	No	No	Yes	No	Yes	No	
1	58	Male	No	No	No	Yes	No	No	Yes	No	No	No	Yes	
2	41	Male	Yes	No	No	Yes	Yes	No	No	Yes	No	Yes	No	
3	45	Male	No	No	Yes	Yes	Yes	Yes	No	Yes	No	Yes	No	
4	60	Male	Yes	Yes	Yes	Yes	Yes	No	Yes	Yes	Yes	Yes	Yes	

data.info()

```
<class 'pandas.core.frame.DataFrame'>
     RangeIndex: 520 entries, 0 to 519
    Data columns (total 17 columns):
     # Column
                            Non-Null Count Dtype
     ---
                            520 non-null
     0 Age
                                           int64
     1
         Gender
                            520 non-null
                                           object
                           520 non-null
     2
         Polyuria
                                           object
     3
         Polydipsia
                            520 non-null
                                           object
         sudden weight loss 520 non-null
                                           object
                      520 non-null
520 non-null
                                           object
         Polyphagia
                                           object
         Genital thrush
                            520 non-null
                                           object
         visual blurring 520 non-null
                                           object
                           520 non-null
520 non-null
         Itching
                                           object
     10 Irritability
                                           obiect
     11 delayed healing 520 non-null
                                           object
     12 partial paresis
                            520 non-null
                                           object
     13
         muscle stiffness 520 non-null
                                           object
     14 Alopecia
                            520 non-null
                                           object
     15 Obesity
                            520 non-null
                                           obiect
                             520 non-null
     16 class
                                           object
    dtypes: int64(1), object(16)
    memory usage: 69.2+ KB
from sklearn.model_selection import train_test_split
train, test = train_test_split(data, test_size=0.2, random_state=122)
# dimension of train and test dataset
print('Dimension of training data: ', train.shape)
print('Dimension of test data: ', test.shape)
    Dimension of training data: (416, 17)
    Dimension of test data: (104, 17)
```

```
# segregate the feature matrix and target vector from train and test data
Xtrain = train.drop(columns=['class'], axis=1)
ytrain = train['class']
Xtest = test.drop(columns=['class'], axis=1)
ytest = test['class']
\mbox{\tt\#} encode the target/label for train and test dataset
from sklearn.preprocessing import LabelEncoder
encoder = LabelEncoder()
ytrain_encoded = encoder.fit_transform(ytrain)
ytest_encoded = encoder.transform(ytest)
from \ sklearn.compose \ import \ ColumnTransformer
from sklearn.pipeline import Pipeline
from \ sklearn.preprocessing \ import \ One HotEncoder, \ MinMaxScaler
# extracted the list of categorical columns to be encoded using OneHotEncoder
excluded_col = 'Age'
categorical_col = [col for col in Xtrain.columns if col != excluded_col]
# define the column transformer
preprocessor = ColumnTransformer(
    transformers=[
        ('cat', OneHotEncoder(drop='first'), categorical_col)
    ٦.
    remainder='passthrough'
)
# create the pipeline
pipeline = Pipeline([
    ('preprocessor', preprocessor),
    ('scaler', MinMaxScaler())
1)
# process the train and test data
Xtrain_transformed = pipeline.fit_transform(Xtrain)
Xtest_transformed = pipeline.transform(Xtest)
## Classification
from \ sklearn.tree \ import \ Decision Tree Classifier
from sklearn.model_selection import GridSearchCV
# create a Decision Tree Classifier
tree = DecisionTreeClassifier(random state=122)
# Define the hyperparameter grid
param_grid = {
    'criterion': ['gini', 'entropy'],
'splitter': ['best', 'random'],
    'max_depth': [10, 20, 30],
    'min_samples_split': [2, 5, 10],
    'min_samples_leaf': [2, 4]
}
# create the GridSearchCV object
grid_search_tree = GridSearchCV(tree, param_grid, cv=5, scoring='accuracy', n_jobs=-1)
# fit the grid search to the data
grid_search_tree.fit(Xtrain_transformed, ytrain_encoded)
# print the best parameters and the corresponding accuracy
print('Best Parameters: ', grid_search_tree.best_params_)
print('Best Accuracy: ', grid_search_tree.best_score_)
# get the best model
best_tree = grid_search_tree.best_estimator_
     Best Parameters: {'criterion': 'entropy', 'max_depth': 10, 'min_samples_leaf': 2, 'min_samples_split': 2, 'splitter': 'best'}
     Best Accuracy: 0.935140562248996
```