3 4 76 76	4 0 137 40 35 168 43.1
76	766 1 126 60 0 0 30.1 767 1 93 70 31 0 30.4 DiabetesPedigreeFunction Age Outcome 9 0.627 50 1 1 0.351 31 0 1 0.672 32 1 1 0.167 21 0
76 76 76 76	
# >	# splitting the dataset into training, validation and testing wihtout normalising the data X_train, X_test, y_train, y_test = train_test_split(X,y,random_state=104,train_size=0.8,shuffle=True) X_train, X_val, y_train, y_val = train_test_split(X_train, y_train, random_state=104, train_size=5/8, shuffle=True) Making the decision Tree
t #	A class Node determines the nodes for the decision tree where each tree stores the entropy of the training data that it has received, the feature that is best to split upon for that node thresfold value to make the split and also what the majority outcome of the dataset is that node # function to find the entropy def entropy(y): x=np.count_nonzero(y==1)/len(y) if x==1 or x==0:
	<pre>return 0 e= -x*np.log2(x) - (1-x)*np.log2(1-x) return e # creating a node class for the tree class NODE :</pre>
	<pre>definit(self,depth,ind): self.id = ind self.type="internal" self.entropy = None self.left = None self.right = None self.feature = None self.label = None self.label = None</pre>
	<pre># finding the feature_split value for a feature def feature_split(X,y,feature) : m = X.shape[0] x=X[:,feature]</pre>
	<pre>y_copy = np.copy(y) ind = np.argsort(x) x=x[ind] y_copy = y_copy[ind] feature_split_list = []</pre>
	<pre>for i in range(1,m) : if y_copy[i]!=y_copy[i-1] : feature_split_list.append((x[i-1]+x[i])/2) entropy_list=[]</pre>
	<pre>for feature_split in feature_split_list : y_left=np.array([]) y_right=np.array([]) for i in range(m): if(x[i]<=feature_split) : y_left=np.append(y_left,y_copy[i]) else :</pre>
	<pre>y_right=np.append(y_right,y_copy[i]) if(len(y_left)==0 or len(y_right) == 0) :</pre>
	<pre>if len(entropy_list) == 0 : return None ,None min_entropy_sum = np.min(entropy_list) t=np.argmin(entropy_list) feature_split = feature_split_list[t] return min_entropy_sum , feature_split</pre>
	<pre># function to best feature def best_feature(X,y) : m=X.shape[0] n=X.shape[1] # stores the smallest entropy possible for each feature</pre>
	<pre>entropy_list=[] label_list = [] for i in range(n) : entropy , label = feature_split(X,y,i) if entropy != None and label != None : entropy_list.append(entropy) label_list.append(label)</pre>
	<pre>min_entropy = np.min(entropy_list) feature = np.argmin(entropy_list) label = label_list[feature] return feature, min_entropy, label</pre>
r	<pre>global node_id node_id=0 # function to build the decision tree def DecisionTree(X, y, depth=0) : m=X.shape[0] n=X.shape[1] ribbal ride</pre>
	<pre>global node_id root = NODE(depth, node_id) node_id = node_id+1 # base case if m <10 or y.ptp() == 0 or n==0 : root.type = "leaf"</pre>
	<pre>values , counts = np.unique(y,return_counts=True) root.pred=values[counts.argmax()] return root root.entropy = entropy(y) feature,entropy_sum,label = best_feature(X,y)</pre>
	<pre>information_gain = root.entropy - entropy_sum root.feature = feature root.label = label values , counts = np.unique(y,return_counts=True) root.pred=values[counts.argmax()] # splitting X and y based on the feature X_left = np.empty((0,n))</pre>
	<pre>X_left = np.empty((0,n)) X_right = np.empty((0,n)) y_left = np.array([]) for i in range(m) : if(X[i][root.feature] <= root.label) : X_left = np.r_[X_left,[X[i]]] y_left = np.append(y_left,y[i]) else :</pre>
	<pre>else : X_right = np.r_[X_right,[X[i]]] y_right = np.append(y_right,y[i]) root.left = DecisionTree(X_left,y_left,depth+1) root.right = DecisionTree(X_right,y_right,depth+1)</pre>
#	<pre>return root root=DecisionTree(X_train, y_train) # code to finding number of nodes in the tree def nodes(root):</pre>
	<pre>if root.type == "leaf" : return 1 return 1 + nodes(root.left) + nodes(root.right) initial_nodes = nodes(root) Testing the decision tree on a Validation set</pre>
t #	The decision tree is then tested on the validation set and we calculate the accuracy precision and recall values for them We also try this for various values of max depth that the test to and based on this we decide whats the best size for our tree. We use this info to prune the tree # making a new prediction function with max depth def predicted(root, x, max_depth=100): if root.type == "leaf" or root.depth == max_depth : rooturn root production production in the control of the cont
	<pre>return root.pred if x[root.feature] <= root.label : pred = predicted(root.left,x,max_depth) else : pred = predicted(root.right,x,max_depth) return pred # obtaining the predicted values over the test set for different max_depths</pre> ## The tree was found to began a double of 10 as we are tablished may depths from 1 to 10.
c a p r	<pre># The tree was found to have a depth of 13 so we are taking max_depths from 1 to 13 depth = [i for i in range(1,14)] accuracy = np.array([]) precision = np.array([]) recall = np.array([]) for max_depth in depth: y_pred = np.array([]) m = X_val.shape[0]</pre>
	<pre>for i in range(m): test = predicted(root, X_val[i], max_depth) y_pred = np.append(y_pred, test) # making the confusion matrix c = np.zeros((2,2)) for i in range(m):</pre>
	<pre>c[y_val[i]][int(y_pred[i])] =c[y_val[i]][int(y_pred[i])] + 1 # calculating accuracy acc = (c[0][0] + c[1][1])/(c[0][0] + c[0][1] + c[1][0] + c[1][1]) accuracy=np.append(accuracy,acc) # calculating precision</pre>
	<pre>pre = c[1][1] / (c[1][1] + c[0][1]) precision=np.append(precision, pre) #calculating recall rec = c[1][1] / (c[1][1] + c[1][0]) recall=np.append(recall, rec) # plotting the accuracy, precision and recall vs max_depth</pre>
p p	<pre>plt.plot(depth, accuracy, label="accuracy") plt.plot(depth, precision, label="precision") plt.plot(depth, recall, label="recall") plt.xlabel("max_depth") plt.ylabel("accuracy/precision/recall") plt.legend() plt.show()</pre>
/recall	0.75 - accuracy — precision — recall
accuracy/precision	0.65 - 0.60 - 0.
	0.55 - 2 4 6 8 10 12 max_depth
#	Now we need to prune this tree so that we dont overfit the model A reduced error pruning is done to achieve this # pruning the tree using reduced error pruning def prune(root): if root.type == "leaf": return prune(root.left)
	<pre>prune(root.right) if root.left.type == "leaf" and root.right.type == "leaf" : left = root.left right = root.right # calculating the accuracy before pruning y_pred = np.array([]) m = X yal_shape[8]</pre>
	<pre>m = X_val.shape[0] for i in range(m) : test = predicted(root, X_val[i]) y_pred = np.append(y_pred, test) # making the confusion matrix c = np.zeros((2,2)) for i in range(m) :</pre>
	<pre>c(y_val[i])[int(y_pred[i])] =c(y_val[i])[int(y_pred[i])] + 1 # calculating accuracy acc_bfr = (c[0][0] + c[1][1])/(c[0][0] + c[0][1] + c[1][0] + c[1][1]) # now make root as the leaf node root.type = "leaf" root.left = None</pre>
	<pre>root.right = None # calculating the accuracy after pruning y_pred = np.array([]) m = X_val.shape[0] for i in range(m) : test = predicted(root, X_val[i])</pre>
	<pre>y_pred = np.append(y_pred, test) # making the confusion matrix c = np.zeros((2,2)) for i in range(m): c[y_val[i]][int(y_pred[i])] = c[y_val[i]][int(y_pred[i])] + 1</pre>
	<pre># calculating accuracy acc_aft = (c[0][0] + c[1][1])/(c[0][0] + c[0][1] + c[1][0] + c[1][1]) if acc_bfr > acc_aft : root.type = "internal" root.left = left root.right = right return</pre>
f #	<pre>return prune(root) final_nodes = nodes(root) # the number of nodes before pruning and after pruning print("Number of nodes before pruning : ",initial_nodes) print("Number of nodes after pruning : ",final_nodes)</pre>
Nu Nu	<pre>print("Number of nodes after pruning : ",Tinal_nodes) Number of nodes before pruning : 83 Number of nodes after pruning : 63 # plotting the tree def plot(root): dot = graphviz.Digraph() dot.node(str(root.id) + "\n" + str(root.type) + "\n" + str(columns[root.feature]) + "<=" + str(root.label))</pre>
	<pre>def add_node_edges(node) : if node.left : if node.left.type == "leaf" :</pre>
	<pre>dot.edge(str(node.id) + "\n" + str(node.type) + "\n" + str(columns[node.feature]) + "<=" + str(node.label), str(node.left.id) + "\n</pre>
	<pre>dot.edge(str(node.id) + "\n" + str(node.type) + "\n" + str(columns[node.feature]) + "<=" + str(node.label), str(node.right.id) + "\</pre>
F	plot(root) Output internal Glucose<=126.0 True False
	$\begin{array}{c c} & & & & & & & \\ & & & & & \\ & & & & & $
	Pregnancies<=6.0 BMI<=26.75 True False
	True False
16	$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$
le	0.0 1.0 BloodPressure<=67.0 0.0 True False True False 32 leaf 1.0 DiabetesPedigreeFunction<=0.34 BloodPressure<=67.0 0.0 Age<=37.5 Insulin<=333.5 Go 61 internal internal Glucose<=163.0 0.0 BloodPressure<=67.0 0.0 Age<=37.5 Insulin<=333.5 Go 65 76 leaf Internal Glucose<=163.0 Glucose<=163.0 0.0 Output DiabetesPedigreeFunction<=0.34 0.0 DiabetesPedigreeFunctio
16	True False True False
16	34 37 1 62 63 1 66 1 1 60 67 70 67 70 67 70 67 70 67 70 67 70 67 70 67 70 67 70 67 70 67 70 67 70 67 70 67 70 67 70 67 70 67 70 67 70 67 70 70
le	34 internal Glucose<=122.5 leaf 1.0 True False True False

print("\n\n\n")

Assignment 1

import graphviz

Name: Navaneeth Shaji

Roll Number: 21CS30032

In []: # import all the necessary libraries here
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler
from sklearn.tree import plot_tree
import graphviz

from IPython.display import Image, display

In []: df = pd.read_csv('../../dataset/decision-tree.csv')
 X = df.iloc[:, :-1].values
 y = df.iloc[:, -1].values