In [1]: import numpy as np import pandas as pd import matplotlib.pyplot as plt import scipy.io as sio from pyitlib import discrete\_random\_variable as drv from tabulate import tabulate from graphviz import Source import os import warnings warnings.filterwarnings('ignore') from sklearn.datasets import load iris from sklearn.metrics import accuracy\_score from sklearn.tree import export graphviz from sklearn.tree import DecisionTreeClassifier **Loading Data** In [2]: data = sio.loadmat('covtype reduced.mat') X train = data['X train'] X test = data['X test'] y\_train = data['y\_train'][0] y\_test = data['y\_test'][0] print (X\_train.shape, X\_test.shape, y\_train.shape, y\_test.shape) data = sio.loadmat('covtype reduced.mat') X train = data['X train'] X test = data['X test'] y\_train = data['y\_train'].T y test = data['y test'].T print (X\_train.shape, X\_test.shape, y\_train.shape, y\_test.shape) (468, 54) (116202, 54) (468,) (116202,) (468, 54) (116202, 54) (468, 1) (116202, 1) computing Entropy and conditionalEntropy In [3]: def entropy(y): **if** len(y) == 0: return 0 unique, count = np.unique(y, return counts=True, axis=0) individualProbabilities = count/len(y) entropy = -np.sum(individualProbabilities\*np.log2(individualProbabilities)) return entropy #Additional Helper function def jointEntropy(y,x):  $yx = np.c_[y,x]$ return entropy(yx) def cond\_entropy(y, yhat): return jointEntropy(y, yhat) - entropy(yhat) random\_sequences = sio.loadmat('random\_sequences.mat') s1 = random\_sequences['s1'][0] s2 = random\_sequences['s2'][0] print ('entropy = ', entropy(s1)) print ('conditional entropy = ', cond\_entropy(s1,s2)) print("\n====Verification with in-built functions") print("verification entropy = :", drv.entropy(s1)) print("verification conditional entropy = :",drv.entropy\_conditional(s1,s2)) entropy = 3.3141823231610834conditional entropy = 3.3029598816135173====Verification with in-built functions verification entropy = : 3.3141823231610834 verification conditional entropy = : 3.3029598816135173 Tree Class and its functions In [4]: class Tree: def \_\_init\_\_(self,  $max_depth = 10,$  $minimum_gain = 1e-7$ , min samples split = 2): self.max\_depth = max\_depth self.minimum\_gain = minimum\_gain self.min\_samples\_split = min\_samples\_split def fit(self, X, y): self.numberOfClasses = np.unique(y).shape[0] self.feature\_importance = np.zeros(X.shape[1]) # 1st call to create the Decision Tree self.tree = getDecisionTree(X, y, self.max\_depth, self.minimum\_gain, self.min samples split, self.numberOfClasses, self.feature\_importance, X.shape[0]) self.feature importance /= np.sum(self.feature importance) return self def predict(self, X): Traverse each row and predict the relevant class. predictions =[] for i in range(X.shape[0]): temp = self.classifyExample(X[i, :], self.tree) predictions.append(temp) return predictions def classifyExample(self, example, tree): classification is done recurssively until you reach the leaf node # base case if tree['is leaf']: return np.argmax(tree['prob']) # recurssion else: featureName, value = tree['split col'], tree['threshold'] splitColumn = tree['split col'] featureType = FEATURE TYPES[splitColumn] # Differentiating between continuous and categorical values if featureType == "continuous": if example[featureName] <= value:</pre> return self.classifyExample(example, tree['left']) else: return self.classifyExample(example, tree['right']) else: if example[featureName] == value: return self.classifyExample(example, tree['left']) else: return self.classifyExample(example, tree['right']) def printTree(self): Helper function to print the tree for debugging. print ('printing tree...') def printNode(parent, tree, childType): if not tree: return if parent is None: print(', ROOT', ) # Differentiating between continuous and categorical values if tree['featureType'] == "continuous": print(', ROOT, Condition: ' +str(tree['colName'])+ '<= '+ str(tree['threshold']))</pre> else: print(', ROOT, Condition: ' +str(tree['colName']) + '== '+ str(tree['threshold'])) # Differentiating between Leaf Node and Non-Leaf Node if tree['is leaf']: print(', LEAF, ', 'Total Samples '+str(sum(tree['counts'])) + ' , Distribution' + str(t ree['counts'])) # Differentiating between continuous and categorical values if tree['featureType'] == "continuous": if childType == "left": print(', NONLEAF, Condition: ' +str(tree['colName']) + '<= '+ str(tree['threshol</pre> **d'**])) print(', NONLEAF, Condition: ' +str(tree['colName']) + '> '+ str(tree['threshol **d'**])) else: if childType == "left": print(', NONLEAF, Condition: ' +str(tree['colName']) + '== '+ str(tree['threshol d'])) print(', NONLEAF, Condition: ' +str(tree['colName']) + '! = ' + str(tree['threshol **d'**])) printNode(tree, tree['left'], "left") printNode(tree, tree['right'], "right") printNode(None, tree.tree,"") **Decision Tree helper functions** In [5]: def getDecisionTree(X, y, max depth, minimum gain, min samples split, numberOfClasses, feature\_importance, n\_row): Recusively constructs a Decision tree 1) Determine if we can split the tree (or) is it a leaf node. 2) If we can split: i) Find best split ii) Recusrivel call Decision tree on both the splits (in our case, it is binary decision tree) 3) If we cannot split, i.e. it is a Leaf Node i) We store the distribution of the data('label') at the leaf node and use this information in our prediction. if max depth>0 and X.shape[0] > min samples split : column, value, informationGain = findBestSplit(X, y) if informationGain > minimum\_gain: feature importance[column] += (X.shape[0] / n row) \* informationGain # computing left and right child left X, right X, left y, right y = splitData(X, y, column, value) left\_child = getDecisionTree(left\_X, left\_y, max depth - 1, minimum gain, min\_samples\_split, numberOfClasses, feature\_importance, n row) right\_child = getDecisionTree(right\_X, right\_y,  $max_depth - 1$ , minimum gain, min samples split, numberOfClasses, feature importance, n\_row) nonLeafNode = { 'is\_leaf': False, 'split\_col': column, 'colName': COLUMN HEADERS[column], 'featureType': FEATURE TYPES[column], 'threshold': value, 'left': left\_child, 'right': right child return nonLeafNode elif X.shape[0] >0 : counts = np.bincount(y, minlength = numberOfClasses) prob = counts / y.shape[0] leafNode = {'is\_leaf': True, 'prob': prob,'counts':counts} return leafNode def findBestSplit(X, y): We try to determine which is the best split for the data based on the Information gain. 1) We find all the unique values for every column, be it continuous (or) categorical 2) We split the data at every unique value for every column and find the information Gain at that value for that column. 3) we pick the column and corresponding split value where we get the maximum information gain. bestSplitColumn, bestSplitValue, maxInformationGain = None, None, None existingEntropy = entropy(y) totalFeatures = X.shape[1] for column in range(totalFeatures): splitValues = np.unique(X[:, column]) for value in splitValues: splits = splitData(X, y, column, value, return\_X = False) informationGain = existingEntropy - computeEntropyAfterSplit(y, splits) if maxInformationGain is None or informationGain > maxInformationGain: bestSplitColumn, bestSplitValue, maxInformationGain = column, value, informationGain return bestSplitColumn, bestSplitValue, maxInformationGain def splitData(X, y, splitColumn, splitValue, return\_X=True): 1) Takes the input data (X,Y). 2) Uses the splitColumn and splitValue to split the data into 2 halves: i) In case of continuous data a) We split them as rowsBelowThreshold and rowsAboveThreshold ii) In case of categorical data (Here we have binary data) a) We split them as rowsWhereValueIsZero and rowsWhereValueIsOne 11 11 11 type\_of\_feature = FEATURE\_TYPES[splitColumn] splitColumnValues = X[:, splitColumn] if type of feature == "continuous": rowsBelowThreshold = splitColumnValues <= splitValue</pre> rowsAboveThreshold = splitColumnValues > splitValue else:#categorical rowsBelowThreshold = splitColumnValues == splitValue rowsAboveThreshold = splitColumnValues != splitValue XBelowThresshold = X[rowsBelowThreshold] YBelowThreshold = y[rowsBelowThreshold] XAboveThresshold = X[rowsAboveThreshold] YAboveThreshold = y[rowsAboveThreshold] if not return X: return YBelowThreshold, YAboveThreshold return XBelowThresshold, XAboveThresshold, YBelowThreshold, YAboveThreshold def entropy(y): Computes entropy for the given column values, counts = np.unique(y, return counts = True) p = counts / y.shape[0] entropy = -np.sum(p \* np.log2(p))return entropy def computeEntropyAfterSplit(y, splits): Computes Entropy of the data after splitting into 2 halves. This information is used to compute the information Gain. splits entropy = 0for split in splits: splits\_entropy += (split.shape[0] / y.shape[0]) \* entropy(split) return splits entropy def getLeafNodeInfo(tree): Gives information about all the leaf nodes present in the tree. Leaf Nodes Information include the following: 1) Total samples in the leaf node 2) Distribution of the samples in the leaf node. 11 11 11 global output output = [] def getDetailedInfo(parent, tree, childType): if not tree: return if tree['is leaf']: a = 'LEAF, ', 'Total Samples '+ "{0:0=3d}".format(sum(tree['counts'])) + ' , Distribution ' + str(tree['counts']) output.append(a) else: getDetailedInfo(tree, tree['left'], "left") getDetailedInfo(tree, tree['right'], "right") getDetailedInfo(None, tree,"") for index, details in enumerate(output): print("Leaf Node "+"{0:0=2d}".format(index+1) + " "+str(details[1])) return def determineTypeOfFeature(data): Determing whether a feature is categorical or continuous Here, we are using our existing knowledge of the dataset, i.e. first 10 columns are continuous and remaining are categorical values. totalColumns = data.shape[1] output = [] for columnIndex in range(totalColumns): if columnIndex<=9:</pre> output.append('continuous') else: output.append('categorical') return output In [6]: data = sio.loadmat('covtype\_reduced.mat') X\_train = data['X\_train'].astype(float) X test = data['X test'].astype(float) y\_train = data['y\_train'][0].astype(int) y\_test = data['y\_test'][0].astype(float) global COLUMN\_HEADERS, FEATURE\_TYPES COLUMN HEADERS = ['Elevation', 'Aspect', 'Slope', 'Horizontal Distance To\_Hydrology', 'Vertical\_Distance e\_To\_Hydrology', 'Horizontal\_Distance\_To\_Roadways', 'Hillshade\_9am', 'Hillshade\_Noon', 'Hillshade\_3pm', 'Horizontal\_Distance\_To\_Fire\_Points', 'Wilderness\_Area1', 'Wilderness\_Area2', 'Wilderness\_Area3', 'Wil derness\_Area4', 'Soil\_Type1', 'Soil\_Type2', 'Soil\_Type3', 'Soil\_Type4', 'Soil\_Type5', 'Soil\_Type6', 'So il\_Type7', 'Soil\_Type8', 'Soil\_Type9', 'Soil\_Type10', 'Soil\_Type11', 'Soil\_Type12', 'Soil\_Type13', 'Soil\_Type14', 'Soil\_Type15', 'Soil\_Type16', 'Soil\_Type17', 'Soil\_Type18', 'Soil\_Type19', 'Soil\_Type20', 'S oil Type21', 'Soil\_Type22', 'Soil\_Type23', 'Soil\_Type24', 'Soil\_Type25', 'Soil\_Type26', 'Soil\_Type27', 'Soil\_Type28', 'Soil\_Type29', 'Soil\_Type30', 'Soil\_Type31', 'Soil\_Type32', 'Soil\_Type33', 'Soil\_Type34' , 'Soil Type35', 'Soil Type36', 'Soil Type37', 'Soil Type38', 'Soil Type39', 'Soil Type40', 'label'] COLUMN\_HEADERS = COLUMN HEADERS FEATURE\_TYPES = determineTypeOfFeature(X\_train) results = [] for depth in range(1,6): tree = Tree(max depth=depth, min samples split=2) tree.fit(X\_train, y\_train) y\_pred = tree.predict(X\_train) y pred2 = tree.predict(X test) results.append((2\*\*depth, "{0:.2f}".format(100\*accuracy\_score(y\_train, y\_pred))+' %', "{0:.2f}".format(100\*(1-accuracy score(y train, y pred)))+' %', "{0:.2f}".format(100\*accuracy\_score(y\_test, y\_pred2))+' %', "{0:.2f}".format(100\*(1-accuracy\_score(y\_test, y\_pred2)))+' %')) columns = ['No of Splits','Train Accuracy','Train Error', 'Test Accuracy', 'Test Error'] df = pd.DataFrame(results, columns=columns) df.reset\_index() print(tabulate(df, headers='keys', tablefmt='psql')) | No of Splits | Train Accuracy | Train Error | Test Accuracy | Test Error | | 0 | 2 | 49.79 % | 50.21 % | 50.40 % | 49.60 % | 1 | 1 | 4 | 62.82 % | 37.18 % | 73.32 % | 26.68 % | 2 | 8 | 63.68 % | 36.32 % | 70.29 % | 29.71 % | 3 | 16 | 67.31 % | 32.69 % | 73.01 % | 26.99 % | 4 | 32 | 72.65 % | 27.35 % | 63.50 % | 36.50 % **Leaf Nodes Info** In [7]: getLeafNodeInfo(tree.tree) Leaf Node 01 Total Samples 001 , Distribution [0 0 0 0 1 0 0] Leaf Node 02 Total Samples 003 , Distribution [0 0 2 1 0 0 0] Leaf Node 03 Total Samples 003 , Distribution [0 0 0 3 0 0 0] Leaf Node 04 Total Samples 001 , Distribution [0 0 1 0 0 0 0] Leaf Node 05 Total Samples 004 , Distribution [0 0 0 4 0 0 0] Leaf Node 06 Total Samples 006 , Distribution  $[0\ 0\ 0\ 1\ 0\ 0\ 5]$ Leaf Node 07 Total Samples 019 , Distribution [ 0 0 0 15 0 Leaf Node 08 Total Samples 009 , Distribution [0 0 6 0 0 0 3] Leaf Node 09 Total Samples 004 , Distribution [0 0 0 4 0 0 0] Leaf Node 10 Total Samples 026 , Distribution [ 0  $\,$  4 22  $\,$  0  $\,$  0 Leaf Node 11 Total Samples 009 , Distribution [0 1 4 0 0 0 4] Leaf Node 12 Total Samples 001 , Distribution [0 0 1 0 0 0] Leaf Node 13 Total Samples 009 , Distribution [0 0 9 0 0 0 0] Leaf Node 14 Total Samples 007 , Distribution [0 2 4 0 0 1 0] Leaf Node 15 Total Samples 014 , Distribution [0 0 4 6 0 4 0] Leaf Node 16 Total Samples 003 , Distribution [0 3 0 0 0 0] Leaf Node 17 Total Samples 055 , Distribution [ 0 11 42 0 0 Leaf Node 18 Total Samples 001 , Distribution [0 0 1 0 0 0] Leaf Node 19 Total Samples 015 , Distribution [ 0 0 15 0 0 0 0] Leaf Node 20 Total Samples 015 , Distribution [ 0 13 2 0 0 Leaf Node 21 Total Samples 121 , Distribution [ 0 62 59 Leaf Node 22 Total Samples 001 , Distribution [0 0 1 0 0 0] Leaf Node 23 Total Samples 009 , Distribution [0 0 9 0 0 0 0] Leaf Node 24 Total Samples 015 , Distribution [ 0 15 0 0 0 0 0] Leaf Node 25 Total Samples 050 , Distribution [ 0 38 6 0 0 0 6] Leaf Node 26 Total Samples 013 , Distribution [ 0 13 0 Leaf Node 27 Total Samples 021 , Distribution [ 0 7 14 Leaf Node 28 Total Samples 005 , Distribution [0 2 0 0 0 0 3] Leaf Node 29 Total Samples 014 , Distribution [ 0 12 2 0 Leaf Node 30 Total Samples 002 , Distribution [0 2 0 0 0 0] Leaf Node 31 Total Samples 012 , Distribution [ 0  $\,$  2  $\,$  0  $\,$  0  $\,$  0  $\,$  0  $\,$  10] Printing a sample tree for visualisation In [8]: tree = Tree(max\_depth=2, min\_samples\_split=2) tree.fit(X train, y train) tree.printTree() printing tree... , ROOT , ROOT, Condition: Elevation <= 2843.0 , NONLEAF, Condition: Elevation> 2843.0 , NONLEAF, Condition: Elevation <= 2524.0 , LEAF, Total Samples 37 , Distribution[ 0 0 3 24 1 0 , LEAF, Total Samples 79 , Distribution[ 0  $\,$  7  $\,$  50  $\,$  10  $\,$  0 , NONLEAF, Condition: Elevation> 3170.0 , LEAF, Total Samples 220 , Distribution[ 0 89 129 , LEAF, Total Samples 132 , Distribution[ 0 91 22 0 0 0 19] **Algorithm** In [9]: from graphviz import Digraph In [10]: | g = Digraph('G') g.edge('Read data', 'Create Decison Tree', label='Training data') g.edge('Create Decison Tree','Start') g.edge('Start', 'Is the data seperable?') g.edge('Is the data seperable?', 'Leaf Node', label='yes') g.edge('Leaf Node', 'Store the \ndata distribution\n at this point') g.edge('Store the  $\mbox{\ensuremath{n}}$ data distribution $\mbox{\ensuremath{n}}$  at this point', 'end') g.edge('Is the data seperable?', 'Non-Leaf Node', label='no') g.edge('Non-Leaf Node', 'find all \n Potenial Splits') g.edge('find all  $\n$  Potenial Splits', 'find the best  $\n$  Column & Split value  $\n$  which would give  $\n$  hig hest information \n gain') g.edge('find the best  $\n$  Column & Split value  $\n$  which would give  $\n$  highest information  $\n$  gain', 'Spl it the data') g.edge('Split the data', '<=')</pre> g.edge('Split the data', '>') g.edge('<=','Is the data seperable?',label='==0 (in case of  $\n$  categorical data)') g.edge('>','Is the data seperable?', label='==1 (in case of  $\n$  categorical data)') Out[10]: Read data Training data Create Decison Tree Start Is the data seperable? yes no Leaf Node Non-Leaf Node Store the find all data distribution **Potenial Splits** at this point ==0 (in case of ==1 (in case of categorical data) categorical data) find the best Column & Split value end which would give highest information gain Split the data <= Verification with in-built function

In [11]: clf = DecisionTreeClassifier(criterion = 'entropy', min samples split = 2, max depth = 5)

columns = ['Depth','Train Accuracy','Train Error', 'Test Accuracy', 'Test Error']

| 27.35 %

2) https://www.youtube.com/watch?v=y6DmpG PtN0&list=PLPOTBrypY74xS3WD0G uzqPjCQfU6IRK-

"{0:.2f}".format(100\*accuracy\_score(y\_train, y\_pred))+' %',
"{0:.2f}".format(100\*(1-accuracy\_score(y\_train, y\_pred)))+' %',
"{0:.2f}".format(100\*accuracy\_score(y\_test, y\_pred2))+' %',

"{0:.2f}".format(100\*(1-accuracy score(y test, y pred2)))+' %'))

| 64.95 %

| 35.05 %

clf.fit(X\_train, y\_train)
y\_pred = clf.predict(X\_train)
y pred2 = clf.predict(X test)

results.append((depth,

df = pd.DataFrame(results, columns=columns)

os.system('dot -Tpng tree.dot -o tree.jpeg')

export graphviz(clf, filled = True,

5 | 72.65 %

1) https://piazza.com/class/kdhx0iapnk06la?cid=347

&index=1&ab channel=SebastianMantey

with open('tree.dot') as f:
 dot graph = f.read()

print(tabulate(df, headers='keys', tablefmt='psql'))

out file = 'tree.dot')

results = []

df.reset index()

Source (dot graph)

Resources

Out[11]:

In [ ]: