|)]: | import pandas as pd import numpy as np from sklearn.preprocessing import MinMaxScaler, StandardScaler from scipy.stats import zscore from sklearn.preprocessing import LabelEncoder from pyitlib import discrete_random_variable as drv from math import log from sklearn.tree import DecisionTreeClassifier from sklearn.model_selection import train_test_split from sklearn.maive_bayes import GaussianNB |
|-----|---|
| | from sklearn.naive_bayes import GaussianNB from sklearn.metrics import accuracy_score import csv import math import random from sklearn.neighbors import KNeighborsClassifier from sklearn.neighbors import confusion_matrix, classification_report from sklearn.cluster import KMeans import pickle from pandas_profiling import ProfileReport Uploading data |
| : | dic = {'train_path': 'D:\\onedrive\\OneDrive - ac.sce.ac.il\\Desktop\\train.csv', 'test_path': 'D:\\onedrive\\OneDrive - ac.sce.ac.il\\Desktop\\test.csv', 'missing_data': 'By all data', 'normalization': 'Yes', 'discritizatio train_file = pd.read_csv(dic['train_path']) test_file = pd.read_csv(dic['test_path']) EDA: profile = ProfileReport(train_file, title="Pandas Profiling Report") profile |
| | Pandas Profiling Report Overview Variables Interactions Correlations Missing values Samp |
| | Overview Warnings 11 Reproduction Dataset statistics Number of variables Number of variables Aumber of observations 42180 Categorical 6 Missing cells Missing cells Missing cells 29 Boolean 4 Missing cells (%) Ouplicate rows Duplicate rows (%) Total size in memory 5.1 Miß Average record size in memory 128.0 B |
| | Age Real number (R ₂₀) Distinct 76 Distinct (%) 0.2% Maximum 500 Our analysis: |
| | 1. Our data has 16 columns : 6 - numeric columns , 6 - categorical columns , 4 - boolean columns . 2. Columns corrolations: contact is highly correlated with month education is highly correlated with job job is highly correlated with education month is highly correlated with contact and 2 other fields day is highly correlated with month month is highly correlated with housing housing is highly correlated with month Delete classification missing data def Delete_Rows(data): data['class'].replace('', np.nan, inplace=True) |
| | <pre>data.dropna(subset=['class'], inplace=True) Delete_Rows(train_file) Delete_Rows(test_file) Complete Missing Data def numeric(j): for i in j: if type(i) == str and i != '': return False</pre> |
| | <pre>return True def D_Array(data): a = [] no = data[data['class'] == 'no'].dropna(axis = 0) yes = data[data['class'] == 'yes'].dropna(axis = 0) for i in data.columns: if numeric(data[i]): a.append([yes[i].mean(), no[i].mean()]) else: a.append([yes[i].value_counts().index[0]], no[i].value_counts().index[0]])</pre> |
| | <pre>return a def Missing_Data(data): a = D_Array(train_file) c = data.columns.tolist() if dic['missing_data'] == 'By all data': for i in data.columns: if numeric(data[i]) == True:</pre> |
| | <pre>else: data[i].fillna(train_file[i].value_counts().index[0], inplace = True) else: for i,rows in data.iterrows(): for b in data.columns: if pd.isnull(data[b][i]) and data['class'][i] == 'yes':</pre> |
| : | <pre>Missing_Data(train_file) Missing_Data(test_file) Split the data for i in train_file.columns: if numeric(train_file[i]): train_file[i] = pd.to_numeric(train_file[i], downcast = 'float') test_file[i] = pd.to_numeric(test_file[i], downcast = 'float')</pre> |
| | <pre>Normalization def Normalization(train_file, test_file): if dic['normalization'] != 'No': for i in train_file.columns: if numeric(train_file[i]): avg = train_file[i].mean() std = train_file[i].std() for j in range(0, len(train_file[i])):</pre> |
| | train_file.at[j, str(i)] = ((train_file[i][j] - avg) / std) for j in range(0, len(test_file[i])): test_file.at[j, str(i)] = ((test_file[i][j] - avg) / std) Normalization(train_file, test_file) Discritization def Equal_width_binning(data, bins, label): for i in data: |
| | <pre>if numeric(data[i]):</pre> |
| | <pre>#Equal-frequency discretization def Equal_frequency_binning_implementation(): def Equal_frequency(data, test, columns , m): a = [] for i in data[columns]: a.append(i) a.sort()</pre> |
| | <pre>length = len(a) n = int(length / m) arri=[] for i in range(0, m): arr = [] for j in range(1 * n, (i + 1) * n): if j >= length: break arr = arr + [a[j]] arri.append(arr) for i in range(0, len(arri)): arri[i] = list(set(arri[i])) for i in range(0, len (arri)): if test[columns][j] in arri[j]: test.at[i,str(columns]) = j for i in range(0, len(data[columns])): for j in range(0, len (data[columns])): for j in range(0, len (arri)):</pre> |
| | <pre>if data[columns][i] in arri[j]:</pre> |
| | <pre>a.append(i) a.sort() w = int((max(a) - min(a)) / m)+1 min1 = min(a) arr = [] for i in range(0, m + 1): arr = arr + [min1 + w * i] arri=[] for i in range(0, m): temp = [] for j in a:</pre> |
| | <pre>for j in a: if j >= arr[i] and j <= arr[i+1]:</pre> |
| | <pre>test.at[i,str(columns)] = j for i in range(0, len(data[columns])): for j in range(0, len (arri)): if data[columns][i] in arri[j]: data.at[i,str(columns)] = j for i in train_file.columns: if numeric(train_file[i]): Equal_width(train_file, test_file, i , dic['number_of_bins']) def Discritization(data): bins = dic['number_of_bins']</pre> |
| | <pre>label = [] for i in list(range(bins)): label.append(i) if dic['discritization'] == 'Equal Width Binning': Equal_width_binning(data, bins, label) elif dic['discritization'] == 'Equal Frequency Binning': Equal_frequency_binning(data, bins) elif dic['discritization'] == 'Equal Width Binning-Implementation': Equal_width_binning_implementation() elif dic['discritization'] == 'Equal Frequency Binning-Implementation': Equal_frequency_binning_implementation() elif dic['discritization'] == 'Discritization based Entropy': for i in data.columns: if numeric(data[i]): val = np.array(data[i]) new_val = val.astype(int) </pre> |
| : | Tis = drv.entropy(new_val) for j in range(0, len(data[i])): if data[i][j] < IG: data.at[j,str(i)] = 0 else: data.at[j,str(i)] = 1 Discritization(train_file) Discritization(test_file) Encoder |
| | <pre>def Encoder(data, dat): for i in data.columns: if not numeric(data[i]): le = LabelEncoder() data[i] = le.fit_transform(data[i]) dat[i] = le.fit_transform(dat[i])</pre> Encoder(train_file, test_file) Saving files |
| | <pre>train_file.to_csv('train_file_clean.csv') test_file.to_csv('test_file_clean.csv') Model: def Naive_bayes(): X_train = train_file.iloc[:, : -1] Y_train = train_file.iloc[:, -1] X_test = test_file.iloc[:, -1] X_test = test_file.iloc[:, -1] Y_test = test_file.iloc[:, -1]</pre> |
| | <pre>gnb = GaussianNB() y_pred = gnb.fit(X_train, Y_train).predict(X_test) print("Number of mislabeled points out of a total %d points : %d"% (X_test.shape[0], (Y_test != y_pred).sum())) accuracy = accuracy_score(Y_test, y_pred)*100 print("Precentence: ", accuracy) print("Confusion matrix: ") print(confusion_matrix(Y_test, y_pred)) print() print() print("Report: ") print("Report: ") print(classification_report(Y_test, y_pred)) pickle.dump(gnb, open('Naive_bayes.sav', 'wb'))</pre> |
| : | <pre>def Decision_Tree(): X_train = train_file.iloc[:, :-1] Y_train = train_file.iloc[:, :-1] X_test = test_file.iloc[:, :-1] Y_test = test_file.iloc[:, :-1] tree = DecisionTreeClassifier(criterion = 'entropy', random_state = 0) tree.fit(X_train, Y_train) pred = tree.predict(X_test) print("The prediction accuracy is: ", tree.score(X_test,Y_test)*100, "%") print("Confusion matrix: ")</pre> |
| | <pre>print(confusion_matrix(Y_test, pred)) print() print("Report: ") print() print(classification_report(Y_test, pred)) pickle.dump(tree, open('Id3.sav', 'wb')) def Naive_bayes_implementation(): py = 0 pn = 0</pre> |
| | <pre>total = len(test_file['class']) correct = 0 y = 0 n = 0 for i in train_file['class']: if i == 0: n = n + 1 else: y = y + 1</pre> |
| | <pre>py = y / len(train_file['class']) pn = n / len(train_file['class']) for i,rows in test_file.iterrows(): arryes = [] arrno = [] for j in test_file.columns:</pre> |
| | <pre>dyes = train_file[train_file[j] == test_file[j][i]] dyes = train_file[train_file['class'] == 1] arryes.append(len(dyes['class']) / y) dno = train_file[train_file[j] == test_file[j][i]] dno = train_file[train_file[j] == 0] arrno.append(len(dno['class']) / n) yes = 0 no = 0</pre> |
| | <pre>for j in arryes: if yes == 0: yes = j else: yes = yes*j yes = yes * py for j in arrno: if no == 0: no = j else: no = no*j</pre> |
| | <pre>no = no * pn if yes > no and 1 == test_file['class'][i]:</pre> |
| | <pre>def Decision_Tree_implementation(): dataset=train_file test_dataset = test_file def entropy(data_set): """ this function calculate data set entropy """ Probability_set = [] for i in data_set: counter = 0 counter = 0</pre> # create a list of Probability |
| | <pre>for j in data_set: if j == i: counter += 1 Probability_set.append(counter / len(data_set)) return sum(list(map(lambda x: (-1) * (x * log(x,2)), Probability_set))) # return entropy def InfoGain(data, split_attribute_name, target_name="class"): #Entropy of the total dataset total_entropy = entropy(data[target_name])</pre> |
| | <pre>##Calculate the entropy of the dataset values, counts= np.unique(data[split_attribute_name], return_counts=True) #weighted entropy W_Entropy = np.sum([(counts[i]/np.sum(counts))*entropy(data.where(data[split_attribute_name]==values[i]).dropna()[target_name]) for i in range(len(values))]) #Calculate the information gain Info_Gain = total_entropy - W_Entropy return Info_Gain</pre> def ID3(data, originaldata, features, target_attribute="class", Parent_Node = None): |
| | <pre>if len(np.unique(data[target_attribute])) <= 1: return np.unique(data[target_attribute])[0] elif len(data)==0: return np.unique(originaldata[target_attribute])[np.argmax(np.unique(originaldata[target_attribute],return_counts=True)[1])] elif len(features) ==0: return Parent_Node else:</pre> |
| | Parent_Node = np.unique(data[target_attribute])[np.argmax(np.unique(data[target_attribute],return_counts=True)[1])] item = [InfoGain(data,feature,target_attribute) for feature in features] index_ofBestFeature = np.argmax(item) Bestfeature = features[index_ofBestFeature] #Create the tree structure. The root gets the name of the feature (best_feature) with the maximum information #gain in the first run tree = {Bestfeature:{}} |
| | <pre>features = [i for i in features if i != Bestfeature] for value in np.unique(data[Bestfeature]): value = value sub_data = data.where(data[Bestfeature] == value).dropna() #Call the ID3 algorithm for each of those sub_datasets with the new parameters, Here the recursion comes in! subtree = ID3(sub_data, dataset, features, target_attribute, Parent_Node) tree[Bestfeature][value] = subtree return(tree)</pre> |
| | <pre>def predict(query, tree, default = 1): for key in list(query.keys()): if key in list(tree.keys()): try:</pre> |
| | <pre>result = tree[key][query[key]] if isinstance(result, dict): return predict(query, result) else: return result def testing(data, tree): queries = data.iloc[:,:-1].to_dict(orient = "records")</pre> |
| | <pre>predicted = pd.DataFrame(columns=["predicted"]) for i in range(len(data)): predicted.loc[i, "predicted"] = predict(queries[i], tree, 1.0) print('The prediction accuracy is: ', (np.sum(predicted["predicted"] == data["class"])/len(data))*100, '%') train_dataset=dataset.iloc[:100].reset_index(drop=True) """</pre> |
| | <pre>Train the tree, Print the tree and predict the accuracy """ tree = ID3(train_dataset, train_dataset.columns[:-1]) testing(test_dataset, tree) def Model(train_file, test_file): if dic['model_type'] == 'Naive bayes': Naive_bayes() elif dic['model_type'] == 'Naive bayes-Implementation': Naive_bayes_implementation()</pre> |
| | <pre>Naive_bayes_implementation() elif dic['model_type'] == 'Decision tree': Decision_Tree() elif dic['model_type'] == 'Decision tree-Implementation': Decision_Tree_implementation()</pre> Model(train_file, test_file) |
| | Mumber of mislabeled points out of a total 3031 points : 1332 precentence: 56.05410755526229 Confusion matrix: [[1279 188] |
| | accuracy |
| | <pre>arr = [] for i in range(2, 10): classifier = KNeighborsClassifier(n_neighbors = i, p = 2, metric = 'euclidean') classifier.fit(X_train, Y_train) y_pred = classifier.predict(X_test) arr.append(accuracy_score(Y_test, y_pred)) index = 0 mx = 0 for i in range(len(arr)): if arr[1] >= mx: mx = arr[i] index = i best_n_neighbors = (index + 2)</pre> |
| | The best n_neighbors is: 3 The prediction accuracy is: 51.303200263939296 |
| | The prediction accuracy is: 51.303200263939296 Confusion matrix: [[1402 65] [1452 112]] Report: precision recall f1-score support 0 0.49 0.96 0.65 1467 1 0.63 0.07 0.13 1564 accuracy macro avg 0.56 0.51 0.39 3031 |
| | |
| | <pre>for i in range(2, 10): classifier = KMeans(n_clusters = i) classifier.fit(X_train, Y_train) y_pred = classifier.predict(X_test) arr.append(accuracy_score(Y_test, y_pred)) index = 0 mx = 0 for i in range(len(arr)): if arr[i] >= mx:</pre> |
| | |
| | <pre>print("Confusion matrix: ") print(confusion_matrix(Y_test, y_pred)) print() print("Report: ") print() print(classification_report(Y_test, y_pred)) prickle.dump(classifier, open('K_means.sav', 'wb')) K_means() The best n_clusters is: 2 The prediction accuracy is: 52.29297261629825 Confusion matrix:</pre> |
| | Confusion matrix: [[615 852] [593 971]] Report: precision recall f1-score support 0 0.51 0.42 0.46 1467 1 0.53 0.62 0.57 1564 accuracy 0.52 3031 |
| | macro avg 0.52 0.52 0.52 3031 weighted avg 0.52 0.52 0.52 3031 |