TASK1 classes = np.array(["business", "entertainment", "politics", "sport", "tech"]) class_article_count = np.zeros(classes.shape[0]) for i in range(len(classes)): direc = "BBC/" + classes[i] class_article_count[i] = len(os.listdir(direc)) plt1.bar(classes,class article count) plt1.set title('BBC Distribution') plt1.set xlabel('Class Name') plt1.set_ylabel('Number of Instances') TASK2 df = pd.read csv('drug200.csv') classes = np.array(['drugA', 'drugB', 'drugC', 'drugX', 'drugY']) class_instance_count = np.zeros_like(classes) for i in range(classes.shape[0]): class_instance_count[i] = (df.Drug.values == classes[i]).sum() plt2.bar(classes,class_instance_count, color='orange') plt2.set title('Drug Distribution') plt2.set xlabel('Class Name') plt2.set_ylabel('Number of Instances') Out[2]: Text(0, 0.5, 'Number of Instances') BBC Distribution Drug Distribution 91 500 400 Number of Instances 300 200 100 23 business entertainment politics sport drugB drugC drugX drugY tech drugA Class Name Class Name The distributions in Task 1 (BBC Distribution) and Task 2 (Drug Distribution) are very different. The classes in BBC Distribution are evenly distributed. All five classes (business, entertainment, politics, sport, and tech) have a similar number of sample articles (number of instances). While on the other hand, the classes in the Drug Distribution are unbalanced, creating a skewed distribution. Accuracy is well suited for **Task 1** since we have a **well-balanced** dataset. It is easy to see in the BBC Distribution that all article categories have between 400-500 article instances each. Also, it is appropriate to use the accuracy metric for Task 1 as all classes are equally important and equally represented in this case. We are not interested in one type of news category more than another. As for **Task 2**, precision and recall are better-suited metrics to use to evaluate the model. Accuracy is not appropriate in this case since we clearly have a very **unbalanced** distribution of instances. In addition to that, although precision could be used in this task, in a **medical application**, it is preferable to pick a model with the **highest recall**. Since we are dealing with medical data, one type of drug class could be more important than another. Analysis of the results of all the models In [39]: accuracy, f1 macro, f1 weighted = np.zeros(4), np.zeros(4), np.zeros(4) # Q3 - Load the corpus using load files, this assigns a class to each article files info = sklearn.datasets.load files(container path='BBC/', encoding='latin1') corpus = files info.data # articles from all classes vectorizer = CountVectorizer() X = vectorizer.fit transform(corpus) # Learns the vocabulary dictionary and return document-term matrix. [article#, word#, word count] y = files info.target X_train, X_test, y_train, y_test = train_test_split(X.toarray(), y, test_size=.2, random_state=None) clf = MultinomialNB().fit(X_train, y_train) # Create a model using training data y_test_pred = clf.predict(X_test) # Use the model created to make predictions on the test data accuracy[0] = accuracy_score(y_test, y_test_pred) f1 macro[0] = f1 score(y test, y test pred, average='macro') f1 weighted[0] = f1 score(y test, y test pred, average='weighted') clf = MultinomialNB().fit(X_train, y_train) # Create a model using training data y test pred = clf.predict(X test) # Use the model created to make predictions on the test data accuracy[1] = accuracy_score(y_test, y_test_pred) f1_macro[1] = f1_score(y_test, y_test_pred, average='macro') f1_weighted[1] = f1_score(y_test, y_test_pred, average='weighted') clf = MultinomialNB(alpha=.0001).fit(X_train, y_train) # Create a model using training data y_test_pred = clf.predict(X_test) # Use the model created to make predictions on the test data accuracy[2] = accuracy_score(y_test, y_test_pred) f1_macro[2] = f1_score(y_test, y_test_pred, average='macro') f1_weighted[2] = f1_score(y_test, y_test_pred, average='weighted') clf = MultinomialNB(alpha=.9).fit(X_train, y_train) # Create a model using training data y_test_pred = clf.predict(X_test) # Use the model created to make predictions on the test data accuracy[3] = accuracy_score(y_test, y_test_pred) f1_macro[3] = f1_score(y_test, y_test_pred, average='macro') f1_weighted[3] = f1_score(y_test, y_test_pred, average='weighted') plt.plot(np.arange(4,dtype=np.int32)+1, accuracy*100, label='accuracy') plt.plot(np.arange(4,dtype=np.int32)+1, f1_macro*100, label='macro_f1') plt.plot(np.arange(4,dtype=np.int32)+1, f1_weighted*100, label='weighted_f1') plt.plot(1, ((accuracy[0]+f1_macro[0]+f1_weighted[0])/3)*100,'bo', label='smoothing=1') plt.plot(2, ((accuracy[1]+f1_macro[1]+f1_weighted[1])/3)*100,'bo') plt.plot(3, ((accuracy[2]+f1_macro[2]+f1_weighted[2])/3)*100,'ro', label='smoothing=0.0001') plt.plot(4, ((accuracy[3]+f1_macro[3]+f1_weighted[3])/3)*100,'go', label='smoothing=0.9') plt.legend() plt.xlabel('Iteration') plt.ylabel('Performance (%)') plt.title('Multinomial Naive Bayes Classifier') Out[39]: Text(0.5, 1.0, 'Multinomial Naive Bayes Classifier') Multinomial Naive Bayes Classifier 97.6 accuracy macro_fl 97.5 weighted_f1 97.4 smoothing=1 smoothing=0.0001 97.3 smoothing=0.9 97.2 ية 97.1 97.0 96.9 1.0 1.5 2.0 2.5 3.0 3.5 Iteration **Iteration 1 and 2** are the **same** since at both times we used the **default values**. The default values in sklearn.naibe_bayes.MultinomialNB are alpha=1.0 (smoothing), fit-prior=True, and class_prior=None. However, in **iteration 3 and 4**, we did **change the smoothing values** (alpha=0.0001 and alpha=0.9), so this time we do see slightly different performance results.

In [5]:

import os

MP 1

In [2]:

import pandas as pd import numpy as np

import sklearn.datasets

import matplotlib.pyplot as plt

from sklearn.model selection import train test split

from sklearn.naive bayes import GaussianNB

from sklearn.linear model import Perceptron

from sklearn.naive bayes import MultinomialNB from sklearn.metrics import confusion matrix

from sklearn.metrics import accuracy score from sklearn.metrics import precision_score from sklearn.metrics import recall score

from sklearn.metrics import f1 score

Inital dataset analysis

from sklearn.tree import DecisionTreeClassifier from sklearn.model_selection import GridSearchCV

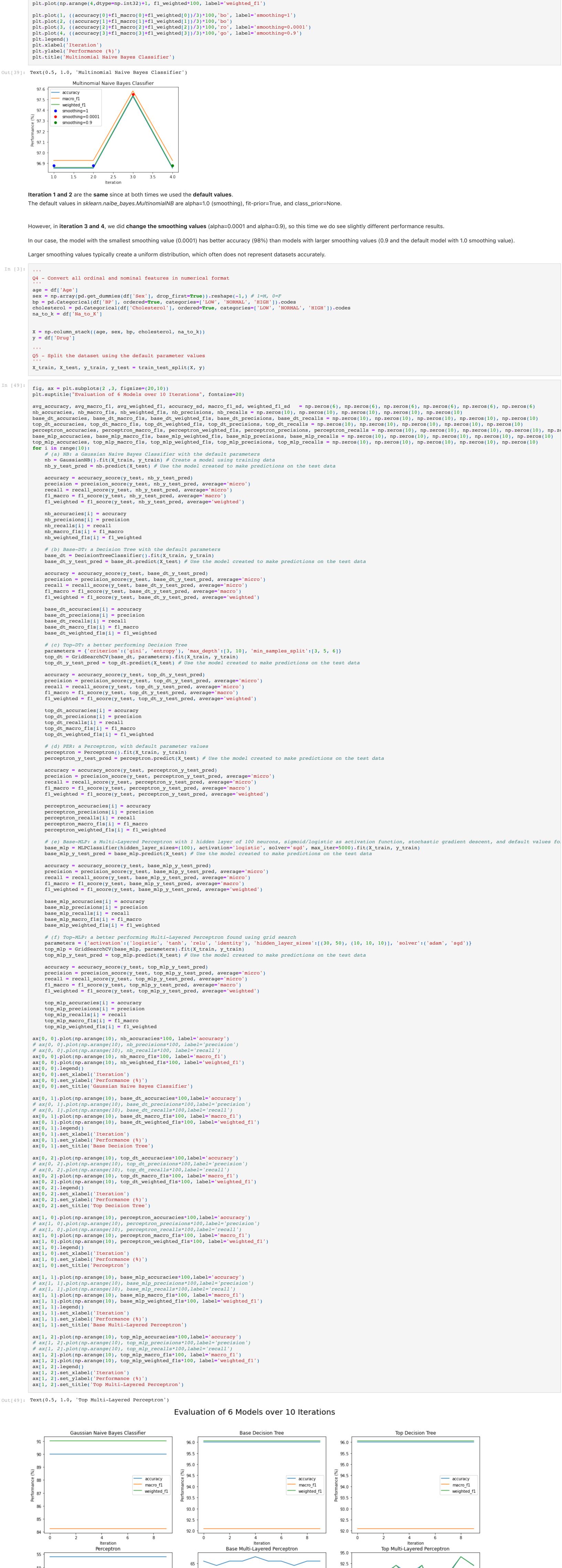
from sklearn.neural network import MLPClassifier

from sklearn.metrics import classification_report

from sklearn.feature_extraction.text import CountVectorizer

Plots of the **distributions** of the instances in each class for Task1 and Task2.

fig, (plt1, plt2) = plt.subplots(1 ,2, figsize=(20,5))



50 60 90.0 45 § 55 87.5 ance 50 accuracy accuracy 85.0 macro f1 macro fl weighted fl weighted fl 82.5 45 35 80.0 30 77.5 35 accuracy macro_fl 25 75.0 30 weighted_f1 Iteration Iteration Iteration Analyzing the plots above, we see that the NB, Base DT, Top DT, and Perceptron have constant performance over all iterations. This is because **no element of randomness** goes into the creation of these models. If the same training and testing data is used every time, the same priors and conditionals will be generated/calculated during the model creation, and in turn used to calculate the score of classes or the entropy in DTs. As for the Perceptron, the initial weights are always set to 1 according to the documentation, so the final weights will be also the same. This is the reason why we do not see any changes in perfromance. For the **Base MLP** and **Top MLP**, we see **changes in performance** over the iterations. This is happening since these models use random weights and bias initialization. So, each iteration will have different final weights. The change in performance is less drastic in the Base MLP because we are not changing the hyper-parameters. While in Top MLP, we are not only using random parameters initialization, we are also experimenting with hyper-parameters.