of Alzheimer's Diseases in MRI Scans packages that need to install to run this code • pip install **tensrflow** || in case of GPU use pip install **tensrflow-gpu** pip install **imblearn** pip install **tensorflow-addons** pip install **matplotlib** pip install **seaborn** pip install **keras** • pip install **scikit-learn** **Dataset Link** File modified to run on colab Follow the below instructions • Instructions to add dataset in colab from kaggle Link • download dataset in your current directory or another and carefully add path in the **WORKING_DIRECTORY** variable In []: !pip install tensorflow !pip install keras !pip install imblearn !pip install matplotlib !pip install seaborn !pip install scikit-learn !pip install tensorflow-addons **Importing Libraries** In []: import numpy as np import random # Plotting import seaborn as sns import matplotlib.pyplot as plt # DataGenerator to read images and rescale images from tensorflow.keras.preprocessing.image import ImageDataGenerator import tensorflow as tf import tensorflow addons as tfa # count each class samples from collections import Counter # callbacks from tensorflow.keras.callbacks import ReduceLROnPlateau # evaluate precison recall and f1-score of each class of model from sklearn.metrics import classification report # Show performance of a classification model from sklearn.metrics import confusion matrix # Different layers from keras.models import Sequential from tensorflow.keras.layers import Input from tensorflow.keras.layers import Flatten from tensorflow.keras.layers import AveragePooling2D from tensorflow.keras.layers import Convolution2D from tensorflow.keras.layers import Dense from tensorflow.keras.layers import Dropout from tensorflow.keras.layers import ReLU from tensorflow.keras.layers import Softmax # split dataset to train, validation and test set from sklearn.model selection import train test split # callbacks from keras import callbacks # SMOTETomek from imblance library from imblearn.combine import SMOTETomek # Optimizer from tensorflow.keras.optimizers import SGD from sklearn.metrics import roc curve, auc from itertools import cycle Define directory of dataset & Classes names In []: ## Set Path Here before running the code WORKING DIRECTORY = ".\\dataset\\" ## Name of classes CLASSES = ['Mild-Demented', 'Moderate-Demented', 'Non-Demented', 'VeryMild-Demented'] Load Images, Rescale Images, and seperate from data generator & Label One Hot encoding In []: X, y = [], []## Images rescaling datagen = ImageDataGenerator(rescale=1.0/255.0) # Load images by resizing and shuffling randomly train_dataset = datagen.flow_from_directory(WORKING_DIRECTORY, target_size=(176, 208),batch_size=6400, shuffle= ### Seperate Dataset from Data Genrator X, y = train_dataset.next() In []: $samples_before = len(X)$ print("Images shape :\t", X.shape) print("Labels shape :\t", y.shape) In []: # Number of samples in classes print("Number of samples in each class:\t", sorted(Counter(np.argmax(y, axis=1)).items())) # class labels as per indices print("Classes Names according to index:\t", train_dataset.class_indices) Show some random samples from the origional dataset In []: | # show some samples from the dataset randomly fig = plt.figure(figsize=(10,8)) rows = 4columns = 4for i in range(rows * columns): fig.add_subplot(rows, columns, i+1) num = random.randint(0, len(X)-1) plt.imshow(X[num]) plt.axis('off') plt.title(CLASSES[(np.argmax(y[num]))], fontsize=8) plt.axis('off') plt.show() Apply SMOTETomek Algorithm to balance the dataset In []: # reshaping the images to 1D X = X.reshape(-1, 176 * 208 * 3)# Oversampling method to remove imbalance class problem X, y = SMOTETomek().fit resample(X, y)reshape images to images size of 208, 176, 3 X = X.reshape(-1, 176, 208, 3)samples after = len(X)print("Number of samples after SMOTETomek :\t", sorted(Counter(np.argmax(y, axis=1)).items())) Show some random samples from the Generated dataset In []: fig = plt.figure(figsize=(10,8)) rows = 4columns = 4for i in range(rows * columns): fig.add subplot(rows, columns, i+1) num = random.randint(samples_before, samples_after - 1) plt.imshow(X[num]) plt.axis('off') plt.title(CLASSES[(np.argmax(y[num]))], fontsize=8) plt.axis('off') plt.show() Splitting dataset for Training, Validation & Testing In []: 20% split to validation and 80% split to train set X_train, x_val, y_train, y_val = train_test_split(X,y, test_size = 0.2) # 20% split to test from 80% of train and 60% remains in train set X_train, x_test, y_train, y_test = train_test_split(X_train, y_train, test_size = 0.2) # Number of samples after train test split print("Number of samples after splitting into Training, validation & test set\n") print("Validation\t", sorted(Counter(np.argmax(y val, axis=1)).items())) print("Test \t", sorted(Counter(np.argmax(y_test, axis=1)).items())) In []: # to free memeory we don't need this one as we split our data del X, y **Model Architecture** from keras.initializers import GlorotUniformV2 init = GlorotUniformV2 model = Sequential() model.add(Input(shape=(176, 208, 3))) model.add(Convolution2D(16, 5, kernel_initializer=init)) model.add(ReLU()) model.add(AveragePooling2D(pool_size=(2,2))) model.add(Convolution2D(32, 5, kernel_initializer=init)) model.add(ReLU()) model.add(AveragePooling2D(pool_size=(2,2))) model.add(Convolution2D(64, 5, kernel_initializer=init)) model.add(ReLU()) model.add(AveragePooling2D(pool_size=(2,2))) model.add(Convolution2D(128, 5, kernel_initializer=init)) model.add(ReLU()) model.add(AveragePooling2D(pool_size=(2,2))) model.add(Dropout(0.01)) model.add(Flatten()) model.add(Dense(256, kernel_initializer=init)) model.add(ReLU()) model.add(Dropout(0.03)) model.add(Dense(4, kernel_initializer=init)) model.add(Softmax()) model.summary() **Compiling the Model** In []: ### Model Compilation model.compile(optimizer=SGD(learning rate=0.01), loss = tf.keras.losses.CategoricalCrossentropy(name='loss'), tf.keras.metrics.CategoricalAccuracy(name='acc'), tf.keras.metrics.AUC(name='auc'), tfa.metrics.F1Score(num classes=4), tf.metrics.Precision(name="precision"), tf.metrics.Recall(name="recall")]) **Defining CALLBACKS to reduce Learning Rate** In []: # callbacks used in model to perform well rop_callback = ReduceLROnPlateau(monitor="val loss", patience=2) CALLBACKS = [rop_callback] Training of the Model In []: declare to run on small gpu create batch sizes of images valAug = ImageDataGenerator() # defining batch size batch size = 8 history = model.fit(valAug.flow(X train, y train, batch size=batch size, shuffle = True), steps per epoch=len(X train) // batch size, validation data=valAug.flow(x val, y val, batch size=batch size, shuffle = True), validation steps=len(x test) // batch size, epochs= 40, batch size=batch size, callbacks = CALLBACKS **Evaluation of Model with the Test data** ### Evaluate Model test_scores = model.evaluate(x_test, y_test, batch_size = 32) print("\n\nTesting Loss : \t\t {0:0.6f}".format(test_scores[0])) print("Testing Accuracy : \t {0:0.6f} %".format(test_scores[1] * 100)) print("Testing AC : \t\t {0:0.6f} %".format(test_scores[2] * 100)) print("Testing F1-Score : \t {0:0.6f} %".format(((test_scores[3][0] + test_scores[3][1] + test_scores[3][2] + t print("Testing Precision : \t {0:0.6f} %".format(test_scores[4] * 100)) print("Testing Recall : \t {0:0.6f} %".format(test_scores[5] * 100)) **Model Training graphs** Accuracy Loss AUC Precision Recall • F1-Score In []: plt.plot(history.history['acc'], 'b') plt.plot(history.history['val_acc'], 'g') plt.title("Model Accuracy") plt.xlabel("Epochs") plt.ylabel("Accuracy") plt.legend(["train", "val"]) plt.show() In []: plt.plot(history.history['loss'], 'b') plt.plot(history.history['val_loss'], 'g') plt.title("Model Loss") plt.xlabel("Epochs") plt.ylabel("Loss") plt.legend(["train", "val"]) plt.show() In []: plt.plot(history.history['auc'], 'b') plt.plot(history.history['val auc'], 'g') plt.title("Model AUC") plt.xlabel("Epochs") plt.ylabel("AUC") plt.legend(["train", "val"]) plt.show() In []: plt.plot(history.history['precision'], 'b') plt.plot(history.history['val precision'], 'g') plt.title("Model Precision") plt.xlabel("Epochs") plt.ylabel("Precision") plt.legend(["train", "val"]) plt.show() In []: plt.plot(history.history['recall'], 'b') plt.plot(history.history['val_recall'], 'g') plt.title("Model Recall") plt.xlabel("Epochs") plt.ylabel("Recall") plt.legend(["train", "val"]) plt.show() In []: plt.plot(history.history['f1_score']) plt.plot(history.history['val_f1_score']) plt.title("Model F1-Score") plt.xlabel("Epochs") plt.ylabel("F1-Score") plt.show() Test set Evaluation Classification Report Confusion Matrix ROC Curve Extension ROC Multiclass In []: pred_labels = model.predict(x_test, batch_size=32) def roundoff(arr): arr[np.argwhere(arr != arr.max())] = 0 arr[np.argwhere(arr == arr.max())] = 1 return arr for labels in pred_labels: labels = roundoff(labels) print(classification_report(y_test, pred_labels, target_names=CLASSES)) In []: pred ls = np.argmax(pred labels, axis=1) test_ls = np.argmax(y_test, axis=1) conf_arr = confusion_matrix(test_ls, pred_ls) plt.figure(figsize=(10, 8), dpi=80, facecolor='w', edgecolor='k') ax = sns.heatmap(conf_arr, cmap='Greens', annot=True, fmt='d', xticklabels= CLASSES, yticklabels=CLASSES) plt.title('Confusion Matrix of Model', fontweight='bold', fontsize=14.0) plt.xlabel('Predictions', fontweight='bold', fontsize=13) plt.ylabel('Ground Truth', fontweight='bold', fontsize=13) plt.tight layout() plt.show(ax) In []: fpr = dict() tpr = dict()roc auc = dict() for i in range(4): fpr[i], tpr[i], _ = roc_curve(y_test[:, i], pred_labels[:, i]) roc_auc[i] = auc(fpr[i], tpr[i]) # Compute micro-average ROC curve and ROC area fpr["micro"], tpr["micro"], = roc_curve(y test.ravel(), pred_labels.ravel()) roc_auc["micro"] = auc(fpr["micro"], tpr["micro"]) plt.figure() lw = 2plt.plot(fpr[2], tpr[2], color="darkorange", lw=lw, label="ROC curve (area = %0.4f)" % roc_auc[2]) plt.plot([0, 1], [0, 1], color="navy", lw=lw, linestyle="--") plt.xlim([0.0, 1.0]) plt.ylim([0.0, 1.05]) plt.xlabel("False Positive Rate") plt.ylabel("True Positive Rate") plt.title("Receiver operating characteristic ") plt.legend(loc="lower right") plt.show() In []: n classes = 4 # First aggregate all false positive rates all_fpr = np.unique(np.concatenate([fpr[i] for i in range(n_classes)])) # Then interpolate all ROC curves at this points mean_tpr = np.zeros_like(all_fpr) for i in range(n classes): mean tpr += np.interp(all_fpr, fpr[i], tpr[i]) # Finally average it and compute AUC mean tpr /= n classes fpr["macro"] = all_fpr tpr["macro"] = mean_tpr roc_auc["macro"] = auc(fpr["macro"], tpr["macro"]) # Plot all ROC curves plt.figure() plt.plot(fpr["micro"], tpr["micro"], label="micro-average ROC curve (area = {0:0.4f})".format(roc_auc["micro"]), color="deeppink", linestyle=":", linewidth=4, plt.plot(fpr["macro"], tpr["macro"], label="macro-average ROC curve (area = {0:0.4f})".format(roc auc["macro"]), color="navy", linestyle=":", linewidth=4, for i in range(n classes): plt.plot(fpr[i], tpr[i], lw=lw, label="ROC curve of class {0} (area = {1:0.4f})".format(i, roc auc[i]), plt.plot([0, 1], [0, 1], "k--", lw=lw) plt.xlim([0.0, 1.0]) plt.ylim([0.0, 1.05]) plt.xlabel("False Positive Rate") plt.ylabel("True Positive Rate") plt.title("Some extension of Receiver operating characteristic to multiclass") plt.legend(loc="lower right") plt.show()

Saving Model for Future Use

model.save(".\\model.h5")

To save the model in the current directory

In []:

ADD-Net: An Effective Deep Learning Model for Early Detection