universal_audio_preprocessing

January 10, 2022

1 AUGMENTED RAVDESS Audio preprocessing

```
[1]: pip install ipynb
    Defaulting to user installation because normal site-packages is not writeable
    Looking in indexes: https://pypi.org/simple, https://pypi.ngc.nvidia.com
    Requirement already satisfied: ipynb in
    /home/19126778/.local/lib/python3.9/site-packages (0.5.1)
    Note: you may need to restart the kernel to use updated packages.
[6]: import librosa as lb
     import numpy as np
     import matplotlib.pyplot as plt
     import os, glob
     import json
     import time
     import pandas as pd
     from multiprocessing import Pool
     from joblib import Parallel, delayed
     import sys
     sys.path.insert(0, './Augmenter')
     from ipynb.fs.full.augmentation import Augmenter
     ModuleNotFoundError
                                                Traceback (most recent call last)
     /tmp/ipykernel_42280/1096819287.py in <module>
           12 import sys
           13 sys.path.insert(0, './Augmenter')
     ---> 14 from ipynb.fs.full.augmentation import Augmenter
     ModuleNotFoundError: No module named 'ipynb.fs.full.augmentation'
[]: TRAIN RAVDESS AUDIO FILES PATH = "../../data/Train/OriginalData/RAVDESS/*.wav"
```

TEST_RAVDESS_AUDIO_FILES_PATH = "../../data/Test/OriginalData/RAVDESS/*.wav"

```
VALIDATE_RAVDESS_AUDIO_FILES_PATH = "../../data/Validate/OriginalData/RAVDESS/*.
-wav"
TRAIN_CREMA_D_AUDIO_FILES_PATH = "../../data/Train/OriginalData/CREMA-D/
→AudioWAV/*.wav"
TEST_CREMA_D_AUDIO_FILES_PATH = ".../.../data/Test/OriginalData/CREMA-D/AudioWAV/
VALIDATE_CREMA_D_AUDIO_FILES_PATH = "../../data/Validate/OriginalData/CREMA-D/
→AudioWAV/*.wav"
PROCESSED_DATA = {
    'features': [],
    'emotions': []
}
focused_emotion_labels = ['neutral', 'happy', 'sad', 'angry', 'fearful', |
# CREMA-D Female samples
CREMA_D_female_samples =__
\rightarrow [1002,1003,1004,1006,1007,1008,1009,1010,1012,1013,1018,1020,1021,1024,1025,1028,1029,1030,
→1052,1053,1054,1055,1056,1058,1060,1061,1063,1072,1073,1074,1075,1076,1078,1079,1082,1084,1
# CREMA-D
CREMA_D_emotion_labels = {
  'NEU': 'neutral',
  'HAP': 'happy',
  'SAD':'sad',
  'ANG': 'angry',
 'FEA': 'fearful',
  'DIS':'disgust'
}
# RAVDESS
RAVDESS emotion labels = {
  '01':'"neutral"',
  '02':'calm',
  '03': 'happy',
  '04':'sad',
  '05': 'angry',
  '06':'fearful',
  '07':'disgust',
  '08':'surprised'
```

```
[]: def load_data_in_pos_neg(path, dataset_name:str):
         data = []
         # https://www.paulekman.com/universal-emotions/what-is-surprise/
         positive = ["neutral", "happy", "calm", "suprised"]
         negative = ["sad", "angry", "fearful", "disgust", "suprised"]
         for i, file in enumerate(glob.glob(path)):
             file_path = os.path.basename(file)
             emotion = ''
             if dataset name == "ravdess":
                  emotion = RAVDESS_emotion_labels[file_path.split("-")[2]]
                  \#emotion = RAVDESS\_emotion\_labels[file\_path.split("-")[3]] \#turn on_{\square}
      \rightarrow for trimmed data
             else:
                  emotion = CREMA_D_emotion_labels[file_path.split("_")[2]]
             if emotion not in focused_emotion_labels:
                  continue
             if emotion in positve:
                  label = "positive"
             else:
                  label = "negative"
             data.append([file, label])
         end_time = time.perf_counter()
         return pd.DataFrame(data, columns=["file", "emotion"])
[]: def load_files_in_df(path, dataset_name:str):
         data = []
         start_time = time.perf_counter()
         for i, file in enumerate(glob.glob(path)):
             file_path = os.path.basename(file)
             emotion = ''
             if dataset_name == "ravdess":
                  emotion = RAVDESS_emotion_labels[file_path.split("-")[2]]
                  \#emotion = RAVDESS\_emotion\_labels[file\_path.split("-")[3]] \#turn on_{\square}
      \hookrightarrow for trimmed data
             else:
                  emotion = CREMA_D_emotion_labels[file_path.split("_")[2]]
```

```
[]: def load_sex_splitted_files(path:str, dataset_name:str):
             Splits the male and female data into seperate datasets
         female_data = []
         male_data = []
         for i, file in enumerate(glob.glob(path)):
             file_path = os.path.basename(file)
             if dataset_name == "ravdess":
                 # Splits RAVDESS sex samples
                 parts = file_path.replace('.','-').split("-")
                 #emotion = RAVDESS_emotion_labels[parts[2]]
                 emotion = RAVDESS_emotion_labels[parts[3]] #turn on for trimmed data
                 if emotion not in focused_emotion_labels:
                     continue
                 if int(parts[7])%2 == 0: #turn on for trimmed data
                 #if int(parts[6])\%2 == 0:
                     # Female sample
                     female_data.append([file, emotion])
                 else:
                     # Male sample
                     male_data.append([file, emotion])
             else:
                 # Splits CREMA-D sex samples
                 parts = file_path.replace('.','_').split("_")
                 emotion = CREMA_D_emotion_labels[parts[2]]
                 if emotion not in focused_emotion_labels:
                     continue
                 if int(parts[0].split('-')[1]) in CREMA D_female_samples: # turn on_
      \hookrightarrow for trimmed data
                 #if int(parts[0]) in CREMA_D_female_samples:
```

```
# Female sample
                    female_data.append([file, emotion])
                else:
                    # Male sample
                    male_data.append([file, emotion])
        female_df = pd.DataFrame(female_data, columns=["file", "emotion"])
        male df = pd.DataFrame(male data, columns=["file", "emotion"])
        return female df, male df
[]: def augment_audio(row, is_augmented: bool = False):
        audio, sr = lb.load(row["file"], sr=22050)
        extracted_features = []
        augmented audios = [audio]
        if is_augmented:
            # Change pitch down
            augmented_audios.append(Augmenter.change_pitch(audio=audio, sr=sr))
            # Change pitch up
            augmented audios.append(Augmenter.change_pitch(audio=audio, sr=sr,_
     →pitch_type="up"))
            # Change speed slow
            augmented_audios.append(Augmenter.change_speed(audio=audio))
            # Change speed fast
            augmented_audios.append(Augmenter.change_speed(audio=audio,__
     # Change speed & pitch down
            augmented_audios.append(Augmenter.change_speed_and_pitch(audio=audio,_
     ⇒sr=sr))
            # Change speed & pitch up
            augmented_audios.append(Augmenter.change_speed_and_pitch(audio=audio,_
     # Add distribution noise
            augmented_audios.append(Augmenter.add_distribution_noise(audio=audio))
        for a in augmented_audios:
```

extracted_features.append(extract_feature(a, sr, row["emotion"]))

```
return extracted_features
```

```
[]: def extract feature(audio, sr, emotion, mfcc=True, chroma=True, mel=True):
         result = np.array([])
         if mfcc:
             # Gets the mean of the MFCC
             # Change to 21 bins instead of 40
             mfccs = np.mean(lb.feature.mfcc(y=audio, sr=sr, n_mfcc=40).T, axis=0)
             result = np.hstack((result, mfccs))
         if chroma:
             # Gets the mean of the chromagram
             stft = np.abs(lb.stft(audio))
             chroma = np.mean(lb.feature.chroma_stft(S=stft, sr=sr).T, axis=0)
             result = np.hstack((result, chroma))
         if mel:
             # Gets the mean of the Mel-frequency spectrogram
             mel = np.mean(lb.feature.melspectrogram(audio, sr=sr).T, axis=0)
             result = np.hstack((result, mel))
         return (result.tolist(), emotion)
```

```
[]: def create_json(df, dataset_type:str, dataset_name:str, is_augmented:bool, sex:

→str = "", augmentation_type:str=""):
         start_time = time.perf_counter()
         n_{proc} = 35
         data = Parallel(n_jobs=n_proc)(delayed(augment_audio)(row, is_augmented)_u
      →for _, row in df.iterrows())
         formatted_data = {
             "features": [],
             "emotions": []
         }
         for d in data:
             for audio in d:
                 formatted_data["features"].append(audio[0])
                 formatted_data["emotions"].append(audio[1])
         is_sex = "" if not sex else f"{sex}_"
         is_agumented = "augmented_" if is_augmented else ""
         json_path = f'../data/{dataset_type}/{dataset_name}/
      →{dataset_type}_{is_sex}{is_agumented}{augmentation_type}_{dataset_name}_preprocessed_data.
      ⇔json'
         print(json_path)
```

```
with open(json_path, 'w') as outfile:
    json.dump(formatted_data, outfile, indent=4)

end_time = time.perf_counter()
    print(f"{dataset_type} set | DONE PROCESSING DATA!!! | Duration: {end_time_u}

-- start_time:04f}")

print(f'Size: {len(formatted_data["features"])}\n')
```

```
[]: def store_data(datasets, dataset_name:str, is_augmented:bool = False, split_sex:
      →bool = False, augmentation_type:str="", split_pos_neg=False):
         for key in datasets:
             if split_sex:
                 print("Sex splitted")
                 female_df, male_df = load_sex_splitted_files(datasets[key],__
      →dataset_name)
                  # Create female JSON file
                  #create_json(df=female_df, dataset_type=key,__
      \rightarrow dataset_name=dataset_name, is_augmented=is_augmented,__
      → sex='female', augmentation_type=augmentation_type)
                  # Create male JSON file
                  #create_json(df=male_df, dataset_type=key,_
      \rightarrow dataset_name=dataset_name, is_augmented=is_augmented,_u
      → sex='male', augmentation_type=augmentation_type)
             if split_pos_neg:
                 print("Emotions classified")
                 pos_df, neg_df = load_data in_pos_neg(datasets[key], dataset_name)
                 # maak pos neg df met load
             else:
                 print("Default")
                 df = load_files_in_df(datasets[key], dataset_name)
                  #create_json(df=df, dataset_type=key,_
      \rightarrow dataset_name=dataset_name, is_augmented=is_augmented,
      → augmentation_type=augmentation_type)
```

1.1 Combined preprocessing

```
[]: def store_data_combined(datasets, is_augmented:bool = False, split_sex:bool = 

→False, augmentation_type:str=""):
    datasets = datasets
```

```
dataset_name='ravdess'
  df_train_r = load_files_in_df(datasets['train-r'], dataset_name)
  df_test_r = load_files_in_df(datasets['test-r'], dataset_name)
  df_valid_r = load_files_in_df(datasets['validate-r'], dataset_name)
  print(len(df_train_r))
  print(len(df_test_r))
  print(len(df_valid_r))
  print()
  dataset name='crema-d'
  df train c = load files in df(datasets['train-c'], dataset name)
  df_test_c = load_files_in_df(datasets['test-c'], dataset_name)
  df_valid_c = load_files_in_df(datasets['validate-c'], dataset_name)
  print(len(df_train_c))
  print(len(df_test_c))
  print(len(df_valid_c))
  print()
  df_train = pd.concat([df_train_r,df_train_c])
  df_test = pd.concat([df_test_r,df_test_c])
  df_valid = pd.concat([df_valid_r,df_valid_c])
  print(len(df train))
  print(len(df_test))
  print(len(df valid))
  print()
   create_json(df_train, 'train', 'combined', __
→is_augmented,split_sex,augmentation_type)
   create_json(df_test, 'test', 'combined', __
→is_augmented,split_sex,augmentation_type)
   create_json(df_test, 'validate','combined',_
→is_augmented,split_sex,augmentation_type)
```

1.1.1 RAVDESS preprocessing

1.1.2 Original data - Ravdess

1.1.3 Original data - Crema-d

1.1.4 Trimmed data - Raydess

(turn on commented line in load_files_in_df())

```
[]: datasets = {
    'train': "../../data/Train/Trimmed/RAVDESS/*.wav",
    'test': "../../data/Test/Trimmed/RAVDESS/*.wav",
    'validate': "../../data/Validate/Trimmed/RAVDESS/*.wav",
```

1.1.5 Trimmed data - Crema-d

(turn on commented line in load_files_in_df())

1.1.6 Augmented - pitch up - RAVDESS

1.1.7 Augmented - pitch up - CREMA-D

```
[]: datasets = {
    'train': TRAIN_CREMA_D_AUDIO_FILES_PATH,
    'test': TEST_CREMA_D_AUDIO_FILES_PATH,
    'validate': VALIDATE_CREMA_D_AUDIO_FILES_PATH,
}
store_data(datasets=datasets,
```

```
dataset_name='crema-d',
  is_augmented=True,
  augmentation_type='PitchUp')
```

1.1.8 Augmented - pitch down - RAVDESS

```
[]: datasets = {
    'train': TRAIN_RAVDESS_AUDIO_FILES_PATH,
    'test': TEST_RAVDESS_AUDIO_FILES_PATH,
    'validate': VALIDATE_RAVDESS_AUDIO_FILES_PATH,
}

store_data(datasets=datasets,
    dataset_name='ravdess',
    is_augmented=True,
    augmentation_type='PitchDown')
```

1.1.9 Augmented - pitch down - CREMA-D

1.1.10 Augmented - lower speed - RAVDESS

1.1.11 Augmented - lower speed - CREMA-D

1.1.12 Augmented - higher speed - RAVDESS

1.1.13 Augmented - higher speed - CREMA-D

1.1.14 Augmented - distribution noise - RAVDESS

1.1.15 Augmented - distribution noise - CREMA-D

1.1.16 Augmented - speed & pitch up - RAVDESS

1.1.17 Augmented - speed & pitch up- CREMA-D

1.1.18 Augmented - speed & pitch down - RAVDESS

1.1.19 Augmented - speed & pitch down- CREMA-D

1.1.20 Augmented - all - RAVDESS

1.1.21 Augmented - all - CREMA-D

```
[]: datasets = {
    'train': TRAIN_CREMA_D_AUDIO_FILES_PATH,
    'test': TEST_CREMA_D_AUDIO_FILES_PATH,
    'validate': VALIDATE_CREMA_D_AUDIO_FILES_PATH,
}

store_data(datasets=datasets,
    dataset_name='crema-d',
    is_augmented=True,
    augmentation_type='AllAugmentations')
```

1.1.22 Augmented - all + trimmed - RAVDESS

(turn on commented line in load_files_in_df())

1.1.23 Augmented - all + trimmed - CREMA-D

(turn on commented line in load_files_in_df())

1.1.24 Split Gender No Aug Original - RAVDESS

1.1.25 Split Gender No Aug Original - CREMA-D

1.1.26 Split Gender No Aug Trimmed - RAVDESS

(turn on commented line in load sex splitted files())

1.1.27 Split Gender No Aug Trimmed - CREMA-D

```
(turn on commented line in load_sex_splitted_files())
```

[]: