IE6400 Foundations Data Analytics Engineering

Fall Semester 2023

Project 3:

Topic: EEG Classification Model

Group 9:

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Importing Libraries

```
In [78]: import os
         import wfdb
         import glob
         import random
         import gc
         import mne
         import re
         import tqdm
         import logging
         import numpy as np
         import pandas as pd
         import tensorflow as tf
         from sklearn import metrics
         from tensorflow import keras
         import matplotlib.pyplot as plt
         from keras.utils import plot model
         from sklearn import model selection
         from scipy.signal import find_peaks
         from tensorflow.keras import layers
         from tensorflow.keras.callbacks import ReduceLROnPlateau, EarlyStopping
```

Data Preprocessing

```
In [22]: file path = 'D:/chb-mit-scalp-eeg-database-1.0.0/'
         folders = sorted(glob.glob(file_path+'/*/'))
         n_patient = [m[-2:] for m in [1.rsplit('\\', 2)[-2] for 1 in folders]]
         print(*n_patient)
         01 02 03 04 05 06 07 08 09 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24
         random.seed(2023)
In [23]:
         ratio train = 0.8
         train_patient_str = sorted(random.sample(n_patient, round(ratio_train*len(n_patient))))
         test patient str = sorted([l for l in n patient if l not in train patient str])
         print('Train PT: ', *train_patient_str)
         print('Test PT: ', *test_patient_str)
         Train PT: 02 03 04 05 06 09 11 12 13 14 15 16 17 18 19 20 21 23 24
         Test PT: 01 07 08 10 22
In [24]:
         files train = []
         for l in train patient str:
             files_train = files_train + glob.glob(file_path+'/chb{}/*.edf'.format(1))
         files test = []
         for l in test patient str:
             files test = files test + glob.glob(file path+'/chb{}/*.edf'.format(1))
In [25]: len(files train), len(files test)
Out[25]: (549, 137)
```

```
In [28]: logger = logging.getLogger(__name__)
         fh = logging.FileHandler('read files.log')
         logger.addHandler(fh)
         time window = 8
         time step = 4
         p = 0.01
         counter = 0
         for temp_f in files_train:
             temp_edf = mne.io.read_raw_edf(temp_f)
             temp labels = temp edf.ch names
             if sum([any([0 if re.match(c, 1)==None else 1 for 1 in temp_edf.ch_names]) for c in ch_labels])==len(ch_l
                 time_window = 8
                 time_step = 4
                 fs = int(1/(temp_edf.times[1]-temp_edf.times[0]))
                 step_window = time_window*fs
                 step = time step*fs
                 temp_is_sz = np.zeros((temp_edf.n_times,))
                 if os.path.exists(temp f+'.seizures'):
                     temp_annotation = wfdb.rdann(temp_f, 'seizures')
                     for i in range(int(temp_annotation.sample.size/2)):
                         temp_is_sz[temp_annotation.sample[i*2]:temp_annotation.sample[i*2+1]]=1
                 temp len = temp edf.n times
                 temp_is_sz_ind = np.array(
                     [temp_is_sz[i*step:i*step+step_window].sum()/step_window for i in range((temp_len-step_window)//s
                 temp_0_sample_size = round(p*np.where(temp_is_sz_ind==0)[0].size)
                 temp_1_sample_size = np.where(temp_is_sz_ind>0)[0].size
                 counter = counter + temp_0_sample_size + temp_1_sample_size
             temp_edf.close()
         array_signals = np.zeros((counter, len(ch_labels), step_window), dtype=np.float32)
         array_is_sz = np.zeros(counter, dtype=bool)
         counter = 0
         for n, temp_f in enumerate(tqdm.tqdm(files_train)):
             to_log = 'No. {}: Reading. '.format(n)
```

```
temp edf = mne.io.read raw edf(temp f)
temp labels = temp edf.ch names
n_label_match = sum([any([0 if re.match(c, 1)==None else 1 for l in temp_edf.ch_names]) for c in ch_label
if n label_match==len(ch_labels):
    ch_mapping = {sorted([1 for 1 in temp_edf.ch_names if re.match(c, 1)!=None ])[0]:c for c in ch_labels
    temp_edf.rename_channels(ch_mapping)
    temp is sz = np.zeros((temp edf.n times,))
    temp signals = temp edf.get data(picks=ch labels)*1e6
    if os.path.exists(temp f+'.seizures'):
        to log = to log+'sz exists.'
        temp annotation = wfdb.rdann(temp f, 'seizures')
        for i in range(int(temp_annotation.sample.size/2)):
            temp is sz[temp annotation.sample[i*2]:temp annotation.sample[i*2+1]]=1
    else:
        to_log = to_log+'No sz.'
    temp_len = temp_edf.n_times
    time\_window = 8
    time step = 4
    fs = int(1/(temp_edf.times[1]-temp_edf.times[0]))
    step window = time window*fs
    step = time step*fs
    temp is sz ind = np.array(
        [temp_is_sz[i*step:i*step+step_window].sum()/step_window for i in range((temp_len-step_window)//s
    del temp_is_sz
    temp 0 sample size = round(p*np.where(temp is sz ind==0)[0].size)
    temp 1 sample size = np.where(temp is sz ind>0)[0].size
    # sz data
    temp_ind = list(np.where(temp_is_sz_ind>0)[0])
    for i in temp ind:
        array_signals[counter, :, :] = temp_signals[:, i*step:i*step+step_window]
        array is sz[counter] = True
        counter = counter+1
    # no sz data
    temp_ind = random.sample(list(np.where(temp_is_sz_ind==0)[0]), temp_0_sample_size)
```

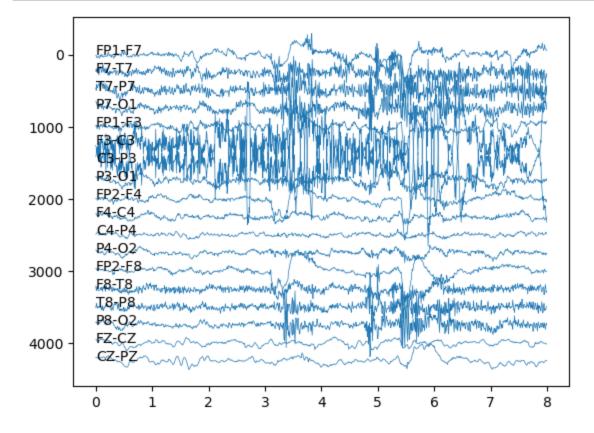
```
for i in temp_ind:
                     array_signals[counter, :, :] = temp_signals[:, i*step:i*step+step_window]
                     array is sz[counter] = False
                     counter = counter+1
                 to_log += '{} signals added: {} w/o sz, {} w/ sz.'.format(
                     temp_0_sample_size+temp_1_sample_size, temp_0_sample_size, temp_1_sample_size
             else:
                 to log += 'Not appropriate channel labels. Reading skipped.'.format(n)
             logger.info(to_log)
             temp_edf.close()
             if n%10==0:
                 gc.collect()
         gc.collect()
         np.save('signal_samples', array_signals)
         np.save('is_sz', array_is_sz)
         100%
                                                                                                 549/549 [06:10<00:00,
         1.48it/s]
In [29]: array_signals.shape
Out[29]: (9505, 18, 2048)
In [30]: | array_signals = array_signals[:, :, ::2]
```

Feature Extraction

```
In [31]: # Displaying a sample of extracted features of the EEG signals

vertical_width = 250
signals = array_signals[-1, :, :]
fs = 128

fig, ax = plt.subplots()
for i in range(signals.shape[0]):
    ax.plot(np.arange(signals.shape[-1])/fs, signals[i, :]+i*vertical_width, linewidth=0.5, color='tab:blue')
    ax.annotate(ch_labels[i], xy=(0, i*vertical_width))
ax.invert_yaxis()
plt.show()
```



```
In [32]: array_n = np.where(array_is_sz>.0)[0]
    print('Number of all the extracted signals: {}'.format(array_is_sz.size))
    print('Number of signals with seizures: {}'.format(array_n.size))
    print('Ratio of signals with seizures: {:.3f}'.format(array_n.size/array_is_sz.size))
```

Number of all the extracted signals: 9505 Number of signals with seizures: 2581 Ratio of signals with seizures: 0.272

```
In [33]: # Displaying samples with seizures.
         for n in random.sample(list(array_n), 10):
             vertical_width = 300
             temp_signals = array_signals[n, :, :]
             fs = 128
             fig, ax = plt.subplots(2, 1, figsize=(10, 6), gridspec_kw={'height_ratios': [3, 1]})
             for i in range(temp_signals.shape[0]):
                 ax[0].plot(np.arange(temp_signals.shape[-1])/fs, temp_signals[i, :]+i*vertical_width, linewidth=0.5,
                 ax[0].annotate(ch_labels[i], xy=(0, i*vertical_width))
             ax[0].invert_yaxis()
             ax[0].set_xlim(0, 8)
             ax[0].set_title('sample no. {}'.format(n))
             ax[1].pcolormesh(np.arange(temp_signals.shape[-1])/fs, np.arange(len(ch_labels)), temp_signals[:, :], cma
             ax[1].invert_yaxis()
             plt.show()
             10
                                                         sample no. 3746
           -1000
            1000
            2000 -
            3000
```

Data Splitting

Model Creation

```
In [38]: ## Creating Deep Learning model
         model = keras.models.Sequential()
         model.add(layers.Conv2D(filters=64, kernel_size=(2, 4), padding='same', activation='relu', input_shape=X_trai
         model.add(layers.Conv2D(filters=64, kernel_size=(2, 4), strides=(1, 2),padding='same', activation='relu'))
         model.add(layers.MaxPooling2D((1, 2)))
         model.add(layers.Conv2D(filters=128, kernel_size=(2, 4), padding='same', activation='relu'))
         model.add(layers.Conv2D(filters=128, kernel_size=(2, 4), strides=(1, 2), padding='same', activation='relu'))
         model.add(layers.MaxPooling2D((2, 2)))
         model.add(layers.Conv2D(filters=256, kernel_size=(4, 4), padding='same', activation='relu'))
         model.add(layers.Conv2D(filters=256, kernel size=(4, 4), strides=(1, 2), padding='same', activation='relu'))
         model.add(layers.MaxPooling2D((1, 2)))
         model.add(layers.GlobalAveragePooling2D())
         #model.add(Layers.Flatten())
         model.add(layers.Dense(256, activation='relu'))
         model.add(layers.Dropout(0.25))
         model.add(layers.Dense(128, activation='relu'))
         model.add(layers.Dense(64, activation='relu'))
         model.add(layers.Dropout(0.25))
         model.add(layers.Dense(1, activation='sigmoid'))
```

D:\Programfiles\anaconda3\Lib\site-packages\keras\src\layers\convolutional\base_conv.py:99: UserWarning: Do not pass an `input_shape`/`input_dim` argument to a layer. When using Sequential models, prefer using an `In put(shape)` object as the first layer in the model instead.

super().__init__(

In [39]: model.summary()

Model: "sequential"

Layer (type)	Output Shape	Param #
conv2d (Conv2D)	(None, 18, 1024, 64)	576
conv2d_1 (Conv2D)	(None, 18, 512, 64)	32,832
max_pooling2d (MaxPooling2D)	(None, 18, 256, 64)	0
conv2d_2 (Conv2D)	(None, 18, 256, 128)	65,664
conv2d_3 (Conv2D)	(None, 18, 128, 128)	131,200
max_pooling2d_1 (MaxPooling2D)	(None, 9, 64, 128)	0
conv2d_4 (Conv2D)	(None, 9, 64, 256)	524,544
conv2d_5 (Conv2D)	(None, 9, 32, 256)	1,048,832
max_pooling2d_2 (MaxPooling2D)	(None, 9, 16, 256)	0
global_average_pooling2d (GlobalAveragePooling2D)	(None, 256)	0
dense (Dense)	(None, 256)	65,792
dropout (Dropout)	(None, 256)	0
dense_1 (Dense)	(None, 128)	32,896
dense_2 (Dense)	(None, 64)	8,256
dropout_1 (Dropout)	(None, 64)	0
dense_3 (Dense)	(None, 1)	65

Total params: 1,910,657 (7.29 MB)

```
Trainable params: 1,910,657 (7.29 MB)
```

Non-trainable params: 0 (0.00 B)

Model Training

```
In [45]: LEARNING_RATE = 1e-4
    OPTIMIZER = tf.keras.optimizers.Adam(learning_rate=LEARNING_RATE)
    model.compile(optimizer=OPTIMIZER, loss='binary_crossentropy', metrics=['accuracy'])
    VERBOSE=1
    es = EarlyStopping(monitor='val_loss', patience=20, verbose=VERBOSE, mode='auto', restore_best_weights=True)
    callbacks = [es]

In [47]: X_train.shape, y_train.shape, X_val.shape, y_val.shape
Out[47]: ((6653, 18, 1024, 1), (6653,), (2852, 18, 1024, 1), (2852,))
```

```
In [61]: hist = model.fit(
             x=X_train, y=y_train,
             validation_data=(X_val, y_val),
             epochs=5,
             batch size=256,
             callbacks=callbacks
         Epoch 1/5
                                     478s 18s/step - accuracy: 0.7984 - loss: 0.5036 - val_accuracy: 0.8193 - val_los
         26/26 -
         s: 0.4577
         Epoch 2/5
                                    465s 18s/step - accuracy: 0.7972 - loss: 0.4930 - val_accuracy: 0.8048 - val_los
         26/26 -
         s: 0.4874
         Epoch 3/5
                                    461s 18s/step - accuracy: 0.8268 - loss: 0.4527 - val_accuracy: 0.8414 - val_los
         26/26 -
         s: 0.4087
         Epoch 4/5
         26/26 -
                                    461s 18s/step - accuracy: 0.8317 - loss: 0.4078 - val_accuracy: 0.8541 - val_los
         s: 0.3892
         Epoch 5/5
                                     462s 18s/step - accuracy: 0.8483 - loss: 0.3866 - val_accuracy: 0.8450 - val_los
         26/26 -
         s: 0.3871
         Restoring model weights from the end of the best epoch: 5.
```

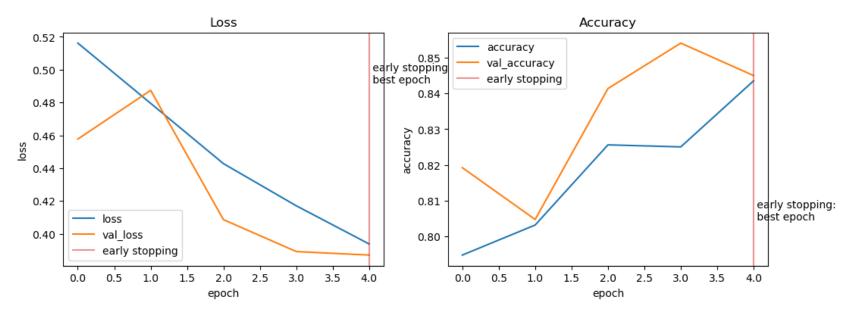
```
In [62]: model.save('CHB_MIT_sz_detec_demo.h5')
```

D:\Programfiles\anaconda3\Lib\site-packages\keras\src\model.py:342: UserWarning: You are saving your model as an HDF5 file via `model.save()`. This file format is considered legacy. We recommend using instead the native Keras format, e.g. `model.save('my_model.keras')`.

warnings.warn(

Model Evaluation and Testing

```
In [63]: fig, ax = plt.subplots(1, 2, figsize=(12, 4))
         ax[0].plot(hist.history['loss'], label='loss')
         ax[0].plot(hist.history['val_loss'], label='val_loss')
         ax[0].set xlabel('epoch')
         ax[0].set_ylabel('loss')
         ax[0].axvline(x=es.best_epoch, label='early stopping', color='tab:red', alpha=0.5)
         r = .2
         temp y = r*min(hist.history['loss'])+(1-r)*max(hist.history['loss'])
         ax[0].annotate(' early stopping:\n best epoch', xy=(es.best epoch, temp y))
         ax[0].set title('Loss')
         ax[0].legend()
         ax[1].plot(hist.history['accuracy'], label='accuracy')
         ax[1].plot(hist.history['val_accuracy'], label='val_accuracy')
         ax[1].set_xlabel('epoch')
         ax[1].set_ylabel('accuracy')
         r = .8
         temp_y = r*min(hist.history['accuracy'])+(1-r)*max(hist.history['accuracy'])
         ax[1].axvline(x=es.best epoch, label='early stopping', color='tab:red', alpha=0.5)
         ax[1].annotate(' early stopping:\n best epoch', xy=(es.best epoch, temp y))
         ax[1].set_title('Accuracy')
         ax[1].legend()
         plt.show()
```



```
def sampling_data_pred(f, verbose=True):
In [64]:
             list signals = []
             list_is_sz = []
             #n sample = 40
             if verbose==True:
                 print('{}: Reading. '.format(f))
             temp_edf = mne.io.read_raw_edf(f)
             temp_labels = temp_edf.ch_names
             if sum([any([0 if re.match(c, 1)==None else 1 for 1 in temp_edf.ch_names]) for c in ch_labels])==len(ch_1
                 ch_mapping = {sorted([1 for 1 in temp_edf.ch_names if re.match(c, 1)!=None ])[0]:c for c in ch_labels
                 temp_edf.rename_channels(ch_mapping)
                 #temp edf = temp edf.pick(ch labels)
                 temp_is_sz = np.zeros((temp_edf.n_times,))
                 temp_signals = temp_edf.get_data(picks=ch_labels)*1e6
                 if os.path.exists(f+'.seizures'):
                     if verbose==True:
                         print('sz exists.', end=' ')
                     temp_annotation = wfdb.rdann(f, 'seizures')
                     for i in range(int(temp_annotation.sample.size/2)):
                         temp_is_sz[temp_annotation.sample[i*2]:temp_annotation.sample[i*2+1]]=1
                 #else:
                     #print('No sz.', end=' ')
                 temp_len = temp_edf.n_times
                 time_window = 8
                 time step = 4
                 fs = int(1/(temp_edf.times[1]-temp_edf.times[0]))
                 step_window = time_window*fs
                 step = time step*fs
                 # sampling all signals
                 temp_array_signals = np.array([temp_signals[:, i*step:i*step+step_window] for i in range((temp_len-st
                 temp_is_sz_ind = np.array([temp_is_sz[i*step:i*step+step_window].sum()/step_window for i in range((te
             else:
                 if verbose==True:
                     print('EEG {}: Not appropriate channel labels. Reading skipped.'.format(n))
```

```
return temp_array_signals, temp_is_sz_ind
```

```
In [65]: # reading files and prediction

list_pred = []
list_true = []

for f in tqdm.tqdm(files_test):
    array_signals, array_is_sz = sampling_data_pred(f, verbose=False)
    array_signals = array_signals[:, :, ::2, np.newaxis]

list_pred.append(model.predict(array_signals, verbose=0))
    list_true.append(array_is_sz)
```

100%|**256**

137/137 [1:03:12<00:00,

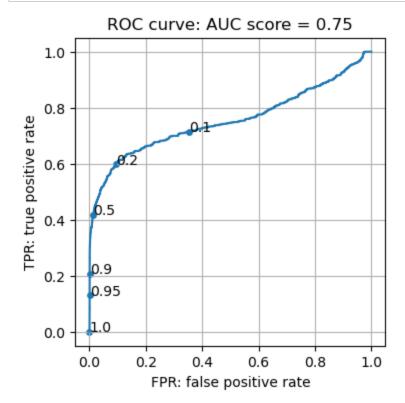
```
In [76]:
    # Setting threshold = 0.5
    report = metrics.classification_report(np.concatenate(list_true)>0, np.concatenate(list_pred)>.5)
    print(report)

# Setting threshold = 0.9
    report = metrics.classification_report(np.concatenate(list_true)>0, np.concatenate(list_pred)>.9)
    print(report)
```

	precision	recall	f1-score	support
False	1.00	0.99	0.99	186865
True	0.10	0.42	0.16	629
accuracy			0.99	187494
macro avg	0.55	0.70	0.58	187494
weighted avg	0.99	0.99	0.99	187494
	precision	recall	f1-score	support
False	precision	recall	f1-score	support 186865
False True	•			
	1.00	1.00	1.00	186865
True	1.00	1.00	1.00 0.30	186865 629

```
In [69]: roc = metrics.roc_curve(np.concatenate(list_true)>0, np.concatenate(list_pred))
auc = metrics.roc_auc_score(np.concatenate(list_true)>0, np.concatenate(list_pred))
```

```
In [70]:
         plt.figure(figsize=(4, 4))
         plt.plot(roc[0][np.argmin(np.abs(roc[2]-1)):], roc[1][np.argmin(np.abs(roc[2]-1)):])
         plt.xlabel('FPR: false positive rate')
         plt.ylabel('TPR: true positive rate')
         plt.title('ROC curve: AUC score = {:.2f}'.format(auc))
         th = [.1, .2, .5, .9, .95, 1.]
         ind = [np.argmin(np.abs(roc[2]-1)) for 1 in th]
         plt.scatter(roc[0][ind], roc[1][ind], s=15)
         for i, l in enumerate(ind):
             plt.annotate("{}".format(th[i]), xy=(roc[0][1], roc[1][1]))
         #plt.plot([0, 1, 1, 0, 0], [0, 0, 1, 1, 0], color='black', linewidth=1)
         plt.ylim(-.05, 1.05)
         plt.xlim(-.05, 1.05)
         plt.grid()
         #plt.axis('off')
         plt.show()
```



```
In [71]: for i, f in enumerate(files_test):
    if os.path.exists(f+'.seizures'):
        print('Index = {} has seizures: {}'.format(i, f))
```

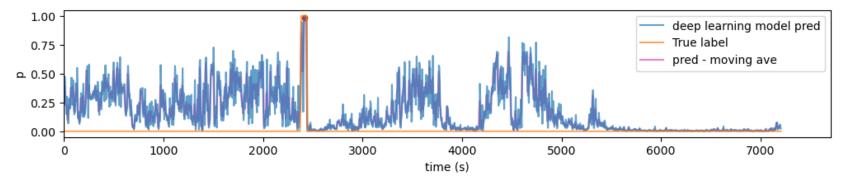
```
Index = 2 has seizures: D:/chb-mit-scalp-eeg-database-1.0.0//chb01\chb01 03.edf
Index = 3 has seizures: D:/chb-mit-scalp-eeg-database-1.0.0//chb01\chb01 04.edf
Index = 14 has seizures: D:/chb-mit-scalp-eeg-database-1.0.0//chb01\chb01 15.edf
Index = 15 has seizures: D:/chb-mit-scalp-eeg-database-1.0.0//chb01\chb01 16.edf
Index = 17 has seizures: D:/chb-mit-scalp-eeg-database-1.0.0//chb01\chb01 18.edf
Index = 20 has seizures: D:/chb-mit-scalp-eeg-database-1.0.0//chb01\chb01 21.edf
Index = 25 has seizures: D:/chb-mit-scalp-eeg-database-1.0.0//chb01\chb01 26.edf
Index = 53 has seizures: D:/chb-mit-scalp-eeg-database-1.0.0//chb07\chb07 12.edf
Index = 54 has seizures: D:/chb-mit-scalp-eeg-database-1.0.0//chb07\chb07 13.edf
Index = 60 has seizures: D:/chb-mit-scalp-eeg-database-1.0.0//chb07\chb07 19.edf
Index = 61 has seizures: D:/chb-mit-scalp-eeg-database-1.0.0//chb08\chb08 02.edf
Index = 64 has seizures: D:/chb-mit-scalp-eeg-database-1.0.0//chb08\chb08 05.edf
Index = 66 has seizures: D:/chb-mit-scalp-eeg-database-1.0.0//chb08\chb08 11.edf
Index = 68 has seizures: D:/chb-mit-scalp-eeg-database-1.0.0//chb08\chb08_13.edf
Index = 76 has seizures: D:/chb-mit-scalp-eeg-database-1.0.0//chb08\chb08 21.edf
Index = 89 has seizures: D:/chb-mit-scalp-eeg-database-1.0.0//chb10\chb10 12.edf
Index = 97 has seizures: D:/chb-mit-scalp-eeg-database-1.0.0//chb10\chb10 20.edf
Index = 100 has seizures: D:/chb-mit-scalp-eeg-database-1.0.0//chb10\chb10_27.edf
Index = 102 has seizures: D:/chb-mit-scalp-eeg-database-1.0.0//chb10\chb10_30.edf
Index = 103 has seizures: D:/chb-mit-scalp-eeg-database-1.0.0//chb10\chb10 31.edf
Index = 104 has seizures: D:/chb-mit-scalp-eeg-database-1.0.0//chb10\chb10_38.edf
Index = 105 has seizures: D:/chb-mit-scalp-eeg-database-1.0.0//chb10\chb10 89.edf
Index = 122 has seizures: D:/chb-mit-scalp-eeg-database-1.0.0//chb22\chb22_20.edf
Index = 127 has seizures: D:/chb-mit-scalp-eeg-database-1.0.0//chb22\chb22 25.edf
Index = 133 has seizures: D:/chb-mit-scalp-eeg-database-1.0.0//chb22\chb22 38.edf
```

Results and Visualization

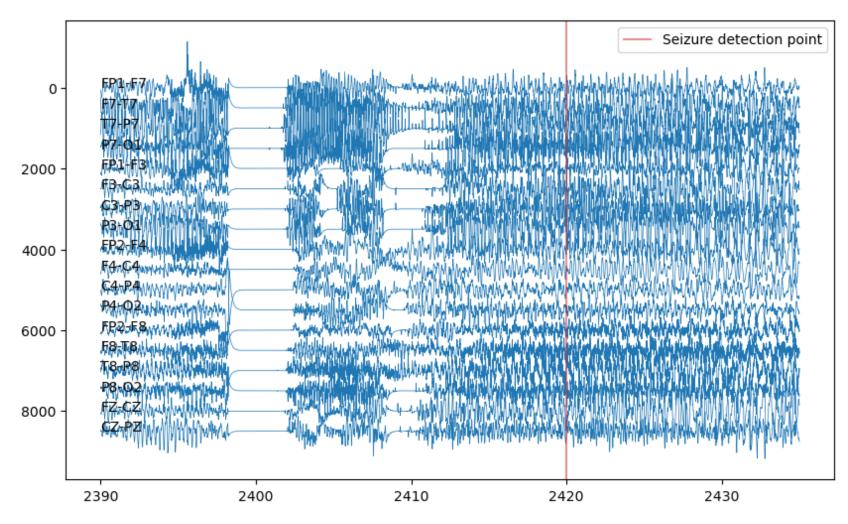
57/57 35s 606ms/step

```
In [72]: def moving_ave(a, n):
             if len(a.shape)!=1:
                 print('Not 1 dimension array. return nothing.')
             temp = np.zeros(a.size-n)
             for i in range(n):
                 temp = temp+a[i:-n+i]
             temp = temp/n
             return temp
In [73]: # Getting signals and labels from test data.
         n=100
         array_signals, array_is_sz = sampling_data_pred(files_test[n])
         # Preprocessing the Test data
         array_signals=array_signals[:, :, ::2, np.newaxis]
         # Prediction using the created deep learning model
         pred = model.predict(array_signals)
         D:/chb-mit-scalp-eeg-database-1.0.0//chb10\chb10_27.edf: Reading.
```

```
In [74]: | time_window = 8
         time step = 4
         mv_win = 3
         fig, ax = plt.subplots(figsize=(12, 2))
         ax.plot(np.arange(pred.size)*time_step, pred.flatten(), alpha=0.7, label='deep learning model pred')
         ax.plot(np.arange(pred.size)*time_step, array_is_sz, alpha=.7, label='True label')
         pred_moving_ave = moving_ave(pred.flatten(), mv_win)
         pred_peaks, _ = find_peaks(pred_moving_ave, height=.95, distance=6)
         ax.plot(np.arange(pred.size-mv_win)*time_step, pred_moving_ave,
                 alpha=.9, label='pred - moving ave', color='tab:pink', zorder=0)
         ax.scatter(pred_peaks*time_step, pred_moving_ave[pred_peaks], s=20, color='tab:red')
         ax.set_xlabel('time (s)')
         ax.set_ylabel('p')
         ax.set_xlim(0, pred.size*time_step+500)
         ax.legend(loc='upper right')
         plt.show()
```



```
In [75]: if pred peaks.size==0:
             print('No seizure detected.')
         else:
             f = files test[n]
             temp edf = mne.io.read raw edf(f)
             temp labels = temp edf.ch names
             if sum([any([0 if re.match(c, 1)==None else 1 for 1 in temp edf.ch names]) for c in ch labels])==len(ch 1
                 ch mapping = {sorted([1 for 1 in temp edf.ch names if re.match(c, 1)!=None ])[0]:c for c in ch labels
                 temp edf.rename channels(ch mapping)
                 #temp edf = temp edf.pick(ch labels)
                 temp_is_sz = np.zeros((temp_edf.n_times,))
                 temp signals = temp edf.get data(picks=ch labels)*1e6
             fs = int(1/(temp_edf.times[1]-temp_edf.times[0]))
             for n peak in range(pred peaks.size):
                 ind peak = pred peaks[n peak]*time step*fs
                 backward steps = 30*fs
                 forward steps = 15*fs
                 vertical width=500
                 fig, ax = plt.subplots(figsize=(10, 6))
                 for i in range(temp_signals.shape[0]):
                     ax.plot(np.arange(ind_peak-backward_steps, ind_peak+forward_steps)/fs,
                             temp_signals[i, ind_peak-backward_steps:ind_peak+forward_steps]+i*vertical width, linewid
                     ax.annotate(ch_labels[i], xy=((ind_peak-backward_steps)/fs, i*vertical_width))
                 ax.axvline(x=ind peak/fs, color='tab:red', alpha=0.5, label='Seizure detection point')
                 ax.invert yaxis()
                 ax.legend(loc='upper right')
                 plt.show()
             \#ax.set xlim(0, 8)
             temp_edf.close()
```



In []: