In [2]:

```
#data structures
import pandas as pd
import pyarrow.parquet as pq
import numpy as np
#used for feature engineering(signal processing tools)
from scipy.fftpack import fft
from scipy.signal import welch
from siml.sk utils import *
from siml.signal analysis utils import *
from sklearn.model selection import StratifiedKFold, train test split, RandomizedSear
from sklearn.metrics import matthews_corrcoef,make_scorer
from catboost import CatBoostClassifier
from lightgbm import LGBMClassifier
from sklearn.linear model import LogisticRegression
from sklearn.ensemble import RandomForestClassifier
from tgdm import tgdm
import ast
import pickle
import warnings
warnings.filterwarnings('ignore')
```

In [14]:

```
def compute spectra(signals, *, m = 1000):
    computes the mean and percentiles by combining different phases belonging to th
   means = []
   percentiles = []
   percentile values = (100,99,95,0,1,5)
    for raw signal in tqdm(signals):
        #normalizing the signal values
        signal = raw signal.astype('float32').reshape(-1, m) / 128.0
        #mean of signal
        mean = np.mean(signal,axis=1)
        #percentiles of signal
        percentile = np.abs(np.percentile(signal,percentile values,axis=1)-mean)
        #calaculating the baseline of percentiles
        baseline = np.percentile(percentile,5.0)
        #subtracting the baseline
        percentile = np.maximum(0.0,percentile-baseline)
        means.append(mean)
        percentiles.append(percentile.T)
    res = \{\}
    res['mean'] = np.array(means)
    res['percentile'] = np.array(percentiles)
    return res
```

In [15]:

```
#https://www.kaggle.com/junkoda/handmade-features
import tensorflow as tf
tf.compat.v1.disable eager execution()
def max windowed(spec, *, width=150, stride=10):
    Smooth the spectrum with a tophat window function and find the
    peak inteval that maximises the smoothed spectrum.
    Returns: d(dict)
      d['w'] (array): smoothed max - mean spectrum
     d['ibegin'] (array): the left edge index of the peak interval
   n = spec.shape[0]
   length = spec.shape[1] \# 800
   nspec = spec.shape[2] # 6 spectra
   n \text{ triplet} = n // 3
   # Reorganize the max spectrum from 8712 data to 2904 triplets with 3 phases
   max spec3 = np.empty((n triplet, length, 3))
    for i triplet in range(n triplet):
        max_spec3[i_triplet, :, 0] = spec[3*i_triplet, :, 0] # phase 0
        max spec3[i triplet, :, 1] = spec[3*i triplet + 1, :, 0] # phase 1
        max spec3[i triplet, :, 2] = spec[3*i triplet + 2, :, 0] # phase 2
   x = tf.compat.v1.placeholder(tf.float32, [None, length, 3]) # input spectra bef
   # 800 -> 80: static convolaution
   # convolution but not CNN, the kernel is static
   # smoothing/convolution kernel
   # tophat window function
   # shape (3, 1) adds up 3 phases to one output
   K = np.ones((width, 3, 1), dtype='float32') / width
   W conv1 = tf.constant(K)
   h conv1 = tf.nn.conv1d(x, W conv1, stride=stride, padding='VALID')
   with tf.compat.v1.Session() as sess:
       w = sess.run(h_conv1, feed_dict={x:max_spec3})
   imax = np.argmax(w[:, :, 0], axis=1) # index of maximum smoothed spectrum
   d = \{\}
    d['w'] = w # smoothed max spectrum
    d['ibegin'] = imax*stride
    return d
```

In [16]:

```
def compute spectra features(spectra, peaks):
    extracts the features from peaks found in percentiles and means
   percentiles = spectra['percentile']
    n = percentiles.shape[0]
    length = percentiles.shape[1]
    nspec = percentiles.shape[2]
   n signals = n//3
   phase percentiles = np.empty((n signals,length,nspec,3))
   #creating an array which combines all 3 phases
    for i signal in range(n signals):
        phase_percentiles[i_signal, :, :, 0] = percentiles[3*i_signal, :, :] # phas
        phase_percentiles[i_signal, :, :, 1] = percentiles[3*i_signal + 1, :, :] #
        phase_percentiles[i_signal, :, :, 2] = percentiles[3*i_signal + 2, :, :] #
   width = 150
   n perc features = 3
   #array to store final features
    spectra_features = np.empty((n_signals,n_perc features*nspec*3 + 3))
   #array to store percentile features
    perc phase features = np.empty((n signals,n perc features,nspec,3))
    for i signal in range(n signals):
        #max of the total percentile features
        perc phase features[i signal,0,:,:] = np.max(phase percentiles[i signal,:,:
        peak_start = peaks['ibegin'][i_signal]
        peak end = peak start +width
        peak_mid = peak_start+width//2
        #mean of the percentle features in peak interval
        perc_phase_features[i_signal,1,:,:] = np.mean(phase_percentiles[i_signal,pe
        #max of the percentle features in peak interval
        perc phase features[i signal,2,:,:] = np.max(phase percentiles[i signal,pea
        #storing the mean value at the mid index of peak interval of each phase
        spectra_features[i_signal,0] = spectra['mean'][3*i_signal,peak_mid]
        spectra_features[i_signal,1] = spectra['mean'][3*i_signal+1,peak_mid]
        spectra features[i signal,2] = spectra['mean'][3*i signal+2,peak mid]
   #storing all the features
    shape = perc_phase_features.shape
    spectra features[:,3:] = perc phase features.reshape(shape[0], shape[1]*shape[2
    return spectra features
```

In [17]:

```
def extract_fourier_features(signal,N=800000,T=1/50):
    converts a signal from time spectrum to frequency spectrum
    and returns only the features required as mentioned above
    fourier_values = fft(signal)
    fourier_values_filtered = 2.0/N * np.abs(fourier_values[0:N//2])
    return fourier_values_filtered
```

In [18]:

```
def filter_features_fourier(features,mph,no_features=8):
    returns fourier transformed features by extracting peaks and
    considering only required number of peaks.
    mph-detect peaks that are greater than minimum peak height
    indices_peaks = detect_peaks(features,mph = mph)
    #print(indices_peaks)
    values = features[indices_peaks]
    if len(values) < no_features:
        return np.append(values , [0]*(no_features-len(values)))
    else:
        return values[:no_features]</pre>
```

In [19]:

```
def extract_psd_features(signal,fs=50):
    returns the features from a time spectrum signal(similiar to fft
    but also considers power spectral density)
    f_values, psd_values = welch(signal, fs=50)
    return psd_values
```

In [82]:

```
def extract_stat_features(signal_features):
    used to extract statistical features from power spectral density
    percentiles = (5,25,50,75)
#array to store
filtered_features = np.zeros(8)
#mean
filtered_features[0] = np.mean(signal_features,axis=0)
#standard deviation
filtered_features[1] = np.std(signal_features,axis=0)
#maximum
filtered_features[2] = np.max(signal_features,axis=0)
#minimum
filtered_features[3] = np.min(signal_features,axis=0)
#mean
filtered_features[4] = np.mean(signal_features,axis=0)
#percentiles
filtered_features[4:] = np.percentile(signal_features,percentiles,axis=0).T
return filtered_features
```

In [114]:

```
def final fun 1(X):
    returns the prediction for raw input
    spectra = compute spectra(X)
    peaks = max windowed(spectra['percentile'])
    spectra features = compute spectra features(spectra, peaks)
    signal features = []
    for i in tqdm(range(0,len(X),3)):
        std = 0
        maxs = 0
        bw low = 0
        #arrays to store features
        fourier_features = np.empty((3,8))
        psd features = np.empty((3,129))
        features = []
        for j in range(3):
            signal = raw signal data[i+j]
            #standard deviation
            std += signal.std()
            #max of signal
            maxs += signal.max()
            #lower bandwidth
            bw low += signal.mean()-signal.std()
            #minimum peak height which can be used to filter fourier features
            mph = signal.min() + (signal.max() - np.abs(signal.min()))/10
            #fourier features
            fourier_features_ = extract_fourier_features(signal)
            fourier features[j,:] = filter features fourier(fourier features ,mph)
            #power spectral density features
            psd_features[j,:] = extract_psd_features(signal)
        #calculating the average of above features
        features.append(std/3)
        features.append(maxs/3)
        features.append(bw_low/3)
        features.extend(np.mean(fourier features,axis=0))
        features.extend(extract_stat_features(np.mean(psd_features,axis=0)))
        signal features.append(features)
```

```
total_features = np.concatenate((spectra_features, signal_features), axis=1)
#loading the saved models
models = []
for i in range(20):
    filename = 'finalmodels/model'+str(i)+'.sav'
    model = pickle.load(open(filename, 'rb'))
    models.append(model)
#prediction
y_test_probas = np.empty((total_features.shape[0], 20))
for i, model in enumerate(models):
    y_test_probas[:, i] = model.predict_proba(total_features)[:, 1]
#taking mean of all the predicted
y_test_proba = np.mean(y_test_probas, axis=1)
# Converting to 0 1 with a threshold 0.25, then replicating 3 copies for 3 phas
y_pred = np.repeat(y_test_proba > 0.25, 3)
return y pred
```

In [111]:

```
def final fun 2(X,Y):
    returns the matthews correlation score for raw input
    spectra = compute spectra(X)
    peaks = max windowed(spectra['percentile'])
    spectra features = compute spectra features(spectra, peaks)
    signal features = []
    for i in tqdm(range(0,len(X),3)):
        std = 0
        maxs = 0
        bw low = 0
        #arrays to store features
        fourier_features = np.empty((3,8))
        psd features = np.empty((3,129))
        features = []
        for j in range(3):
            signal = raw signal data[i+j]
            #standard deviation
            std += signal.std()
            #max of signal
            maxs += signal.max()
            #lower bandwidth
            bw low += signal.mean()-signal.std()
            #minimum peak height which can be used to filter fourier features
            mph = signal.min() + (signal.max() - np.abs(signal.min()))/10
            #fourier features
            fourier_features_ = extract_fourier_features(signal)
            fourier features[j,:] = filter features fourier(fourier features ,mph)
            #power spectral density features
            psd_features[j,:] = extract_psd_features(signal)
        #calculating the average of above features
        features.append(std/3)
        features.append(maxs/3)
        features.append(bw_low/3)
        features.extend(np.mean(fourier features,axis=0))
        features.extend(extract_stat_features(np.mean(psd_features,axis=0)))
        signal features.append(features)
```

```
total_features = np.concatenate((spectra_features, signal_features), axis=1)
#loading the saved models
models = []
for i in range(20):
    filename = 'finalmodels/model'+str(i)+'.sav'
    model = pickle.load(open(filename, 'rb'))
    models.append(model)
#prediction
y test probas = np.empty((total features.shape[0], 20))
for i, model in enumerate(models):
    y_test_probas[:, i] = model.predict_proba(total_features)[:, 1]
#taking mean of all the predicted
y_test_proba = np.mean(y_test_probas, axis=1)
# Converting to 0 1 with a threshold 0.25, then replicating 3 copies for 3 phas
y_pred = np.repeat(y_test_proba > 0.25, 3)
return matthews corrcoef(y pred,Y)
```

In []: