EMATS: EEG based Machine or Deep Learning Algorithms for TBI & Stroke Classification

Predict New Classification from EEG

Michael Caiola (Michael.Caiola@fda.hhs.gov) and Meijun Ye (Meijun.Ye@fda.hhs.gov)

This script contains two sections:

1. Preprocess an .edf file for future classification
2. Predict the classification of a preprocessed EEG

Preprocess EDF

Input: EEG .edf file

- **>4 minutes in length**
- **>= 19 contacts**
- **>= 250 Hz sampling rate**

`Requirements:`

- `EEGLAB with BIOSIG toolbox`
- `Signal Processing Toolbox`

Run this section to convert your EDF into a usable preprocessed .mat file(s). The file will be cut up into 3-minute segments with the first minute discarded. Preprocessed files will be saved as .mat files.

```
LoadEDF();
```

Predict

`For use with a preprocessed EEG .mat file (see above).`

`Requirements:`

- `Signal Processing Toolbox`
- `Deep Learning Toolbox`
- `EEGLAB (only necessary if using TMN)`

Choose model type:

- **STFT**: Short-time Fourier Transform
- **TMN**: Topographic Map Network

- **Sensor Fusion**
- **Feature**: Deep Network using ReliefF Features
- *NOTE: Requires calculation of features: long/expensive.*
- **LDA_SVM**: SVM with LDA Features.
- *NOTE: Requires calculation of features: long/expensive. Will not provide a score*
- **ReliefF_SVM** with ReliefF Features.
- *NOTE: Requires calculation of features: long/expensive.Will not provide a score*

```
model = "STFT";
```

```
[fileLoc,path] = uigetfile(".mat","Select processed EEG file");

%Set up function
switch model
    case {"LDA","ReliefF","Feature"}
        load(fullfile(path,fileLoc))
        F = BigFeats(y);
        F = table(F','VariableNames',{'Features'});

    case "TMN"
        load("chlocs2.mat")
        ds = TopoDatastore(fullfile(path,fileLoc),[],channel_locations);
        y = read(ds);
        y = y.Predictors{1};

    case {"STFT","Fusion"}
        ds = ResampleDatastore(fullfile(path,fileLoc),100,'DataAugmentation',false);
        ds.MiniBatchSize = 1;
        reset(ds)
        y = read(ds);
        y = y.Predictors{1};

end

%Predict
switch model
    case "LDA"
        load("LDA_SVM.mat")
        F = table(F.Features(1,r_logical),'VariableNames',{'Features'});
        [yfit,score] = trainedModel.predictFcn(F);
        disp("EEG prediction: " + string(yfit));
    case "ReliefF"
        load("ReliefF_SVM.mat")
        [yfit,score] = trainedModel1.predictFcn(F);
```

```
        disp("EEG prediction: " + string(yfit));
    case "Feature"
        load("F_DL.mat","net5b")
        load("RelieffScores.mat")
        F = F.Features(1,featureIndex(1:100));
        [yfit,score] =
classify(net5b,F,"ExecutionEnvironment",'cpu',MiniBatchSize=1);
        disp("EEG prediction: " + string(yfit));
        disp("Prediction score: " + num2str(max(score)));
    case "TMN"
        load("Topo_BasicNet.mat","net3")
        [yfit,score] =
classify(net3,y,"ExecutionEnvironment",'cpu',MiniBatchSize=1);
        disp("EEG prediction: " + string(yfit));
        disp("Prediction score: " + num2str(max(score)));
    case "STFT"
        load("STFTNet.mat","net4a")
        [yfit,score] =
classify(net4a,y,"ExecutionEnvironment",'cpu',MiniBatchSize=1);
        disp("EEG prediction: " + string(yfit));
        disp("Prediction score: " + num2str(max(score)));
    case "Fusion"
        load("SFnet.mat","bnet2")
        [yfit,score] =
classify(bnet2,y,"ExecutionEnvironment",'cpu',MiniBatchSize=1);
        disp("EEG prediction: " + string(yfit));
        disp("Prediction score: " + num2str(max(score)));
end
```

```
EEG prediction: HEA
Prediction score: 0.72235
```

CAUTION: EEG classification is set using default thresholding and should be further optimized for `sensitivity/specificity/accuracy` on additional training data prior to use.

Disclaimer

Supporting Functions

Loading and Cleaning

```
function LoadEDF(savefolder)
N=3;
[edfFile,path] = uigetfile("*.edf","Choose .edf File");
if nargin < 1
    savefolder= uigetdir([],"Choose Save Folder");
end
loadEEGLAB();
rawEEG=joinEEG(fullfile(path,edfFile));
if isempty(rawEEG.data)
    warning("Issue finding Channels.")
    return
end
data=cutEEG(rawEEG,N);
if isempty(data)
    warning("Issue splitting EEG.")
    return
end
try
    clndata=filtEEG(rawEEG,data);
catch
    warning("Could not clean EEG.")
    return
end
for kk=1:size(clndata,3)
    y=clndata(:,:,kk);
    if or(isempty(y),y==zeros(size(y)))
        continue
    end
    try
        save(fullfile(savefolder,"ProcessedEEG",erase(string(edfFile),".edf")
+"-"+string(kk)+".mat"),'y')
    catch
        mkdir(fullfile(savefolder,"ProcessedEEG"))
        save(fullfile(savefolder,"ProcessedEEG",erase(string(edfFile),".edf")
+"-"+string(kk)+".mat"),'y')
    end
end
disp("EEG successfully exported!")
end
```

```
function rawEEG=joinEEG(names)
out=0;
cnt = 0;
```

```
try
    [rawEEG] = FindCh(names);
    param_flag = 0;
catch
    rawEEG.comments=[];
    rawEEG.data=[];
    param_flag=1;
end

if param_flag
    rawEGG=[];
else
    if rawEEG.xmax < 3*60
        error(".edf is too short.")
    end
    rawEEG.data=rawEEG.data(:,1:250*60*(floor(length(rawEEG.data)/250/60)-1));
    rawEEG.xmax=length(rawEEG.data)/250;
    rawEEG.pnts=length(rawEEG.data);
    rawEEG.times=0:4:4*(rawEEG.pnts-1);
end
end
```

```
function data=cutEEG(EEG,n)
cuts=floor(EEG.xmax/60/n);
data=zeros(19,n*60*250,cuts);
for i=1:cuts
    data(:,:,i)=EEG.data(:,1+(i-1)*n*60*250:i*n*60*250);
end
end
```

```
function [clndata]=filtEEG(EEG,data)
for i=1:size(data,3)
    rawEEG=EEG;
    rawEEG.data=data(:,:,i);
    rawEEG.pnts=length(rawEEG.data);
    filtEEG = pop_eegfiltnew(rawEEG,1,100);
    EEG = pop_reref(filtEEG,[]);
    [~,W] = fastica(EEG.data,'verbose','off');
    EEG = pop_editset(EEG,'icaweights',W);
    EEG = iclabel(EEG);
    [~,ictype] = max(EEG.etc.ic_classification.ICLabel.classifications,[],2);
    icreject = find(ictype~=1);
    % EEG.reject.gcomprejject(1,icreject') = 1;
    if ~(length(icreject) == size(EEG.icaweights,1))
        clean_EEG = pop_subcomp(EEG,icreject',0,0);
```

```
        %rawdata(:,:,i) = EEG.data;
        clndata(:,:,i) = clean_EEG.data;
    end
end
end
```

```
function [rawEEG,cnt] = FindCh(file)
cnt = 0;
load('chlocs2.mat','channel_locations');
ch_locs = struct2table(channel_locations);
ch_locs = string(ch_locs.labels);
rawEEG = pop_biosig(file,'blockrange',[60
Inf],'importevent','off','importannot','off');
chs = struct2table(rawEEG.chanlocs);
chs = string(chs.labels);
chs = erase(chs,"-REF"|"-LE");
rawEEG = pop_resample(rawEEG,250);
in = [];
for i = 1:length(ch_locs)
    in1 = find(strcmpi(chs,ch_locs(i)));
    in = [in in1];
    if and(isempty(in1),ismember(ch_locs(i),["T3";"T4";"T5";"T6"]))
        switch ch_locs(i)
            case "T3"
                in2 = find(strcmpi(chs,"T7"));
            case "T4"
                in2 = find(strcmpi(chs,"T8"));
            case "T5"
                in2 = find(strcmpi(chs,"P7"));
            case "T6"
                in2 = find(strcmpi(chs,"P8"));
        end
        in = [in in2];
    end
end
if length(in)~=19
    error("Could not locate all channels!");
end
rawEEG.data = rawEEG.data(in,:);
rawEEG.chanlocs = channel_locations;
rawEEG.nbchan = 19;
if ~isequal(in,[1:16,19:21])
    cnt = 1;
    if ~isequal(in,1:19)
        cnt = 2;
    end
end
end
```

```
function loadEEGLAB()
if exist("pop_biosig")~=2
    path = pwd;
    try
        cd(fullfile("..","eeglab"))
    catch
        disp("EEGLAB not found.")
        eeglabpath = uigetdir([],"Locate EEGLAB Directory");
        cd(eeglabpath)
    end
    eeglab
    close
    cd(path);
end
end
```

Feature Calculation

```
function F = BigFeats(data)
srate=250;
numFeats = 19*2 + 171*5;
idx = 0;
h = waitbar(0,"Calculating Features");
tflt = designfilt('bandpassiir','FilterOrder',6, ...
    'HalfPowerFrequency1',4,'HalfPowerFrequency2',8, ...
    'SampleRate',srate);
aflt = designfilt('bandpassiir','FilterOrder',6, ...
    'HalfPowerFrequency1',8,'HalfPowerFrequency2',12, ...
    'SampleRate',srate);
gflt = designfilt('bandpassiir','FilterOrder',6, ...
    'HalfPowerFrequency1',25,'HalfPowerFrequency2',40, ...
    'SampleRate',srate);
F=zeros(380,size(data,3));
for j=1:size(data,3)
    stats = []; %mean, max, min, std 76 values
    spectent = []; % 19 Values
    PACag = []; % 19 Values
    PACtg = []; % 19 Values
    PACta = []; % 19 Values
    [abs_psd,rel_psd]=getPSD(data(:,:,j),srate);
    for i=1:19
        %% Spectral Entropy of each channel
        X = fft(data(i,:,j));
        S = abs(X).^2;
        P = S./sum(S);
        H = 0;
```

```
        for m = 1:length(P)
            H = H + P(m)*log2(P(m));
        end
        spectent = [spectent,-H];
        PACag = [PACag, getPAC(data(i,:,j),aflt,gflt)];
        PACtg = [PACtg, getPAC(data(i,:,j),tflt,gflt)];
        PACta = [PACta, getPAC(data(i,:,j),tflt,aflt)];
        idx = idx + 1;
        waitbar(idx/(numFeats),h)
    end
    stats=[mean(data(:,:,j),2)',max(data(:,:,j),[],2)',min(data(:,:,j),
[],2)',std(data(:,:,j),[],2)'];
    F(:,j)=[stats,reshape(abs_psd,[1,19*6]),reshape(rel_psd,
[1,19*6]),spectent,PACag,PACtg,PACta];
end
tflt = designfilt('bandpassiir','FilterOrder',6, ...
    'HalfPowerFrequency1',4,'HalfPowerFrequency2',8, ...
    'SampleRate',srate);
aflt = designfilt('bandpassiir','FilterOrder',6, ...
    'HalfPowerFrequency1',8,'HalfPowerFrequency2',12, ...
    'SampleRate',srate);
gflt = designfilt('bandpassiir','FilterOrder',6, ...
    'HalfPowerFrequency1',25,'HalfPowerFrequency2',40, ...
    'SampleRate',srate);
dflt = designfilt('bandpassiir','FilterOrder',6, ...
    'HalfPowerFrequency1',1,'HalfPowerFrequency2',4, ...
    'SampleRate',srate);
mflt = designfilt('bandpassiir','FilterOrder',6, ...
    'HalfPowerFrequency1',12,'HalfPowerFrequency2',16, ...
    'SampleRate',srate);
bflt = designfilt('bandpassiir','FilterOrder',6, ...
    'HalfPowerFrequency1',16,'HalfPowerFrequency2',20, ...
    'SampleRate',srate);
RCH=zeros(size(data,3),171*6);
for j=1:size(data,3)
    tdata = zeros(size(data,1),size(data,2));
    adata = zeros(size(data,1),size(data,2));
    mdata = zeros(size(data,1),size(data,2));
    bdata = zeros(size(data,1),size(data,2));
    gdata = zeros(size(data,1),size(data,2));
    ddata = zeros(size(data,1),size(data,2));

    for ch = 1:19
        tdata(ch,:) = filter(tflt,data(ch,:,j));
        adata(ch,:) = filter(aflt,data(ch,:,j));
        mdata(ch,:) = filter(mflt,data(ch,:,j));
        bdata(ch,:) = filter(bflt,data(ch,:,j));
        gdata(ch,:) = filter(gflt,data(ch,:,j));
        ddata(ch,:) = filter(dflt,data(ch,:,j));
        idx = idx + 1;
```

```
        waitbar(idx/(numFeats),h)
    end

    [tCH,idx] = getCOH(tdata,srate,h,idx,numFeats);
    [aCH,idx] = getCOH(adata,srate,h,idx,numFeats);
    [mCH,idx] = getCOH(mdata,srate,h,idx,numFeats);
    [bCH,idx] = getCOH(bdata,srate,h,idx,numFeats);
    [gCH,idx] = getCOH(gdata,srate,h,idx,numFeats);
    [dCH,idx] = getCOH(ddata,srate,h,idx,numFeats);

    close(h)

    RCH(j,:) = [tCH,aCH,mCH,bCH,gCH,dCH];
end
F = [F; RCH'];
end
```

```
function [abs_psd,rel_psd]=getPSD(data,srate)
% data: sample X channel
data=data';
fband=[1 4; 4 8; 8 12; 12 16; 16 20; 25 40];
tot=bandpower(data,srate,[1 srate/2]);

n=size(fband,1);

for i=1:n
    abs_psd(:,i)=bandpower(data,srate,fband(i,:));
end

ch=max(size(abs_psd,1));

for j=1:ch
    rel_psd(j,:)=abs_psd(j,:)./tot(j);
end
end
```

```
function PAC = getPAC(data,pflt,aflt)

fp = filter(pflt,data);
fp = hilbert(fp);
phi = angle(fp).*(180/pi);

fa = filter(aflt,data);
fa = hilbert(fa);
A = abs(fa);
```

```
edges = -180:20:180;
bin_phi = discretize(phi,edges);
A_mean = zeros(1,18);
for b = 1:18
    A_mean(b) = mean(A(bin_phi==b));
end


Pj = zeros(1,18);
for j = 1:18
    Pj(j) = A_mean(j)/sum(A_mean);
end
Dkl = zeros(1,18);
for k = 1:18
    Dkl(k) = log(Pj(k)/(1/18))*Pj(k);
end
PAC = sum(Dkl)/log(18);
end
```

```
function [C,idx] = getCOH(data,srate,h,idx,numFeats)
    ind = [ 0,  0,  0,  0,  0,  0,  0,  0,  0,  0,  0,  0,  0,  0,  0,...
        0,  0,  0,  1,  1,  1,  1,  1,  1,  1,  1,  1,  1,  1,  1,  1,...
        1,  1,  1,  2,  2,  2,  2,  2,  2,  2,  2,  2,  2,  2,  2,  2,...
        2,  2,  2,  3,  3,  3,  3,  3,  3,  3,  3,  3,  3,  3,  3,  3,...
        3,  3,  4,  4,  4,  4,  4,  4,  4,  4,  4,  4,  4,  4,  4,  4,...
        5,  5,  5,  5,  5,  5,  5,  5,  5,  5,  5,  5,  5,  6,  6,  6,...
        6,  6,  6,  6,  6,  6,  6,  6,  6,  7,  7,  7,  7,  7,  7,  7,...
        7,  7,  7,  7,  8,  8,  8,  8,  8,  8,  8,  8,  8,  8,  9,  9,...
        9,  9,  9,  9,  9,  9,  9, 10, 10, 10, 10, 10, 10, 10, 10, 11,...
       11, 11, 11, 11, 11, 11, 12, 12, 12, 12, 12, 12, 13, 13, 13, 13,...
       13, 14, 14, 14, 14, 15, 15, 15, 16, 16, 17; 1,  2,  3,  4,  5,...
        6,  7,  8,  9, 10, 11, 12, 13, 14, 15, 16,...
       17, 18,  2,  3,  4,  5,  6,  7,  8,  9, 10, 11, 12, 13, 14, 15,...
       16, 17, 18,  3,  4,  5,  6,  7,  8,  9, 10, 11, 12, 13, 14, 15,...
       16, 17, 18,  4,  5,  6,  7,  8,  9, 10, 11, 12, 13, 14, 15, 16,...
       17, 18,  5,  6,  7,  8,  9, 10, 11, 12, 13, 14, 15, 16, 17, 18,...
        6,  7,  8,  9, 10, 11, 12, 13, 14, 15, 16, 17, 18,  7,  8,  9,...
       10, 11, 12, 13, 14, 15, 16, 17, 18,  8,  9, 10, 11, 12, 13, 14,...
       15, 16, 17, 18,  9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 10, 11,...
       12, 13, 14, 15, 16, 17, 18, 11, 12, 13, 14, 15, 16, 17, 18, 12,...
       13, 14, 15, 16, 17, 18, 13, 14, 15, 16, 17, 18, 14, 15, 16, 17,...
       18, 15, 16, 17, 18, 16, 17, 18, 17, 18, 18]+1;
   C = zeros(1,length(ind));

    for c = 1:length(ind)
        y = mscohere(data(ind(1,c),:),data(ind(2,c),:),...
                        srate*30,0,1:0.1:100,srate);
        C(c) = mean(y);
```

```
        idx = idx + 1;
        waitbar(idx/numFeats,h)
    end

end
```