

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

## Predict New Classification from EEG

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This script contains two sections:

1. [Preprocess an .edf file for future classification](#)
2. [Predict the classification of a preprocessed EEG](#)

For easier visibility, you can hide the code by going to View > Hide Code

## 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();
```

```
sopen mode is "OVERFLOWDETECTION:OFF"
Reading data in EDF format...
eeg_checkset note: upper time limit (xmax) adjusted so (xmax-xmin)*srate+1 = number of frames
Detected/removing 'EEG' prefix from channel labels
resampling data 250.0000 Hz
1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32
resampling finished
pop_eegfiltnew() - performing 827 point bandpass filtering.
pop_eegfiltnew() - transition band width: 1 Hz
pop_eegfiltnew() - passband edge(s): [1 100] Hz
pop_eegfiltnew() - cutoff frequency(ies) (-6 dB): [0.5 100.5] Hz
pop_eegfiltnew() - filtering the data (zero-phase, non-causal)
firfilt(): |=====| 100%, ETE 00:00
Re-referencing data
eeg_checkset note: upper time limit (xmax) adjusted so (xmax-xmin)*srate+1 = number of frames
Warning: converting input data into regular (double) precision.
ICLabel: extracting features...
```

```

Scaling components to RMS microvolt
Recomputing ICA activations
ICLabel: calculating labels...
ICLabel: saving results...
Computing projection and removing 7 components ....
pop_eegfiltnew() - performing 827 point bandpass filtering.
pop_eegfiltnew() - transition band width: 1 Hz
pop_eegfiltnew() - passband edge(s): [1 100] Hz
pop_eegfiltnew() - cutoff frequency(ies) (-6 dB): [0.5 100.5] Hz
pop_eegfiltnew() - filtering the data (zero-phase, non-causal)
firfilt(): |=====| 100%, ETE 00:00
Re-referencing data
Scaling components to RMS microvolt
Re-referencing ICA matrix
Scaling components to RMS microvolt
Warning: converting input data into regular (double) precision.
ICLabel: extracting features...
Scaling components to RMS microvolt
Recomputing ICA activations
ICLabel: calculating labels...
ICLabel: saving results...
Computing projection and removing 6 components ....
pop_eegfiltnew() - performing 827 point bandpass filtering.
pop_eegfiltnew() - transition band width: 1 Hz
pop_eegfiltnew() - passband edge(s): [1 100] Hz
pop_eegfiltnew() - cutoff frequency(ies) (-6 dB): [0.5 100.5] Hz
pop_eegfiltnew() - filtering the data (zero-phase, non-causal)
firfilt(): |=====| 100%, ETE 00:00
Re-referencing data
Scaling components to RMS microvolt
Re-referencing ICA matrix
Scaling components to RMS microvolt
Warning: converting input data into regular (double) precision.
ICLabel: extracting features...
Scaling components to RMS microvolt
Recomputing ICA activations
ICLabel: calculating labels...
ICLabel: saving results...
Computing projection and removing 7 components ....
pop_eegfiltnew() - performing 827 point bandpass filtering.
pop_eegfiltnew() - transition band width: 1 Hz
pop_eegfiltnew() - passband edge(s): [1 100] Hz
pop_eegfiltnew() - cutoff frequency(ies) (-6 dB): [0.5 100.5] Hz
pop_eegfiltnew() - filtering the data (zero-phase, non-causal)
firfilt(): |=====| 100%, ETE 00:00
Re-referencing data
Scaling components to RMS microvolt
Re-referencing ICA matrix
Scaling components to RMS microvolt
Warning: converting input data into regular (double) precision.
ICLabel: extracting features...
Scaling components to RMS microvolt
Recomputing ICA activations
ICLabel: calculating labels...
ICLabel: saving results...
Computing projection and removing 4 components ....
EEG successfully exported!

```

## 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";
```

## Prediction appears below the following code block.

For easier visibility, you can hide the code by going to View > Hide Code

```
[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")
    load("AllFeatures.mat","r_logical")
    F = table(F.Features(1,r_logical),'VariableNames',{'Features'});
    [yfit,score] = trainedModel.predictFcn(F);
    supportPredict(yfit);
case "ReliefF"
    load("ReliefF_SVM.mat")
    [yfit,score] = trainedModel1.predictFcn(F);
    supportPredict(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);
    supportPredict(yfit,score);
case "TMN"
    load("Topo_BasicNet.mat","net3")
    [yfit,score] =
classify(net3,y,"ExecutionEnvironment",'cpu',MiniBatchSize=1);
    supportPredict(yfit,score);
case "STFT"
    load("STFTNet.mat","net4a")
    [yfit,score] =
classify(net4a,y,"ExecutionEnvironment",'cpu',MiniBatchSize=1);
    supportPredict(yfit,score);
case "Fusion"
    load("SFnet.mat","bnet2")
    warning('off','MATLAB:mir_warning_unrecognized_pragma')
    [yfit,score] =
classify(bnet2,y,"ExecutionEnvironment",'cpu',MiniBatchSize=1);
    supportPredict(yfit,score);
end

```

EEG prediction: Normal  
 Prediction score: 0.72235  
 Score shown is maximum of the three classes and is between 0 and 1.

**CAUTION: EEG classification is set using default thresholding and should be further optimized for sensitivity/specificity/accuracy on additional training data prior to use. This code is not intended to make clinical diagnoses or to be used in any way to diagnose or treat subjects for whom the EEG is taken.**

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## 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"))
    end
end
```

```

        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
    rawEEG=[];
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 [cldata]=filtEEG(EEG,data)
for i=1:size(data,3)
    rawEEG=EEG;
    rawEEG.data=data(:,:,i);
end

```

```

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.gcompreject(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
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
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

```

```

function supportPredict(yfit,score)
if nargin < 2
    score = [];
end
switch yfit
    case "HEA"
        txt = "Normal";
    case "TBI"
        txt = "TBI";
    case "STR"
        txt = "Stroke";
end
disp("EEG prediction: " + string(txt));
if ~isempty(score)
    disp("Prediction score: " + num2str(max(score)));
    disp("Score shown is maximum of the three classes and is between 0 and 1.")
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
Dk1 = zeros(1,18);
for k = 1:18
    Dk1(k) = log(Pj(k)/(1/18))*Pj(k);
end
PAC = sum(Dk1)/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, 0, 0, 1, 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, 2, 3, 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, 4,...
            5, 5, 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, 6, 7, 7, 7, 7, 7, 7, 7,...
            7, 7, 7, 7, 8, 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
```