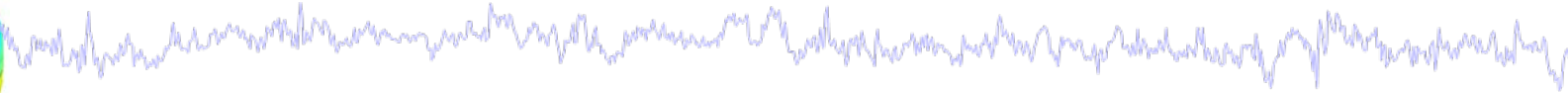


STUDY analysis



Task 1

Search STUDY structure

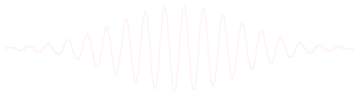
Task 2

Load/plot/use STUDY ERSP data

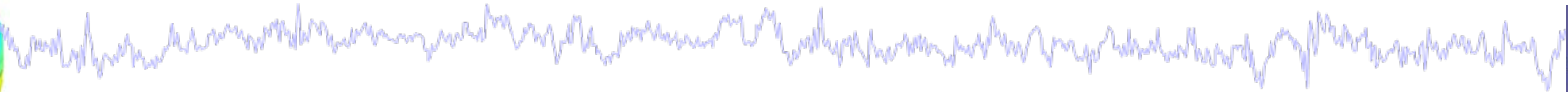
Task 3

Cluster ERP analysis

Exercise...



STUDY analysis



Task 1

Search STUDY structure

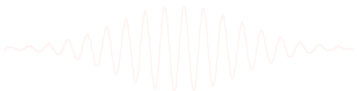
Task 2

Load/plot/use STUDY ERSP data

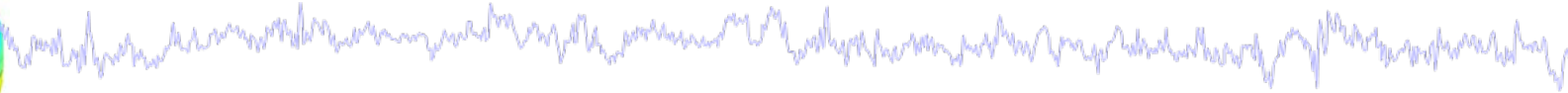
Task 3

Cluster ERP analysis

Exercise...

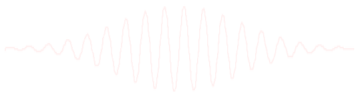


STUDY structure details



Question:

**How are IC and subject information
stored in the **STUDY** structure?**



Understanding STUDY structure



26 = # of clusters

One cluster:



```
>> STUDY.cluster
```

```
1x26 struct array with fields:
```

```
parent
name
child
comps
sets
algorithm
centroid
preclust
dipole
selected
allinds
setinds
topo
topox
topoy
topoall
topopol
```

```
>> STUDY.cluster(6)
```

6 = cluster number

```
ans =
```

```
parent: {'ParentCluster 1'}
```

```
name: 'Cls 6'
```

```
child: []
```

3 = # conditions

IC indices

```
comps: [35 7 12 35 10 23 7 30 4 15]
```

```
sets: [3x10 double]
```

dataset indices

```
algorithm: {'Kmeans' [24]}
```

```
centroid: []
```

```
preclust: [1x1 struct]
```

```
dipole: [1x1 struct]
```

```
selected: 1
```

```
allinds: []
```

67 = # of channels

```
setinds: []
```

```
topo: [67x67 double]
```

```
topox: [67x1 double]
```

```
topoy: [67x1 double]
```

```
topoall: {1x10 cell}
```

```
topopol: [1 1 1 -1 1 1 1 1 1 1]
```

scalp map polarity

Understanding STUDY structure



```
>> STUDY.cluster(6)
ans =
```

```
parent: {'ParentCluster 1'}
name: 'Cls 6'
child: []
comps: [35 7 12 35 10 23 7 30 4 15]
sets: [3x10 double]
algorithm: {'Kmeans' [24]}
centroid: []
preclust: [1x1 struct]
dipole: [1x1 struct]
selected: 1
allinds: []
setinds: []
topo: [67x67 double]
topox: [67x1 double]
topoy: [67x1 double]
topoall: {1x10 cell}
topopol: [1 1 1 -1 1 1 1 1 1 1]
```

Which subject?

Which dataset(s)?

Dataset indices

```
>> STUDY.cluster(6).sets
```

```
ans = Second IC = second column
```

| | | | |
|---|----|----|----------------|
| 2 | 11 | 14 | ...Condition 1 |
| 1 | 10 | 13 | ... |
| 3 | 12 | 15 | ... |

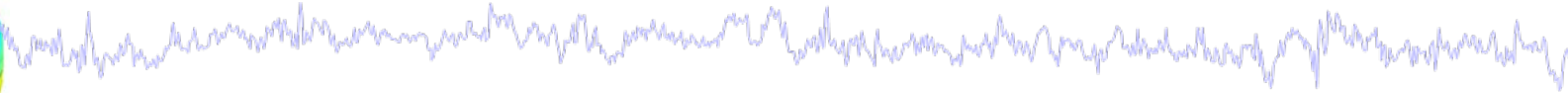
```
>> STUDY.datasetinfo(11) % access dataset 11
ans =
```

```
filepath: [1x61 char]
filename: 'Ignore.set'
subject: 'S04'
session: []
condition: 'ignore'
group: ''
index: 11
comps: [1x24 double]
```

Subject 4!

Condition 1

STUDY analysis



Task 1

Search STUDY structure

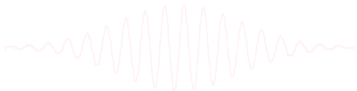
Task 2

Load/plot/use STUDY ERSP data

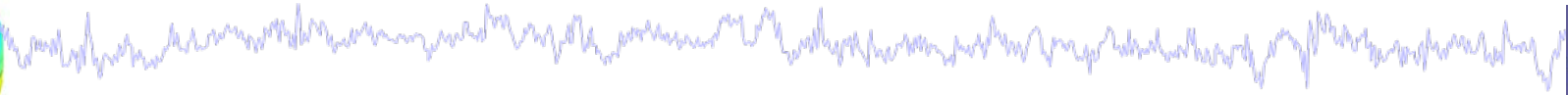
Task 3

Cluster ERP analysis

Exercise...



Load data from commandline



**** Where is the raw data stored?**

Data for each subject is stored in the file path
of that subject (`STUDY.datasetinfo(subj).filepath`)

**** What is it called?**

File name format: `'setname.extension'`

extension = `'.ica*' or '.dat*' (for channel data)`

for example:

`Memorize.icaerp % ERP data`

`Memorize.icaersp % ERSP data`

`Memorize.icaitc % ITC data`

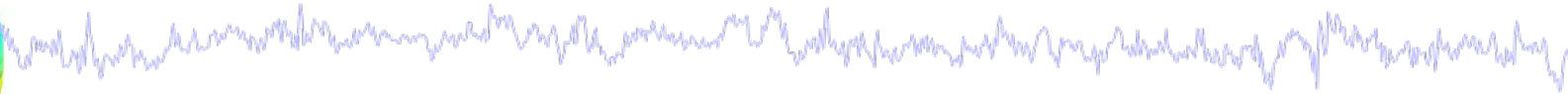
`Memorize.icaspec % Power spectrum data`

`Memorize.icatopo % Scalp map data`

% Example of channel data file name:

`Memorize.daterp % ERP data`

Load individual ERSPs



```
% load ERSP data for all ICs in a single cluster:
```

```
clust = 5; % choose a cluster
```

```
cond = 1; % choose experimental condition
```

```
[STUDY logersp timevals logfreqs pgroup pcond pinter] = ...
```

```
std_erspplot(STUDY,ALLEEG,'clusters',clust, 'plotsubjects', 'off' );
```

```
% Check imported variables in workspace:
```

```
>> whos logersp
```

| Name | Size | Bytes |
|---------|------------|--------------|
| Class | Attributes | |
| logersp | 3x1 | 1983216 cell |

```
>> logersp
```

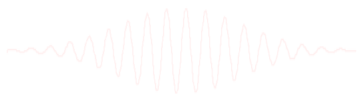
```
logersp =
```

```
[72x153x15 single]
```

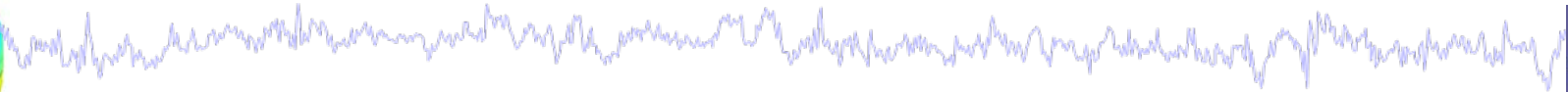
```
[72x153x15 single]
```

```
[72x153x15 single]
```

frequency x times x ICs



Load individual ERSPs



% Check imported variables in workspace:

```
>> whos logersp
```

| Name | Size | Bytes |
|---------|------------|--------------|
| Class | Attributes | |
| logersp | 3x1 | 1983216 cell |

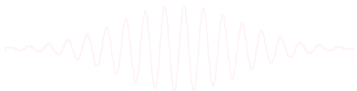
```
>> logersp
```

```
logersp = frequency x times x ICs
```

```
[72x153x15 single]
```

```
[72x153x15 single]
```

```
[72x153x15 single]
```



Plot individual ERSPs



```
% load ERSP data for all ICs in a single cluster:
```

```
clust = 5; % choose a cluster
```

```
cond = 1; % choose experimental condition
```

```
[STUDY logersp timevals logfreqs pgroup pcond pinter] = ...
```

```
std_erspplot(STUDY,ALLEEG,'clusters',clust, 'plotsubjects', 'on' );
```

```
% or plot them yourself from output:
```

```
figure; clim = 3; % standardize color limits
```

```
for ic = 1:size(logersp{cond},3)
```

```
    subplot(row,col,ic);
```


```
    imagesclogy(timevals, logfreqs, logersp{cond}(:, :, ic));
```

```
    set(gca,'clim', [-clim clim]); % adjust the color limits
```

```
    set(gca,'ydir','norm'); % plot low freqs at the bottom
```

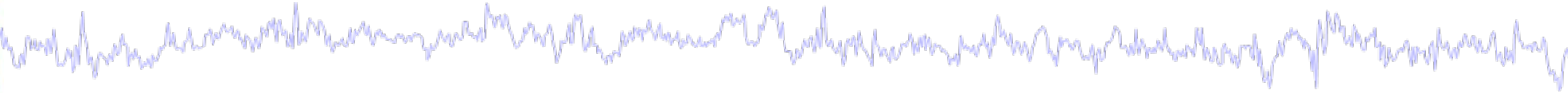
```
    title(['IC ',int2str(STUDY.cluster(clust).comps(ic))]);
```

```
end;
```

```
 textsc(['Cluster ',int2str(clust)], 'title');
```

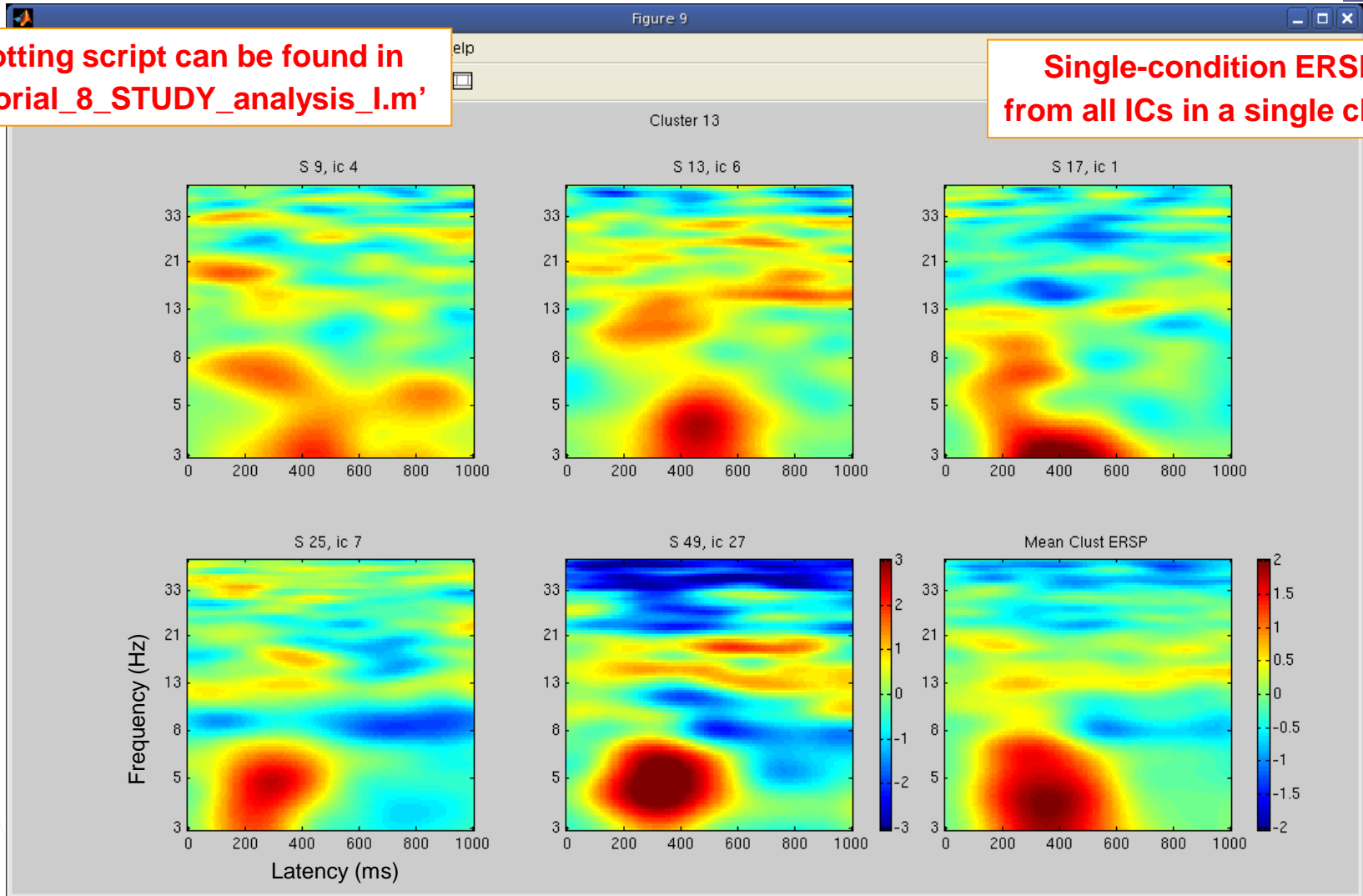
```
cbar; % include a colorbar
```

PLOT individual ERSPs

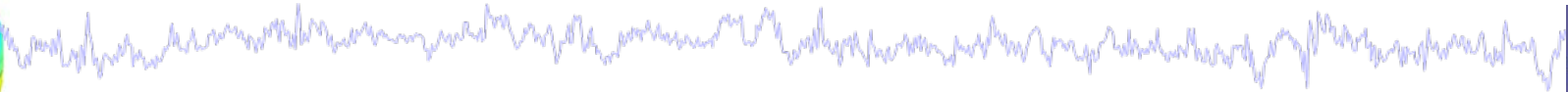


Plotting script can be found in
'Tutorial_8_STUDY_analysis_1.m'

Single-condition ERSPs
from all ICs in a single cluster



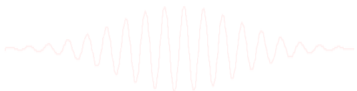
Raw data files



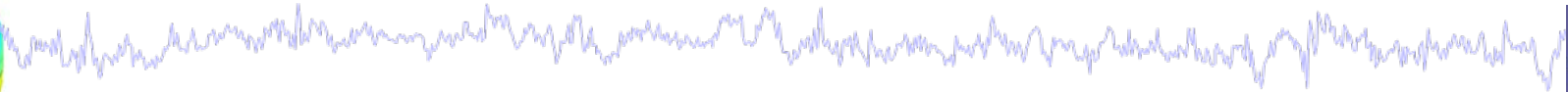
```
% Load *raw* ERSP data
```

```
load_string = 'C:\...\workshop\STUDY\S01\Memorize.icaersp';
```

```
ERSPdata = load('-mat',load_string); % .mat format!
```



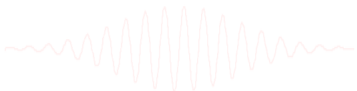
Raw data structure



```
>> ERSPdata
```

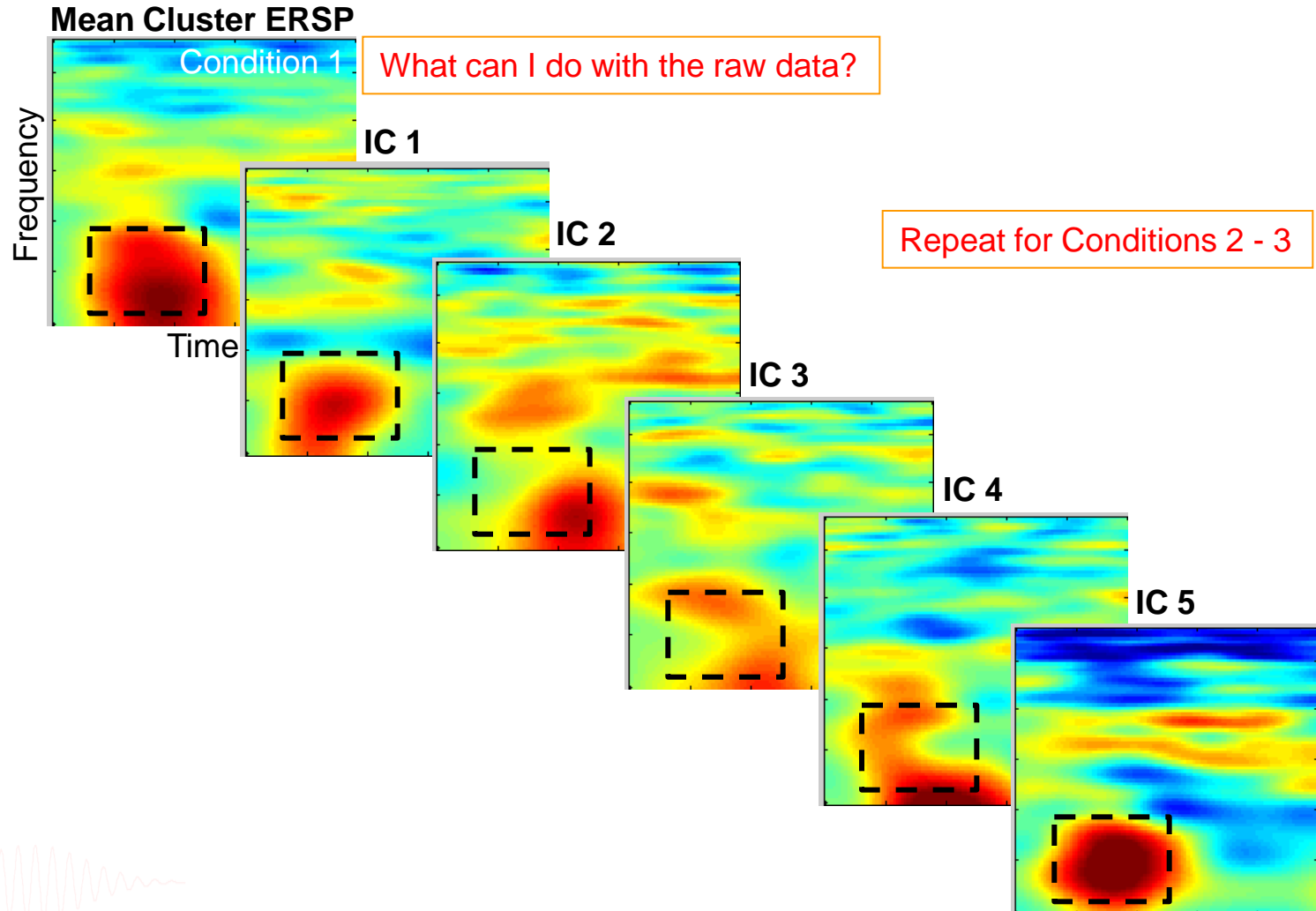
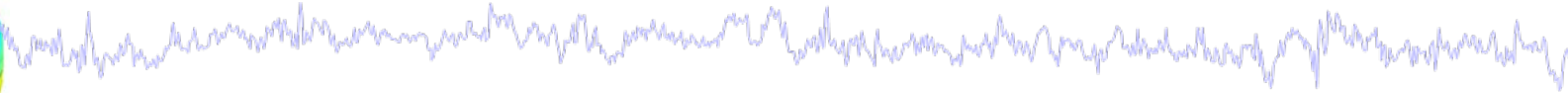
```
ERSP dB data → comp1_ersp: [100 x 200 single]
dB baseline → comp1_erspbase: [1 x 100 single] → 200 time points
bootstrap limits → comp1_erspboot: [100 x 2 single] → upper and lower bootstrap limits
                  comp2_ersp: [100 x 200 single]
                  comp2_erspbase: [1 x 100 single] → 100 frequency bins
                  comp2_erspboot: [100 x 2 single]
100 frequency bins → freqs: [1 x 100 double]
200 time points → times: [1 x 200 double]

      datatype: 'ERSP'
      parameters: {1 x 26 cell}
      datafile: [1 x 57 char]
```

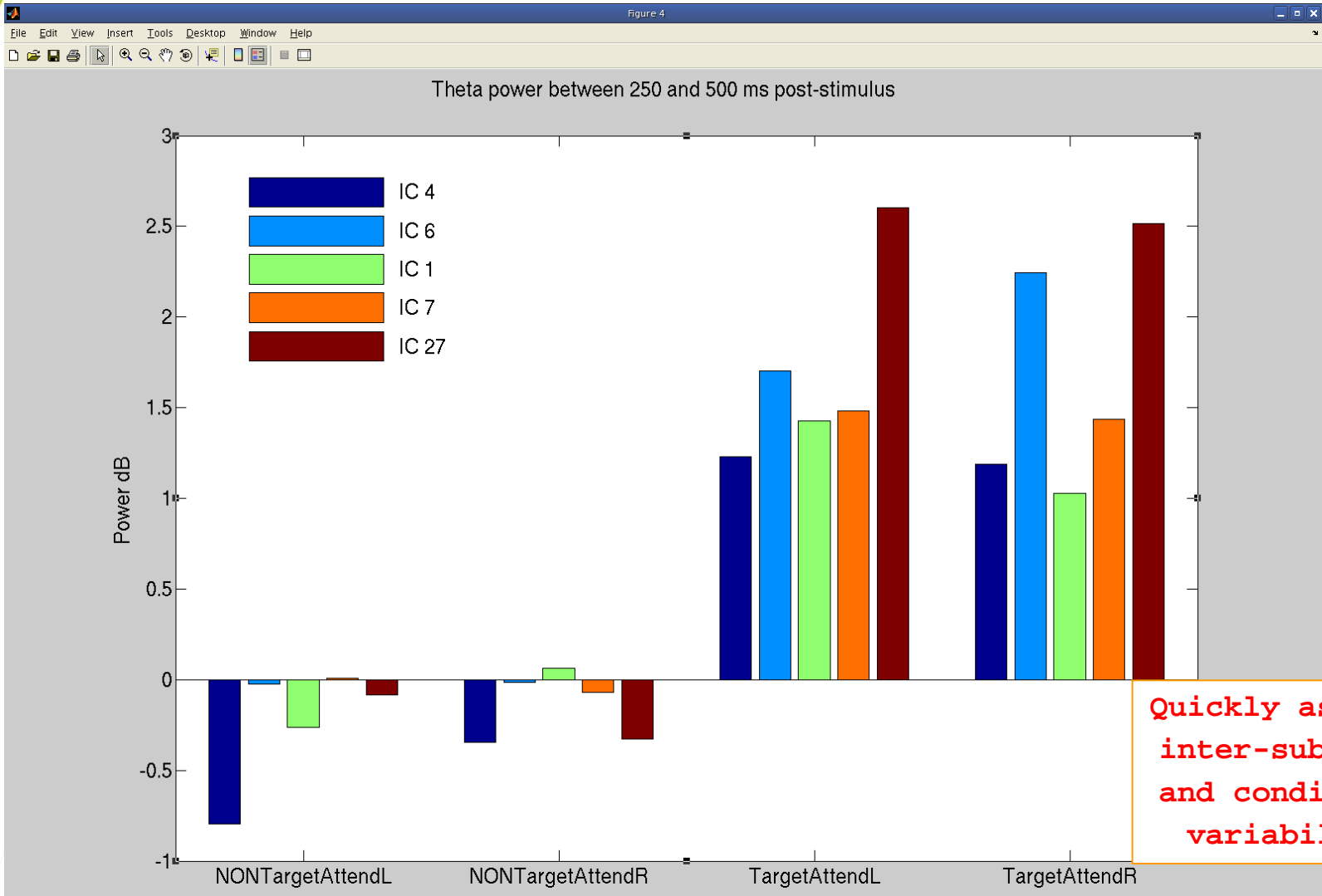
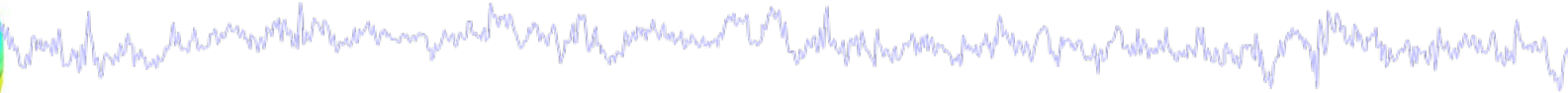


```
>>
```

Use STUDY ERSP data for analysis

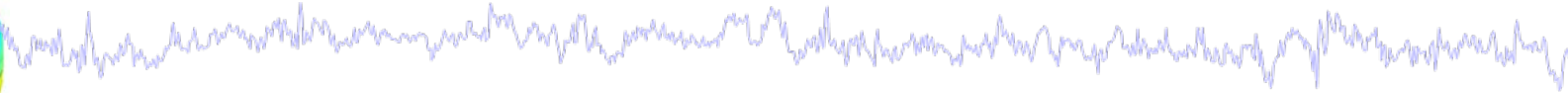


Mean theta power across conditions



Quickly assess inter-subject and condition variability

STUDY analysis



Task 1

Search STUDY structure

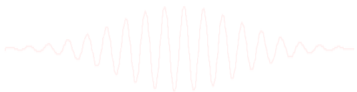
Task 2

Load/plot/use STUDY ERSP data

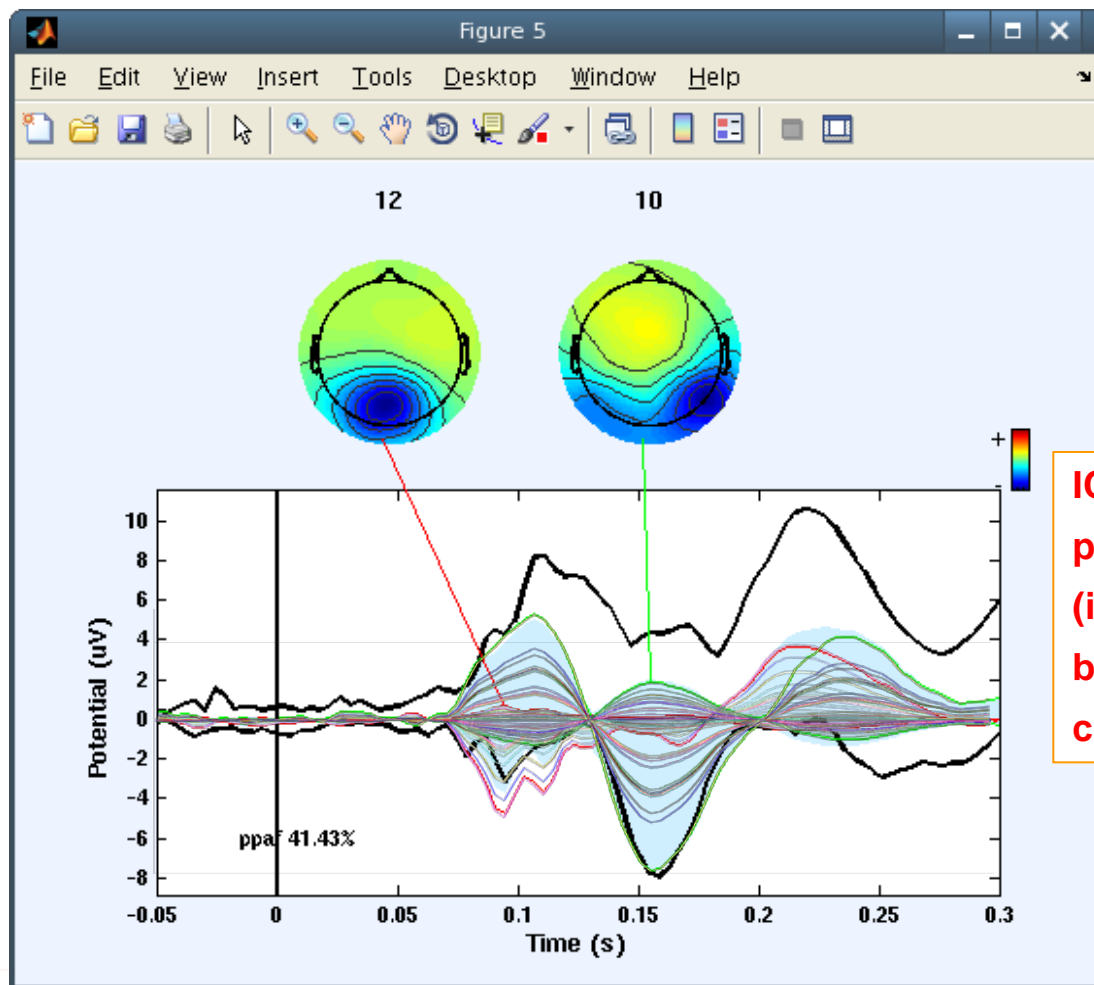
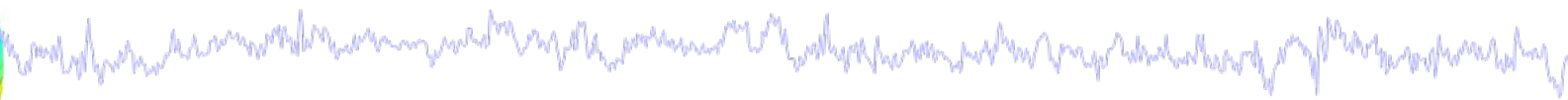
Task 3

Cluster ERP analysis

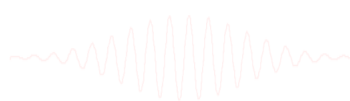
Exercise...



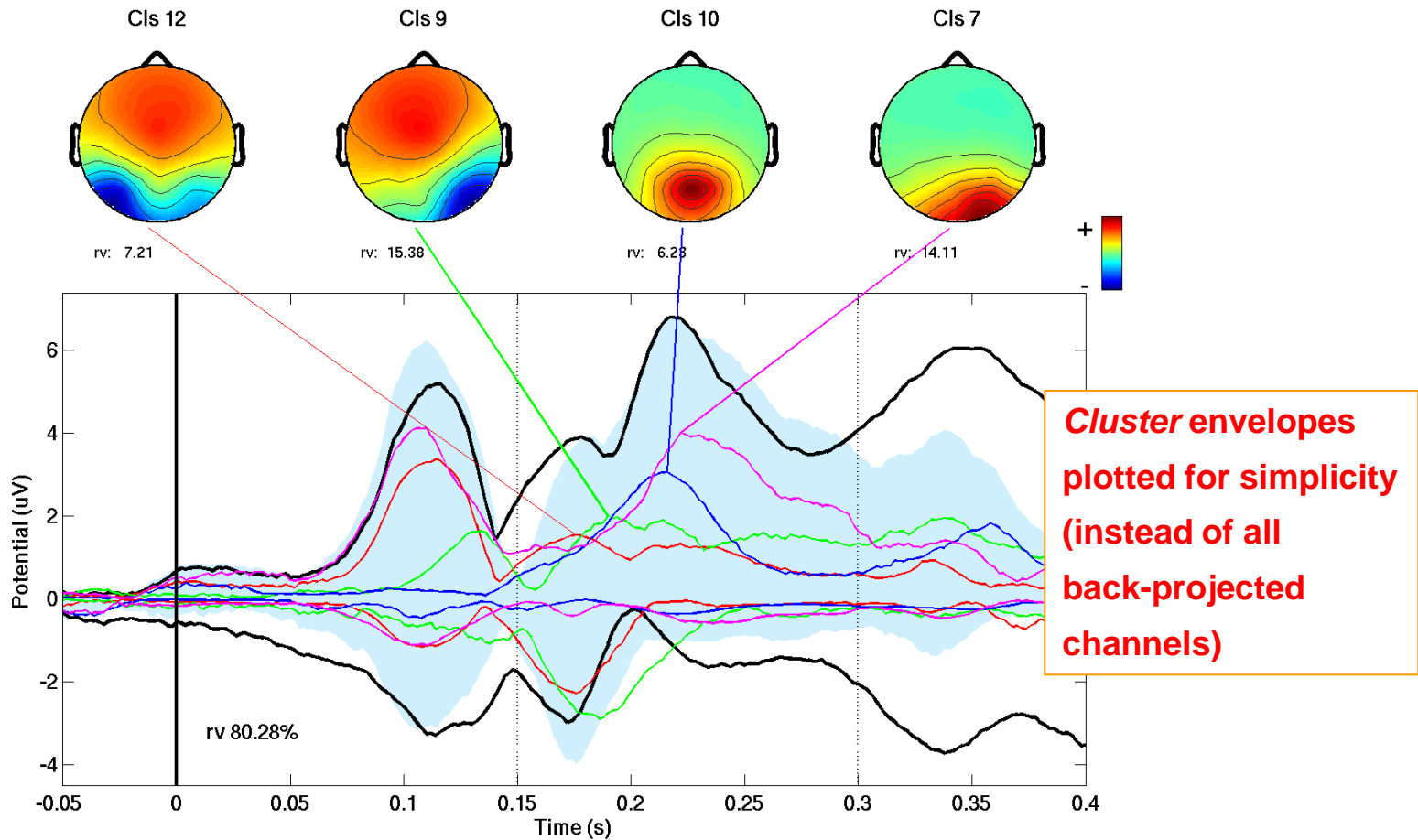
REVIEW: Single-subject IC ERP envelope



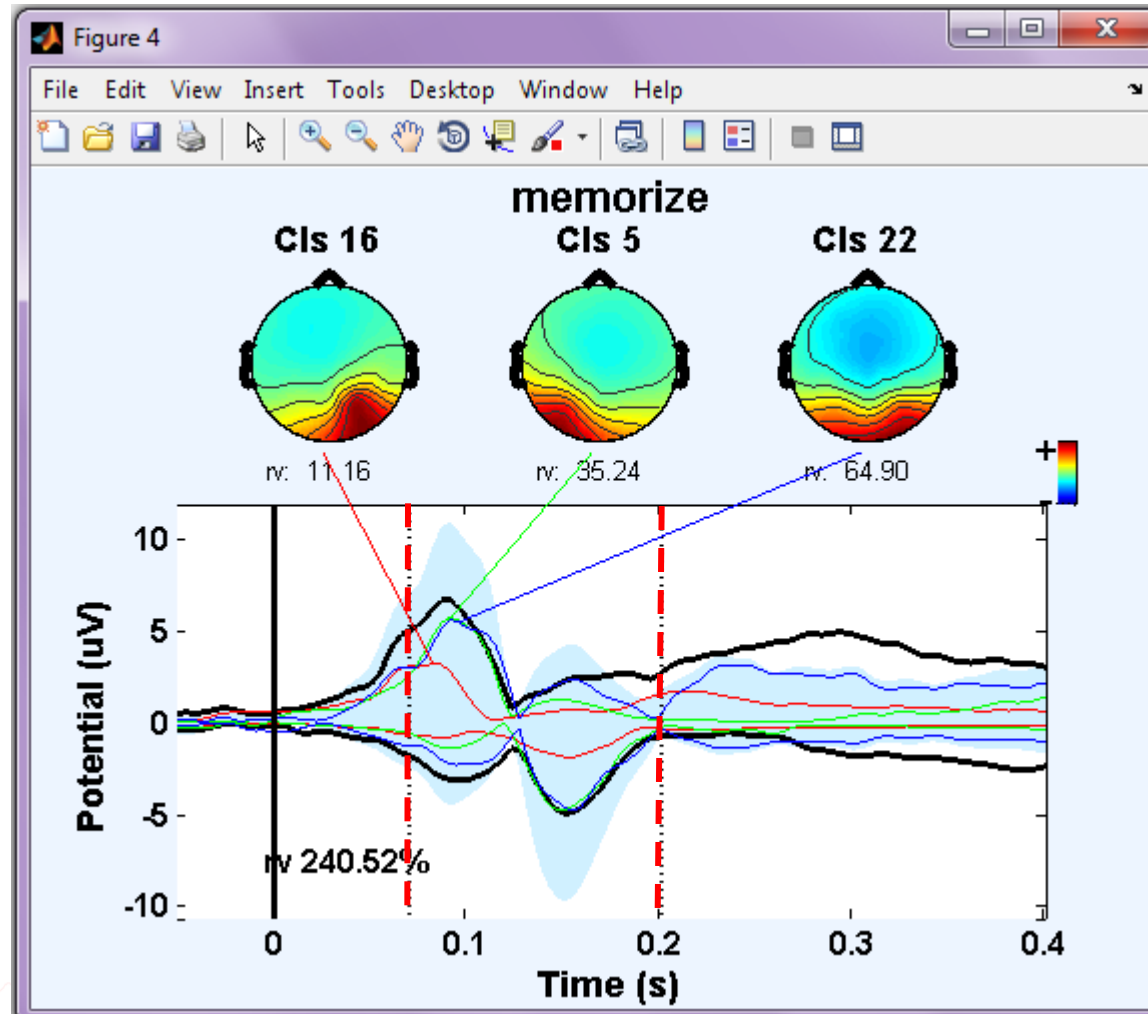
IC envelopes plotted for simplicity (instead of all back-projected channels)



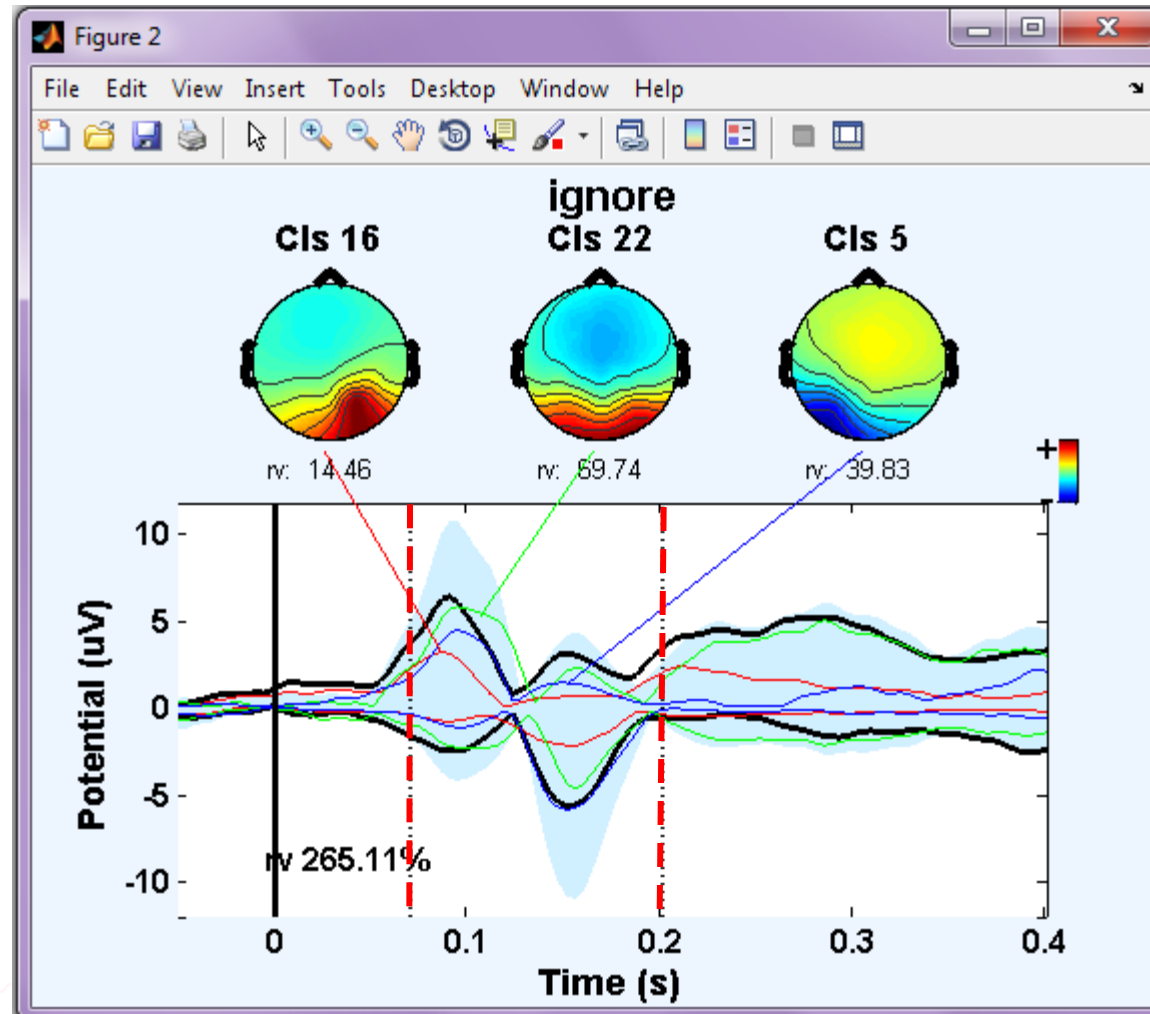
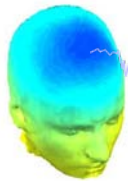
STUDY Cluster ERP analysis



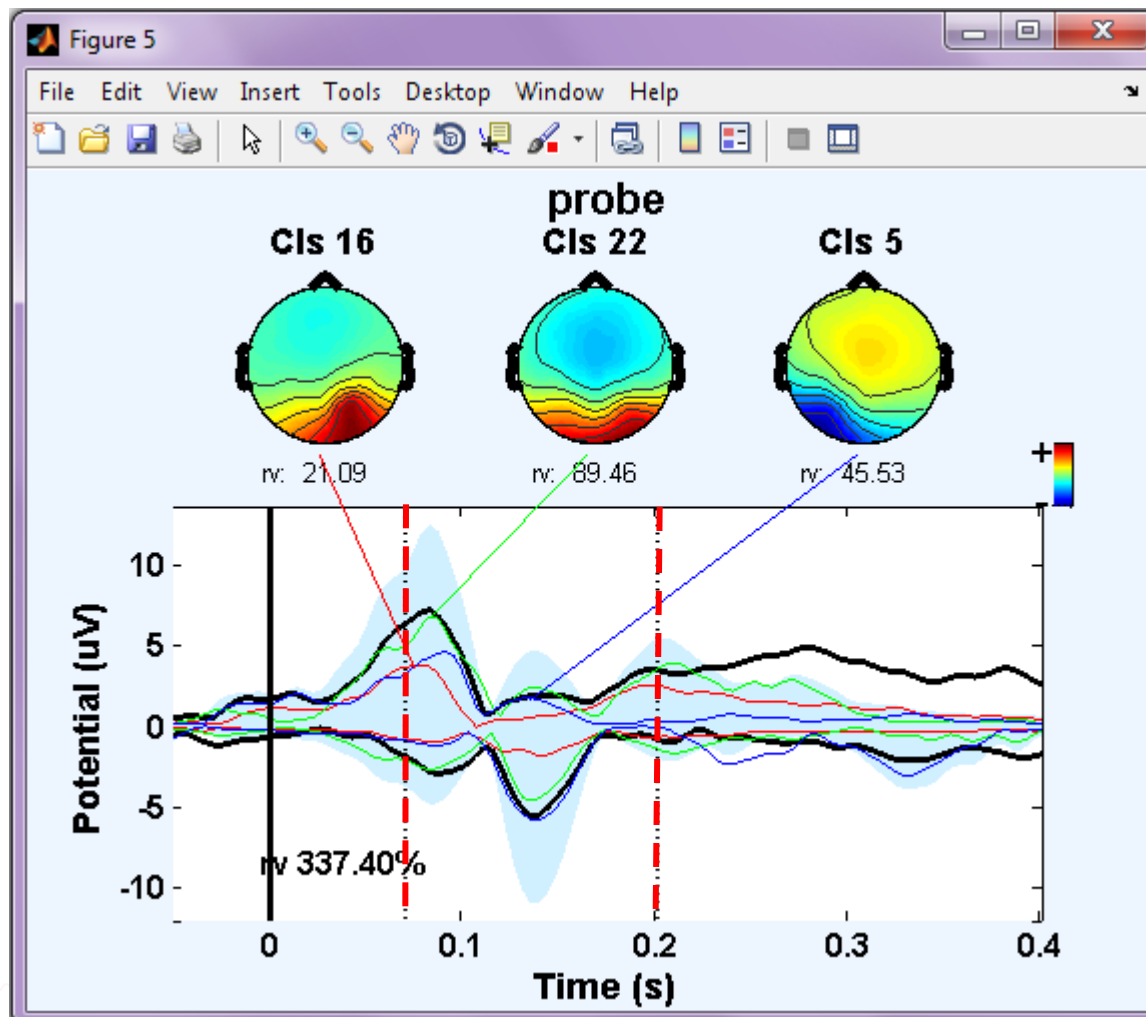
Largest early ERP contributors (Memorize)



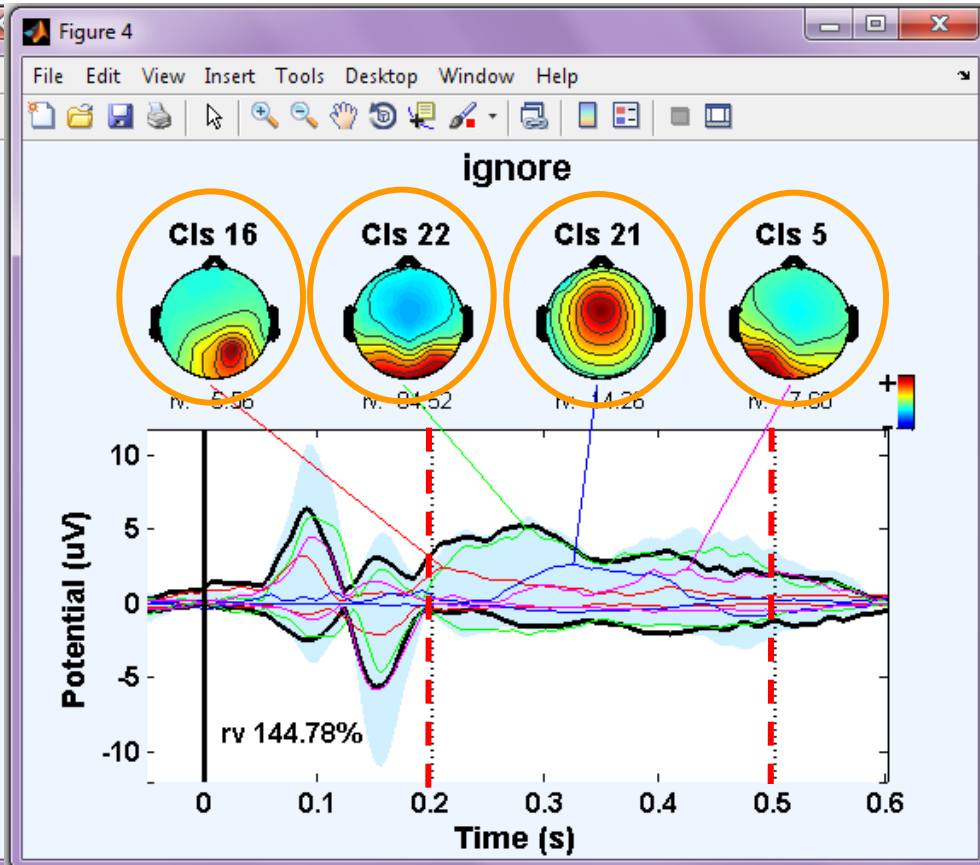
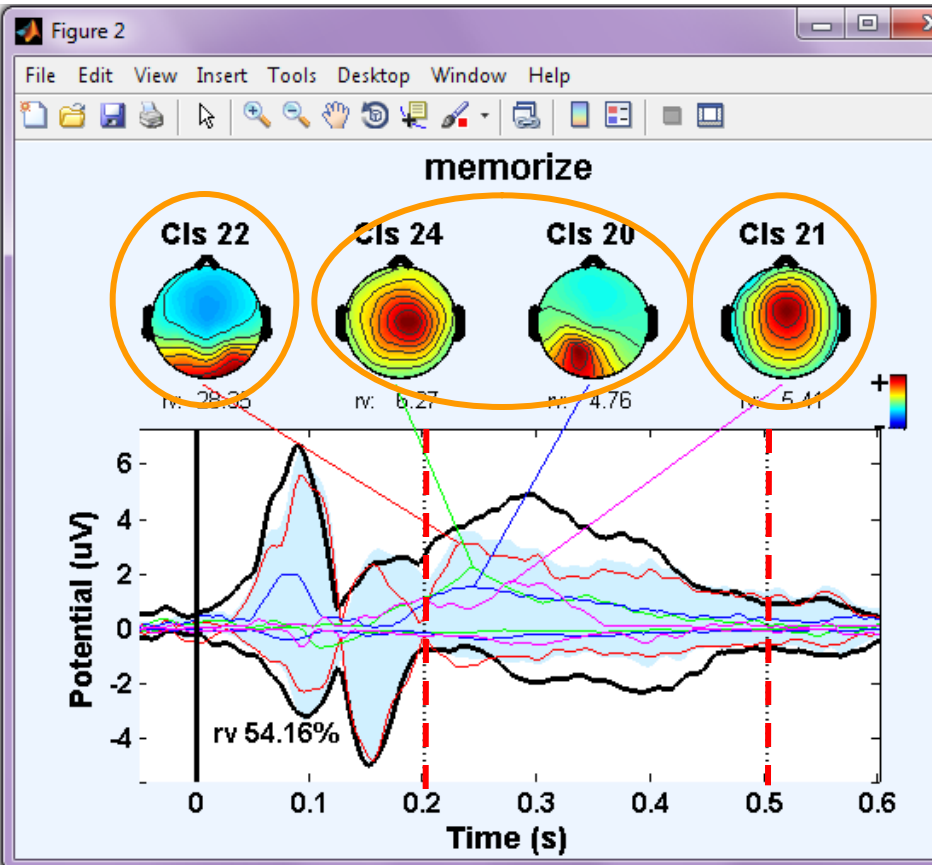
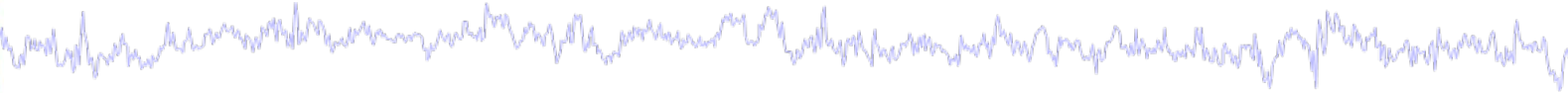
Largest early ERP contributors (Ignore)



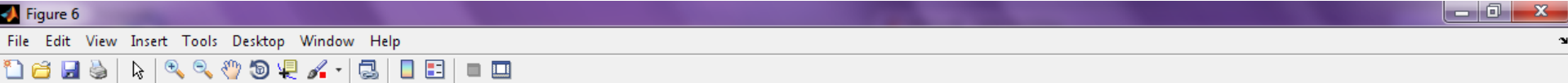
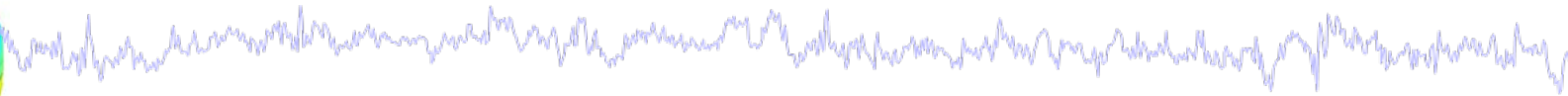
Largest early ERP contributors (Probe)



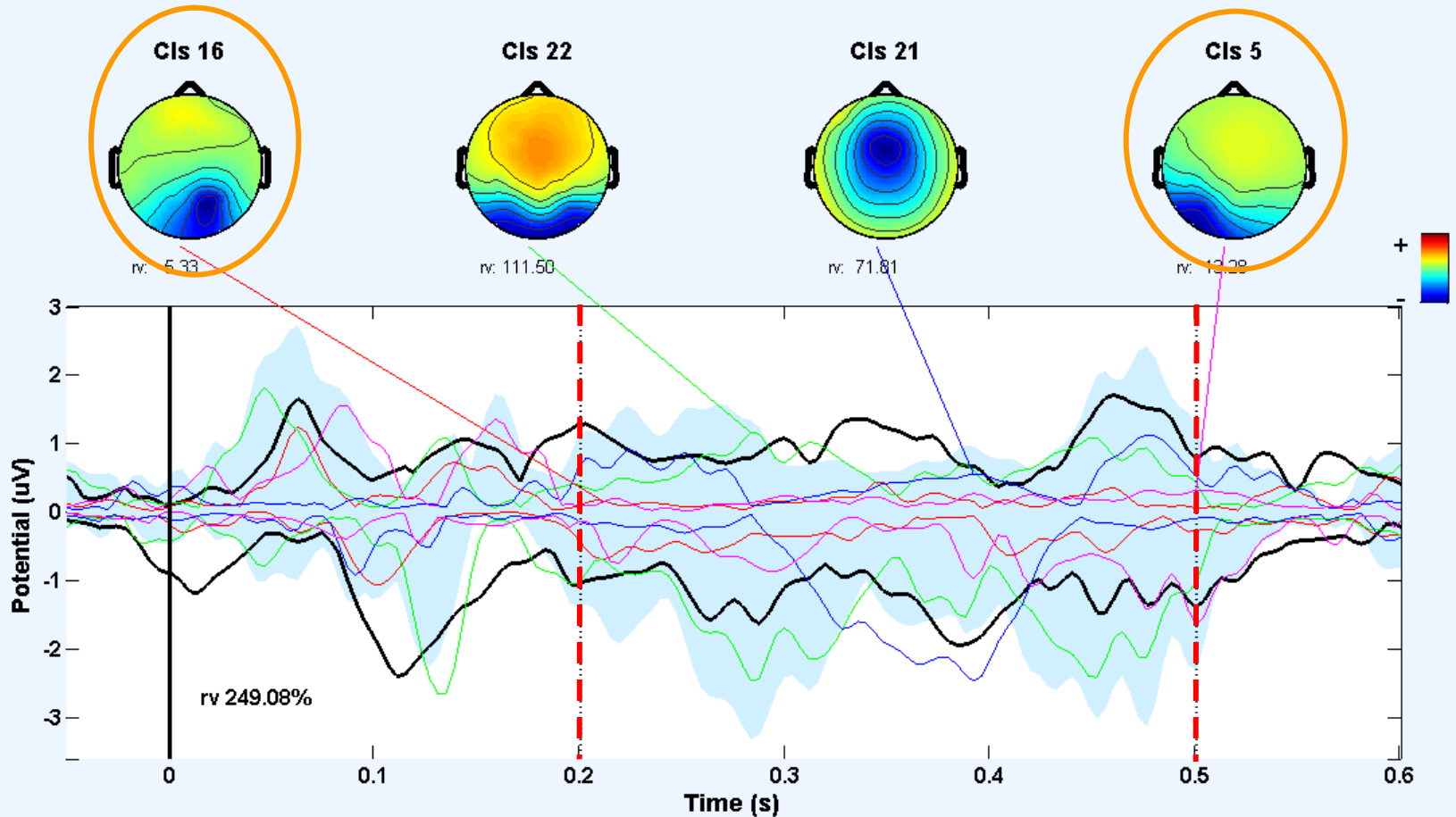
Memorize and Ignore ERP envelopes



Memorize-Ignore ERP envelopes



Difference Between Conditions 2 and 1.



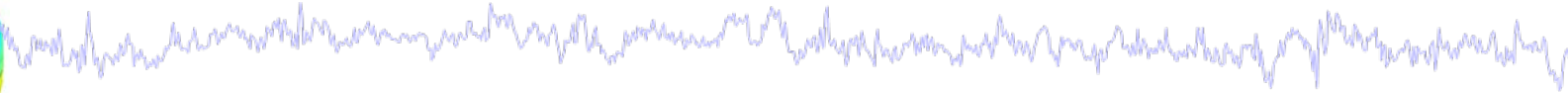
Exercise



- **ALL**
 - Load workshop STUDY
- **Novice**
 - Load and plot individual ERSPs for one or more clusters.
 - How consistent are the ERSPs in these clusters?
- **Intermediate**
 - Choose a cluster to investigate
 - Plot mean power in a small time/frequency window across all ICs and conditions for this cluster
- **Advanced**
 - Plot cluster ERP (std_envtopo) and compare with ERP image
(this function is still under construction and cannot be used at this time)

** All scripts for Intermediate/Advanced exercises can be found in
.../Scripts/Tutorial_8_STUDY_analysis_1.m

STUDY analysis



Task 1

Search STUDY structure

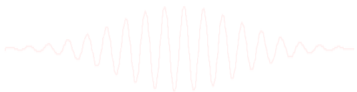
Task 2

Load/plot/use STUDY ERSP data

Task 3

Alternative: saving ICA weights
and applying to another .set

Exercise...



ICA weights are stored in EEG structure



```
Terminal
File Edit View Terminal Tabs Help
>> EEG

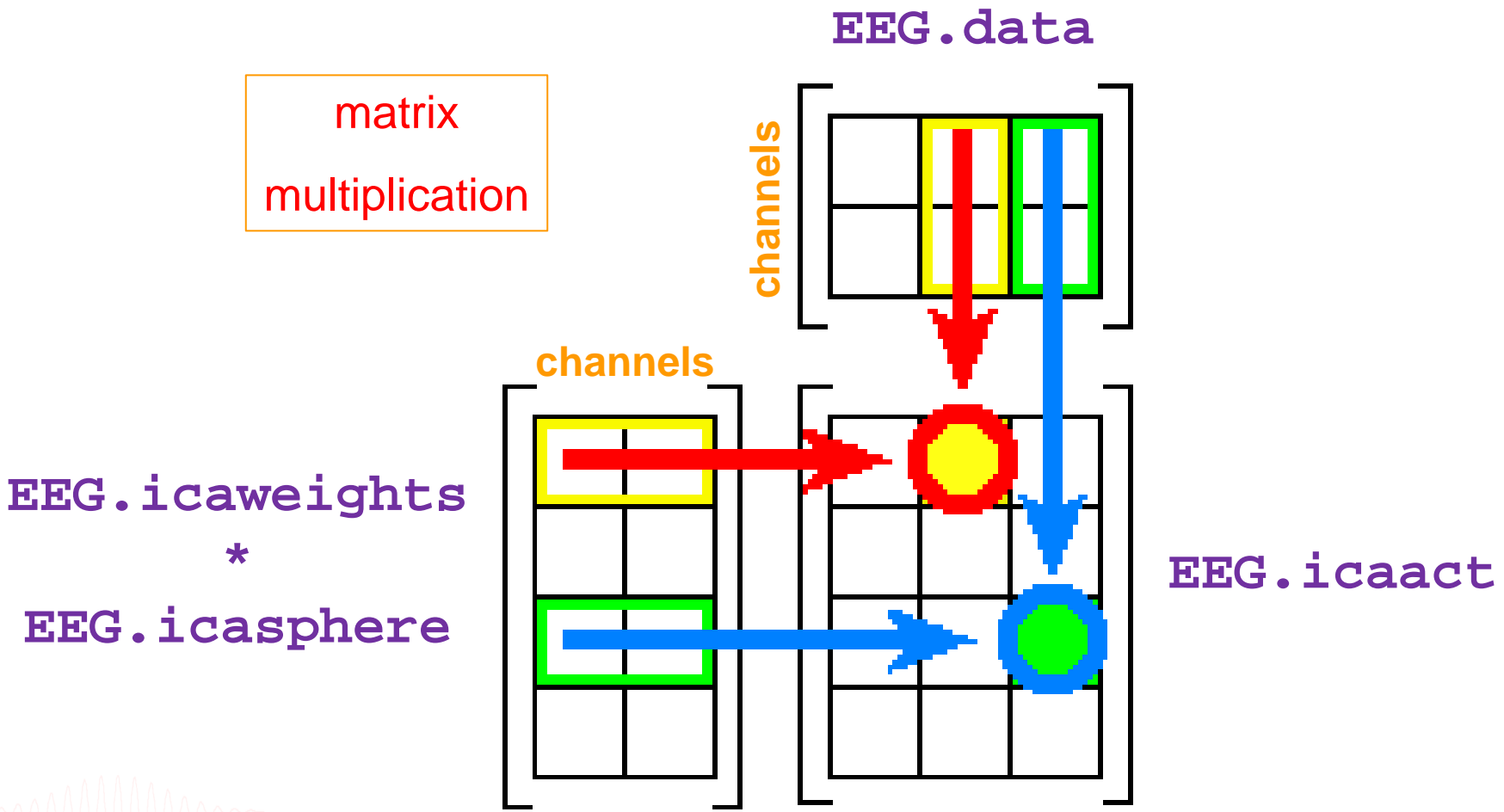
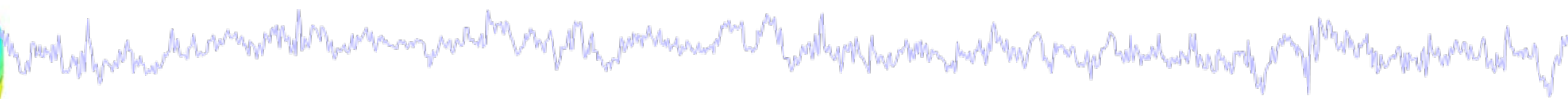
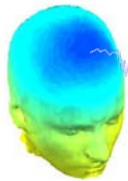
EEG =

    setname: 'faces_4 continuous'
    filename: 'faces_4.set'
    filepath: '/home/julie/workshop06/'
    subject: ''
    group: ''
    condition: ''
    session: []
    comments: [15x48 char]
    nbchan: 33
    trials: 1
        pnts: 133175
    srates: 250
    xmin: 0
    xmax: 532.6960
    times: []
    data: [33x133175 single]
    icaact: [33x133175 single]
    icawinv: [33x33 double]
    icasphere: [33x33 double]
    icaweights: [33x33 double]
    icachansind: [1x33 double]
    chanlocs: [1x33 struct]
    urchanlocs: []
```

ICA weights can be applied to other datasets acquired during the same session with the **same electrode placement**.

$$\text{EEG.icaact} = (\text{EEG.icaweights} * \text{EEG.icasphere}) * \text{EEG.data}$$

Applying ICA weights to EEG data



Saving/applying ICA weights



```
>> wts = EEG.icaweights;
```

```
>> sph = EEG.icasphere;
```

```
>> save subj1_ICA_Weights.mat wts sph
```

```
% close EEGLAB dataset and open another
```

```
% dataset from same subject and same day
```

```
% If you also closed matlab, load the .mat file:
```

```
>> load subj1_ICA_Weights.mat wts sph
```

```
>> EEG.icaweights = wts;
```

```
>> EEG.icasphere = sph;
```

```
>> % delete old activations and scalp maps, if present
```

```
>> EEG.icaact = []; EEG.icawinv = [];
```

```
% recalculate acts and winv
```

```
>> EEG.icaact = eeg_getdataact(EEG,...
```

```
    'component',[1:size(EEG.data,1)]);
```

