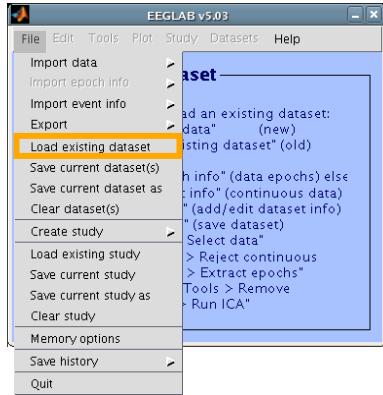


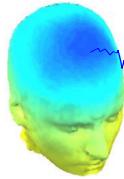
Command line tools



(Menus write both dataset and global history)

- Automated processing on groups of subjects (possibly on several processors).
- Richer options for plotting and processing functions (time-frequency decompositions, ...)
- Selecting data/epoch based on event context
- Custom processing...

Using EEGLAB history for basic scripting



Task 1

Create a script from ‘eegh’ output

Task 2

Adapt your script with variables

Task 3

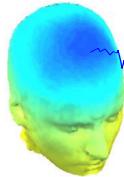
Create a Matlab function

Task 4

Exercise...



Using EEGLAB history for basic scripting



Task 1

Create a script from ‘eegh’ output

Task 2

Adapt your script with variables

Task 3

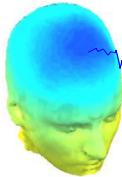
Create a Matlab function

Task 4

Exercise...



Memory options



EEGLAB

File Edit Tools Plot Study
Import data Import epoch info Import event info Export
Load existing dataset Save current dataset(s) Save current dataset as Clear dataset(s)
Create study Load existing study Save current study Save current study as Clear study
Memory and other options Save history ▾ Quit

Memory options - pop_editoptions()

STUDY options (set these checkboxes if you intend to work with studies)

If set, keep at most one dataset in memory. This allows processing hundreds of datasets within studies.
If set, save not one but two files for each dataset (header and data). This allows faster data loading in studies.
If set, write ICA activations to disk. This speeds up loading ICA components when dealing with studies.

Memory options

If set, use single precision under Matlab 7.x. This saves RAM but can lead to rare numerical imprecisions.
If set, use memory mapped array under Matlab 7.x. This may slow down some computation.

ICA options

If set, precompute ICA activations. This requires more RAM but allows faster plotting of component activations.
If set, scale ICA component activities to RMS (Root Mean Square) in microvolt (recommended).

Folder options

If set, when browsing to open a new dataset assume the folder/directory of previous dataset.

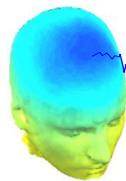
Option file: C:\Users\julie\Documents\MATLAB\functions\adminfunc\eeeg_options.m ...

Help Cancel Ok

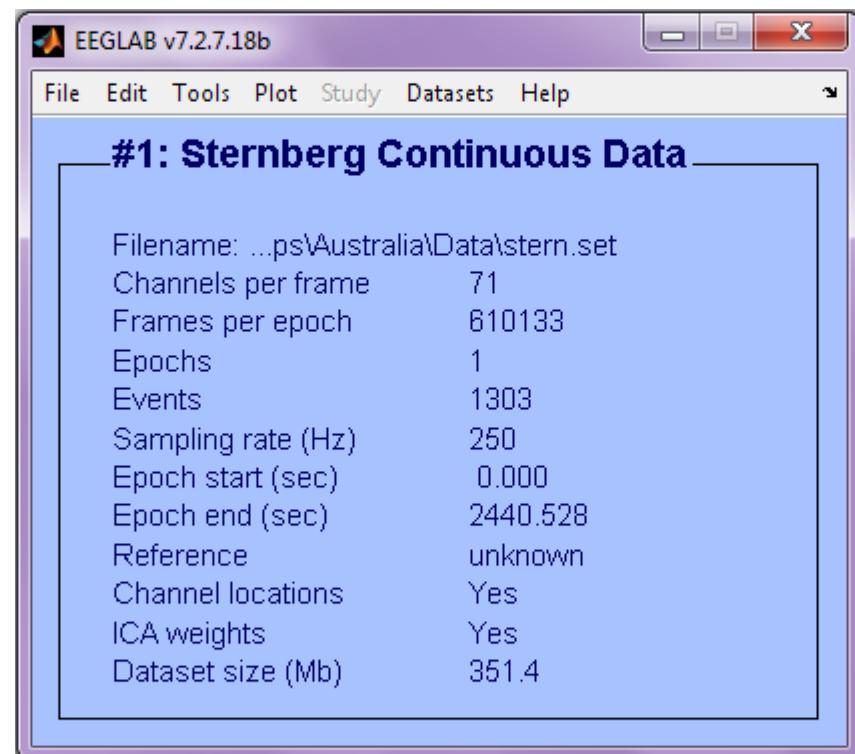
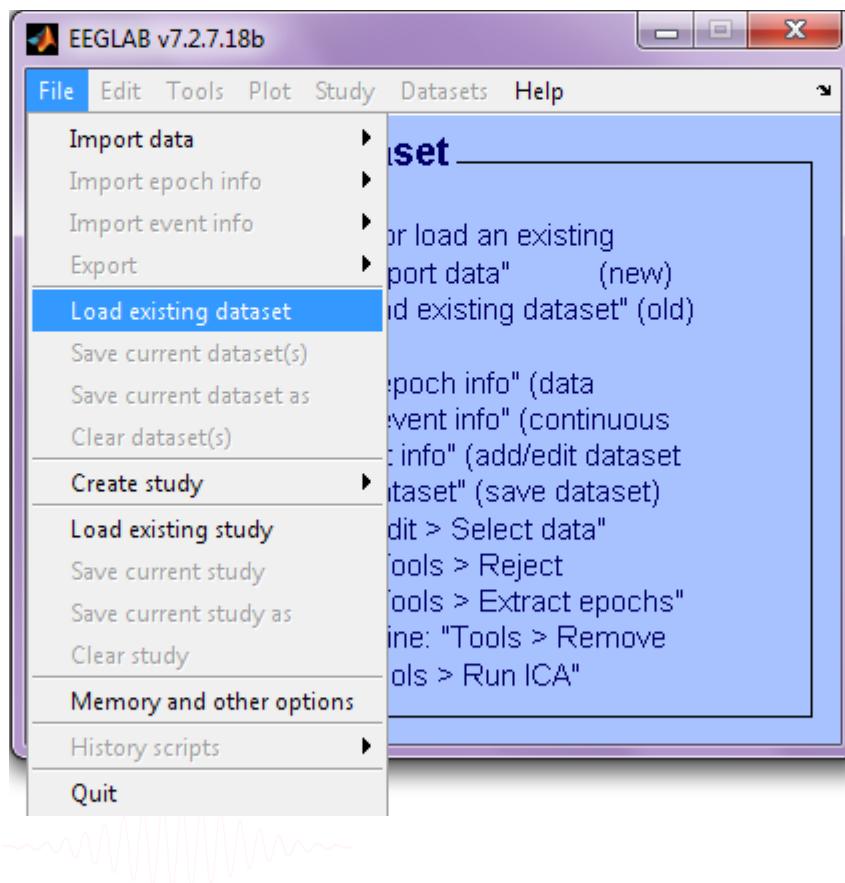
data
ect continuous
ract epochs"
> Remove
ICA"

**Change memory options
to allow more than one dataset in memory**

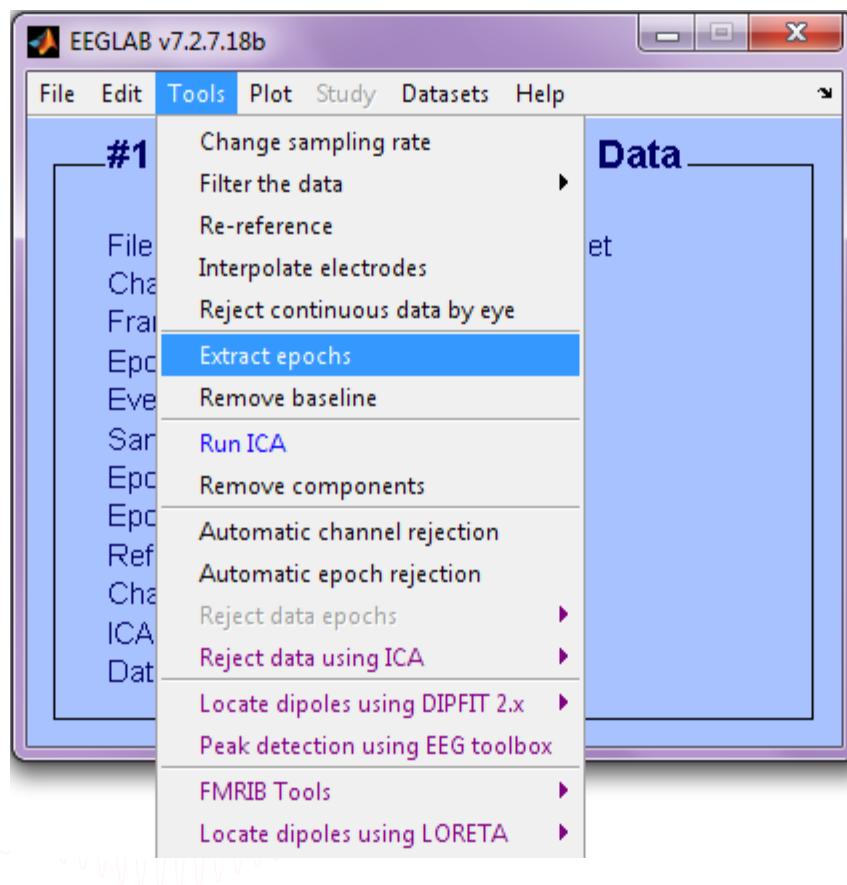
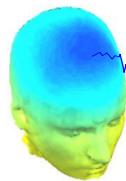
Create a script from ‘eegh’ output



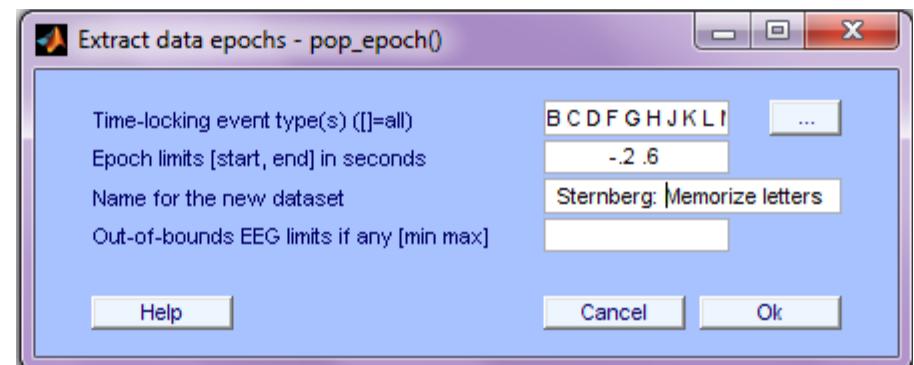
Start by loading a continuous dataset



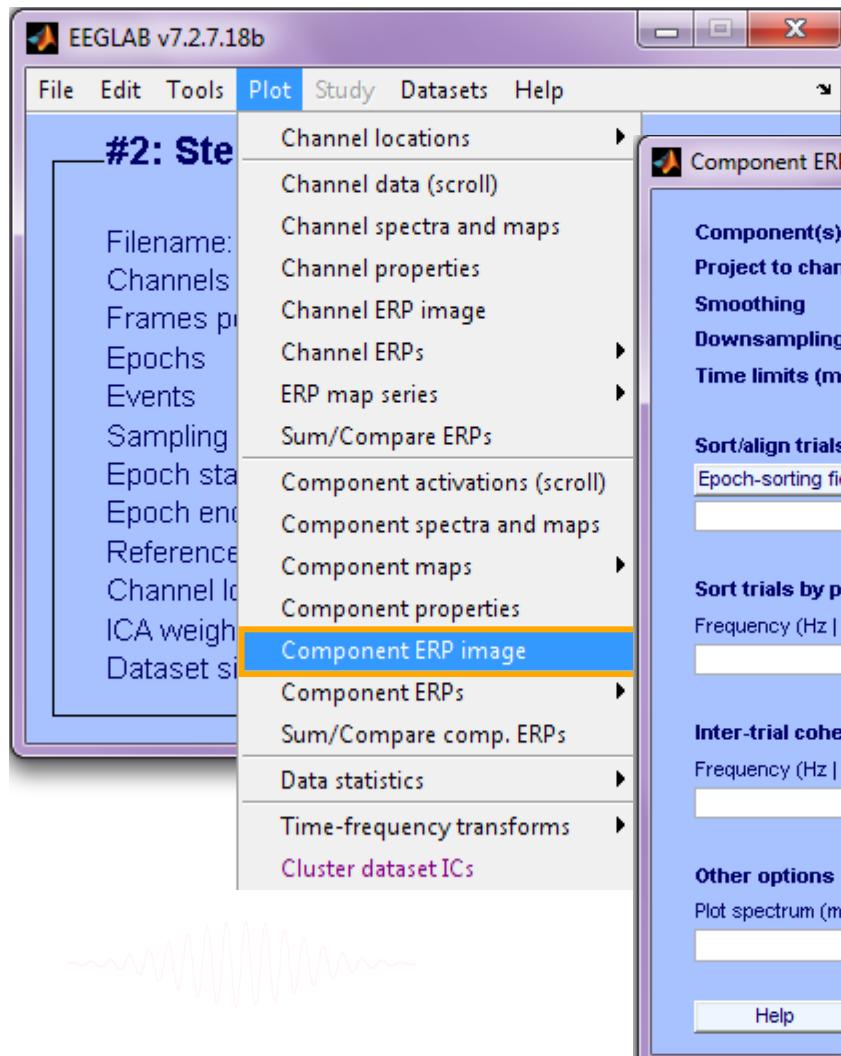
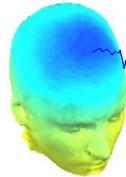
Create a script from ‘eegh’ output



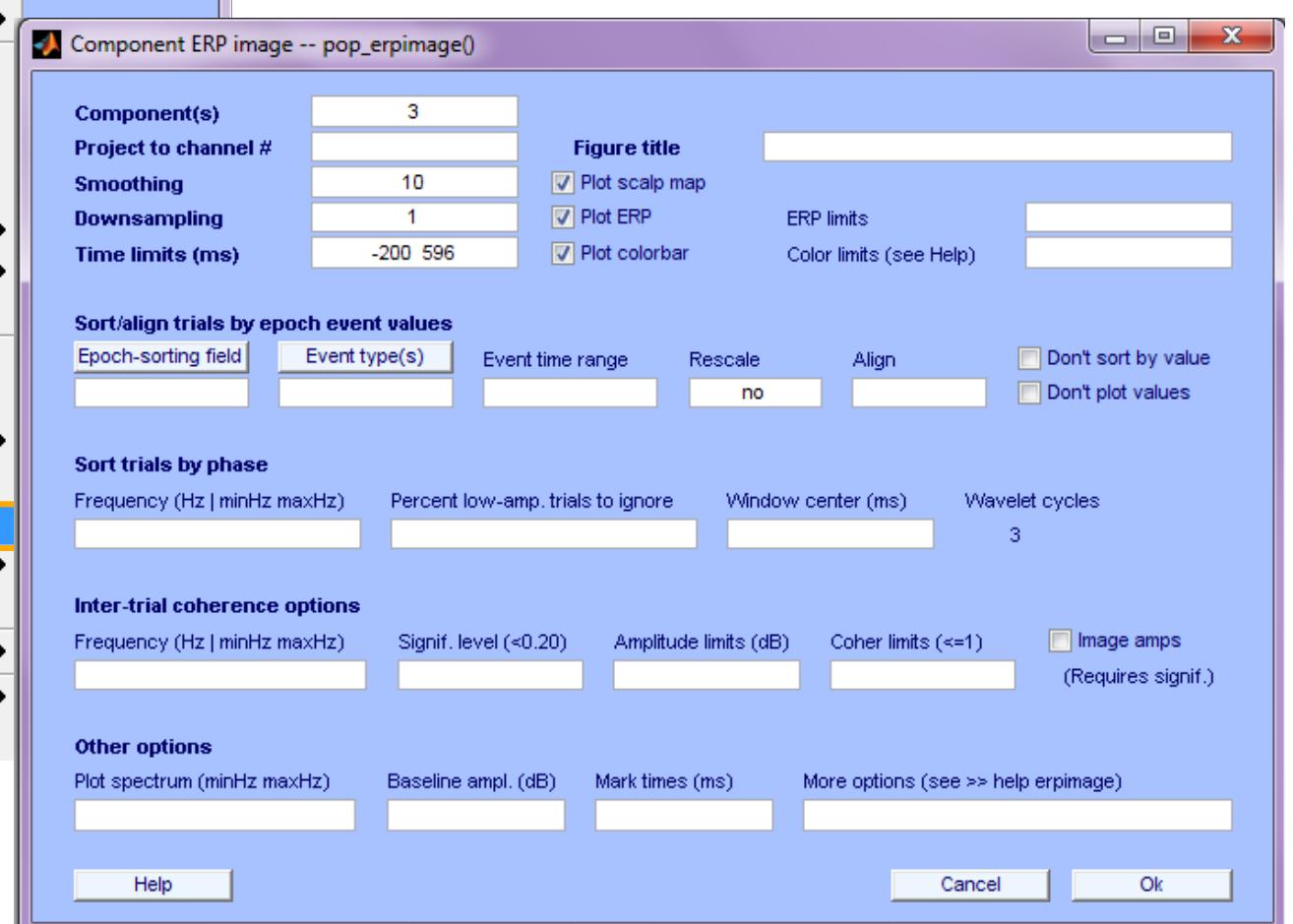
Epoch on Memorize
letters



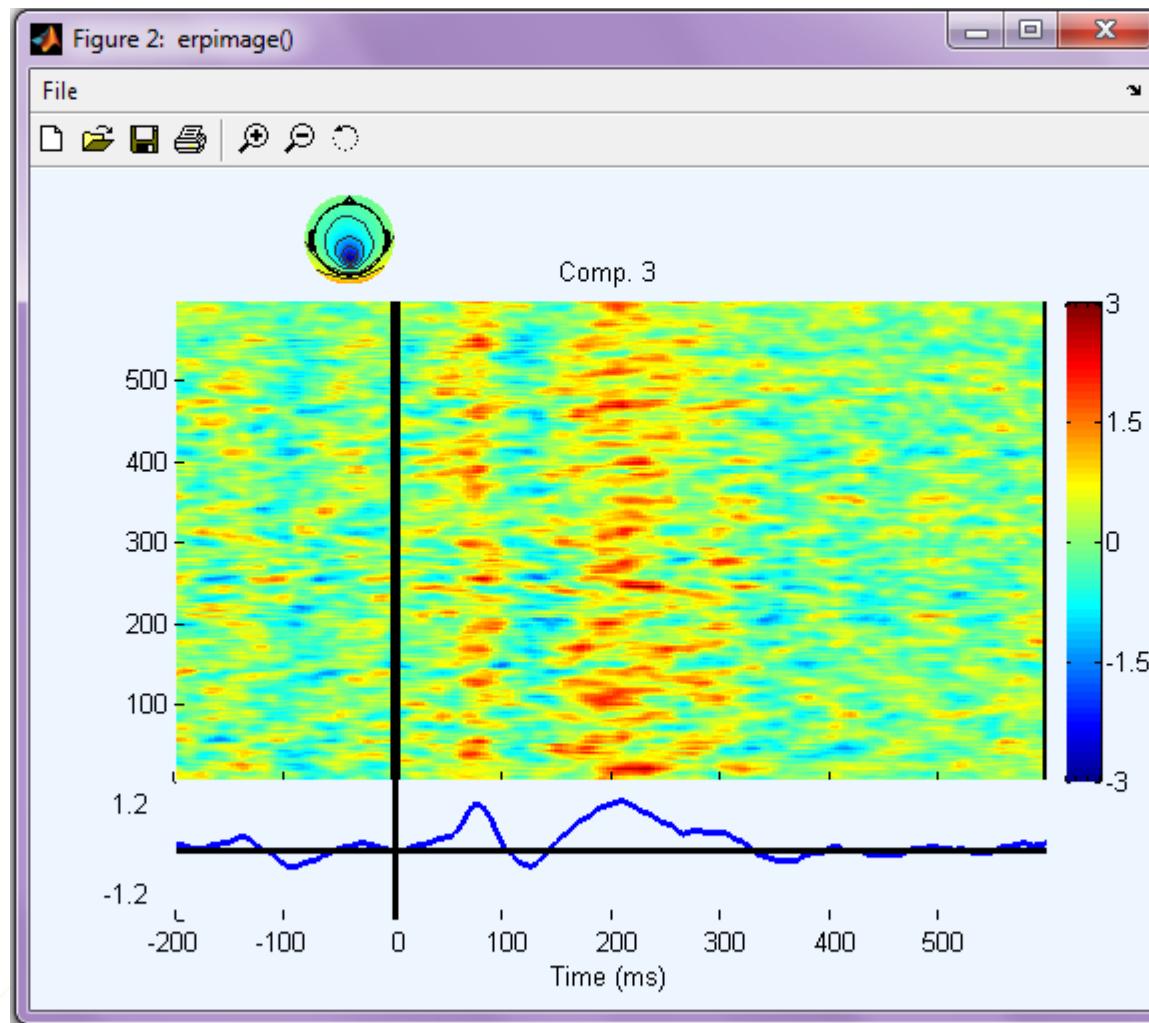
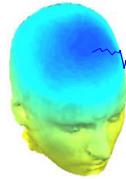
Create a script from ‘eegh’ output



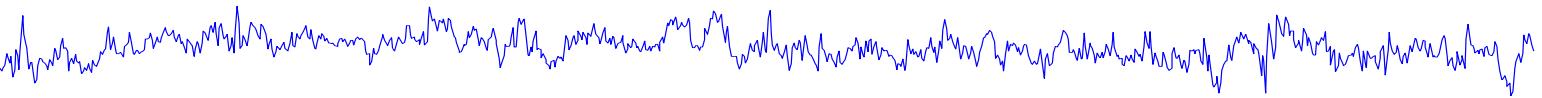
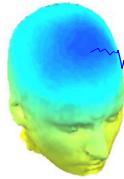
Plot an IC ERP image



Create a script from ‘eegh’ output



Retrieve commands from eegh

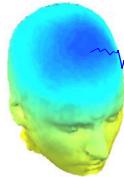


Write a script to do this:

```
>> eegh
```



Retrieve commands from eegh



```
>> eegh
[ALLEEG EEG CURRENTSET ALLCOM] = eeglab;

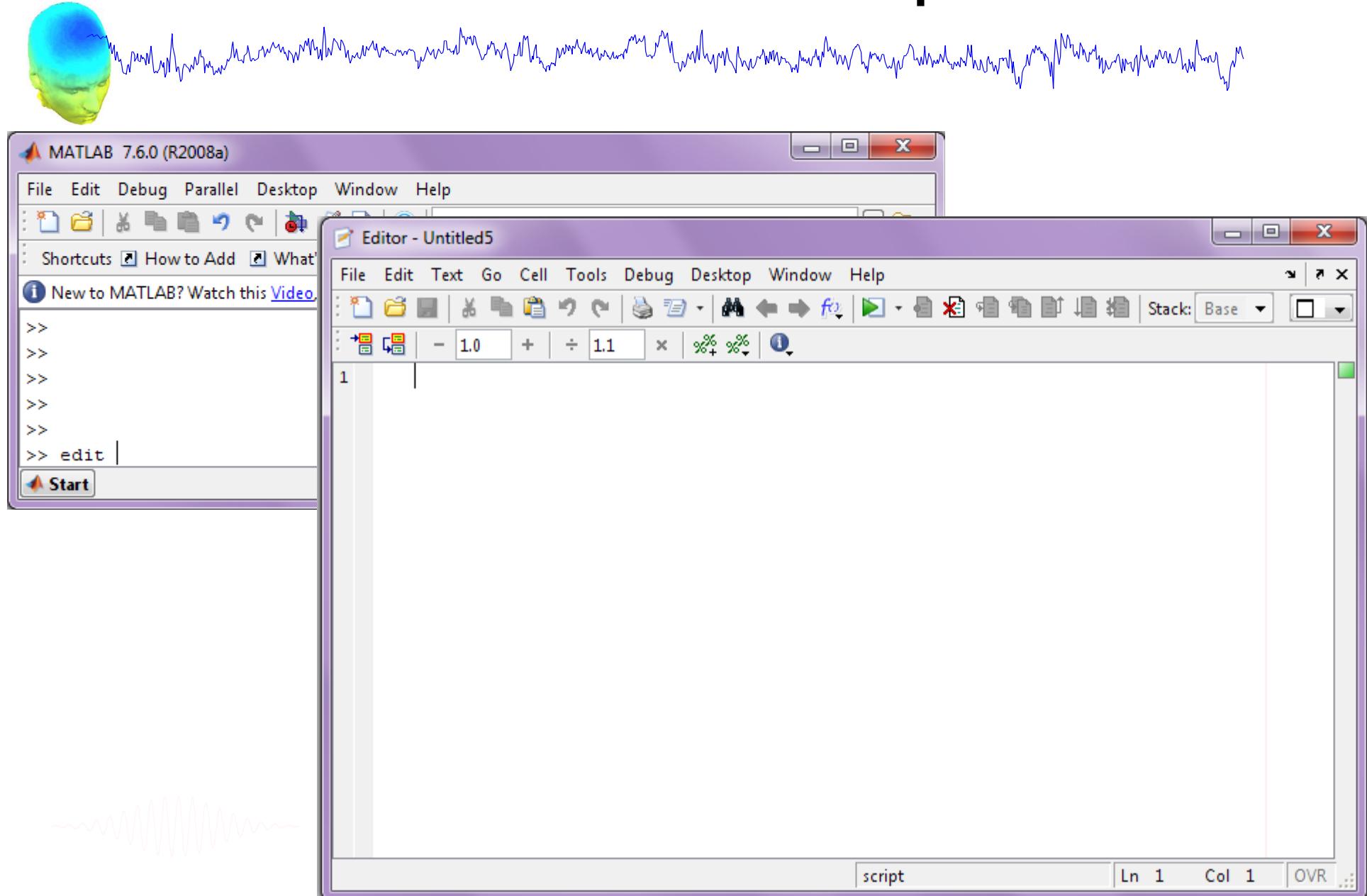
EEG = pop_loadset('filename', 'stern_125Hz.set');
[ALLEEG EEG CURRENTSET] = eeg_store(ALLEEG, EEG, 0);

EEG = pop_epoch( EEG, {'B' 'C' 'D' ... }, [-0.2 0.6], 'newname',
'Memorize epochs', 'epochinfo', 'yes');
[ALLEEG EEG CURRENTSET] = eeg_store(ALLEEG, EEG, 1);
EEG = pop_rmbase( EEG, [-200 0]);
[ALLEEG EEG] = eeg_store(ALLEEG, EEG, CURRENTSET);

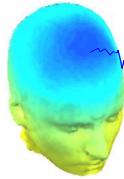
figure; pop_erpimage(EEG,0, [3],[],'Comp. 3',10,1,{},
[],'', 'yerplabel', '', 'erp', 'on', 'cbar', 'on','topo',
{mean(EEG.icawinv(:,[3]),2) EEG.chanlocs EEG.chaninfo});
```



Create a Matlab script



Create a Matlab script



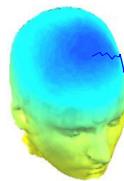
Copy and paste from Matlab window:



```
Editor - Untitled5*
File Edit Text Go Cell Tools Debug Desktop Window Help
Stack: Base
1 [ALLEEG EEG CURRENTSET ALLCOM] = eeglab;
2 EEG = pop_loadset( 'filename', 'stern.set', 'filepath', 'C:\\\\Users\\\\julie\\\\Docum
3 [ALLEEG, EEG, CURRENTSET] = eeg_store( ALLEEG, EEG, 0 );
4 EEG = pop_epoch( EEG, { 'B' 'C' 'D' 'F' 'G' 'H' 'J' 'K' 'L' 'M' 'N'
5 [ALLEEG EEG CURRENTSET] = pop_newset(ALLEEG, EEG, 1, 'gui', 'off');
6 EEG = pop_rmbase( EEG, [-200 0]);
7 [ALLEEG EEG] = eeg_store(ALLEEG, EEG, CURRENTSET);
8 figure; pop_erpimage(EEG,0, [3],[],'Comp. 3',10,1,{},[],'', 'yerplabel','','er
script Ln 7 Col 51 OVR
```

Save as 'ploterpimage.m'
In MATLAB folder

Run your new script



MATLAB 7.6.0 (R2008a)

File Edit Debug Parallel Desktop Window Help

C:\Users\julie\Documents\MATLAB

Shortcuts How to Add What's New

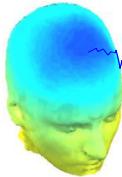
New to MATLAB? Watch this [Video](#), see [Demos](#), or read [Getting Started](#).

```
>>
>>
>>
>>
>> edit
>> ploterpimage|
```

Start OVR



Exercise page 1



```
>> eeglab
```

```
% load dataset,  
% epoch on 'memorize letter' B, C, etc...  
% plot erpimage for component 3
```

```
>> eegh
```

```
% open Matlab editor
```

```
>> edit
```

```
% copy & paste eegh results into a new  
% file and save it (ploterpimage.m)
```

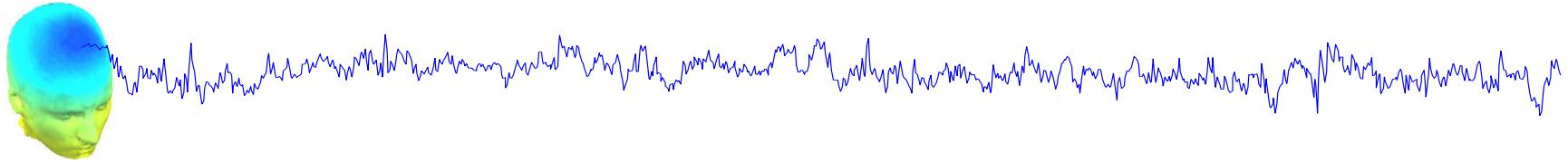
```
>> clear
```

```
>> close all
```

```
>> ploterpimage
```

```
>> eeglab redraw
```

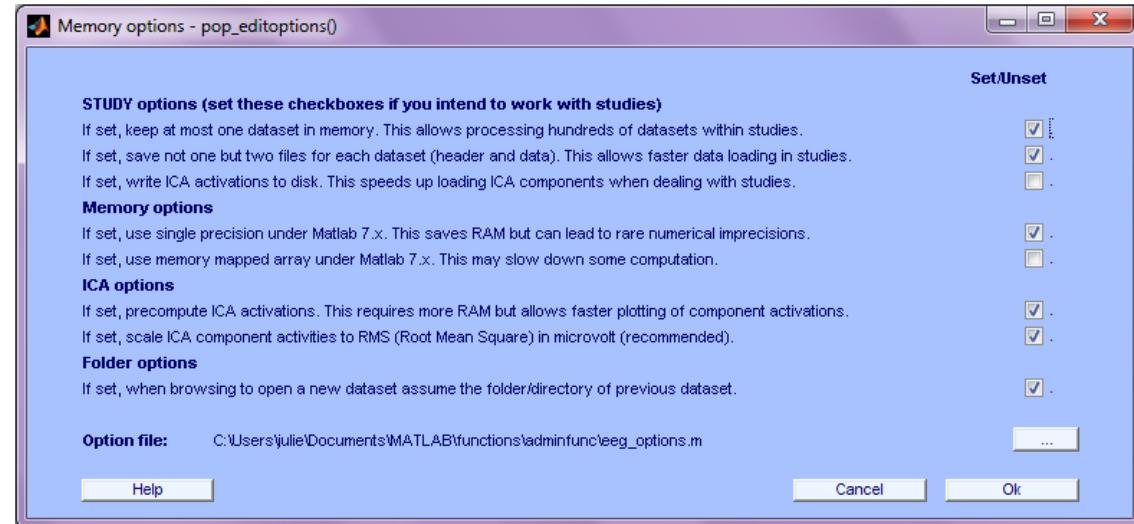
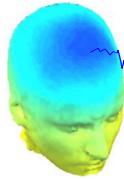




Advanced Scripting in EEGLAB



STUDY scripts



Most important option:

- Allows only one dataset to be loaded at once.
- Most STUDYs are too big to have all data loaded at once.

% Set memory options:

`pop_editoptions('option_storedisk' , 1)`

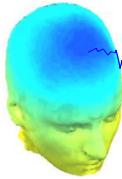
Edit dataset info



The image shows the EEGLAB v13.x dev software interface. On the left, there's a 3D head model with a blue brain surface and a blue waveform plot above it. The main window has a menu bar with File, Edit, Tools, Plot, Study, Datasets, and Help. The 'Study' menu is open, showing options like 'Create study', 'Load existing study', and 'Save current study'. A sub-menu under 'Create study' is also open, showing 'Using all loaded datasets', 'Browse for datasets', and 'Simple ERP STUDY'. To the right, a dialog box titled 'Create a new STUDY set -- pop_study()' is displayed. It asks for 'STUDY set name:' (Stenberg), 'STUDY set task name:', and 'STUDY set notes:'. Below this is a table for dataset management, with columns for 'dataset filename', 'browse', 'subject', 'session', 'condition', 'group', and 'Select by r.v.'. There are 10 rows, each corresponding to a dataset file named 'S01'. The 'condition' column shows values 'Ignore', 'Memorize', and 'Probe'. The 'group' column contains empty fields. To the right of the table are 'Clear' buttons for each row. At the bottom of the dialog, there's an 'Important note: Removed datasets will not be saved before being deleted from EEGLAB memory' and two checkboxes: 'Dataset info (condition, group, ...) differs from study info. [set]' and 'Delete cluster information (to allow loading new datasets, set new components for clustering, etc.)'. At the very bottom are 'Help', 'Cancel', and 'Ok' buttons.

```
[STUDY ALLEEG] = std_editset( STUDY, ALLEEG, 'name', 'Stenberg', 'commands', {{'index' 1
'load' '/data/STUDY/S01/Ignore.set'} {'index' 2 'load' '/data/S01/Memorize.set'}
{'index' 3 'load' '/data/S01/Probe.set'}} {'index' 1 'subject' 'S01'} {'index' 2
'subject' 'S01'} {'index' 3 'subject' 'S01'} {'index' 1 'condition' 'Ignore'} {'index' 2
'condition' 'Memorize'} {'index' 3 'condition' 'Probe'}}), 'updatedat', 'off' );
```

Looking at the function that create STUDY



```
[STUDY ALLEEG] = std_editset( STUDY, ALLEEG, 'name','Stenberg','commands',
  {{'index' 1 'load' '/data/STUDY/S01/Ignore.set'} ...
  {'index' 2 'load' '/data/S01/Memorize.set'} ...
  {'index' 3 'load' '/data/S01/Probe.set'} ...
  {'index' 1 'subject' 'S01'} ...
  {'index' 2 'subject' 'S01'} ...
  {'index' 3 'subject' 'S01'} ...
  {'index' 1 'condition' 'Ignore'} ...
  {'index' 2 'condition' 'Memorize'} ...
  {'index' 3 'condition' 'Probe'}},'updatedat','off' );
```

```
[STUDY ALLEEG] = std_editset( STUDY, ALLEEG, 'name','Stenberg','commands',
  {{'index' 1 'load' '/data/STUDY/S01/Ignore.set' 'subject' 'S01' 'condition' 'Ignore'} ...
  {'index' 2 'load' '/data/S01/Memorize.set' 'subject' 'S01' 'condition' 'Memorize'} ...
  {'index' 3 'load' '/data/S01/Probe.set' 'subject' 'S01' 'condition' 'Probe'} ...
}, 'updatedat','off' );
```



Exercice



```
[STUDY ALLEEG] = std_editset( STUDY, ALLEEG, 'name','Stenberg','commands',
{{'index' 1 'load' '/data/STUDY/S01/Ignore.set' 'subject' 'S01' 'condition' 'Ignore'} ...
{'index' 2 'load' '/data/S01/Memorize.set' 'subject' 'S01' 'condition' 'Memorize'} ...
{'index' 3 'load' '/data/S01/Probe.set' 'subject' 'S01' 'condition' 'Probe'} ...
}, 'updatedat', 'off' );
```

If not present, add it by hand because
some dataset might not have it

1- Start EEGLAB and import the 3 datasets for Subject 1 (Ignore.set, Memorize.set and Probe.set) in a STUDY (menu Tools > Create STUDY > Browse for datasets)

2- Look in the history

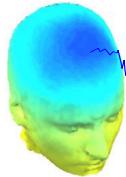
3- Copy to a script, add “eeglab redraw” at the end of your script

4- Restart Matlab, execute the script, look at your STUDY info and design (menu *STUDY > Edit STUDY info* and *STUDY > Select/Edit STUDY design*)

5- Modify the script to import subject 1 to 4

6- Restart Matlab, execute the script, look at your STUDY info and design

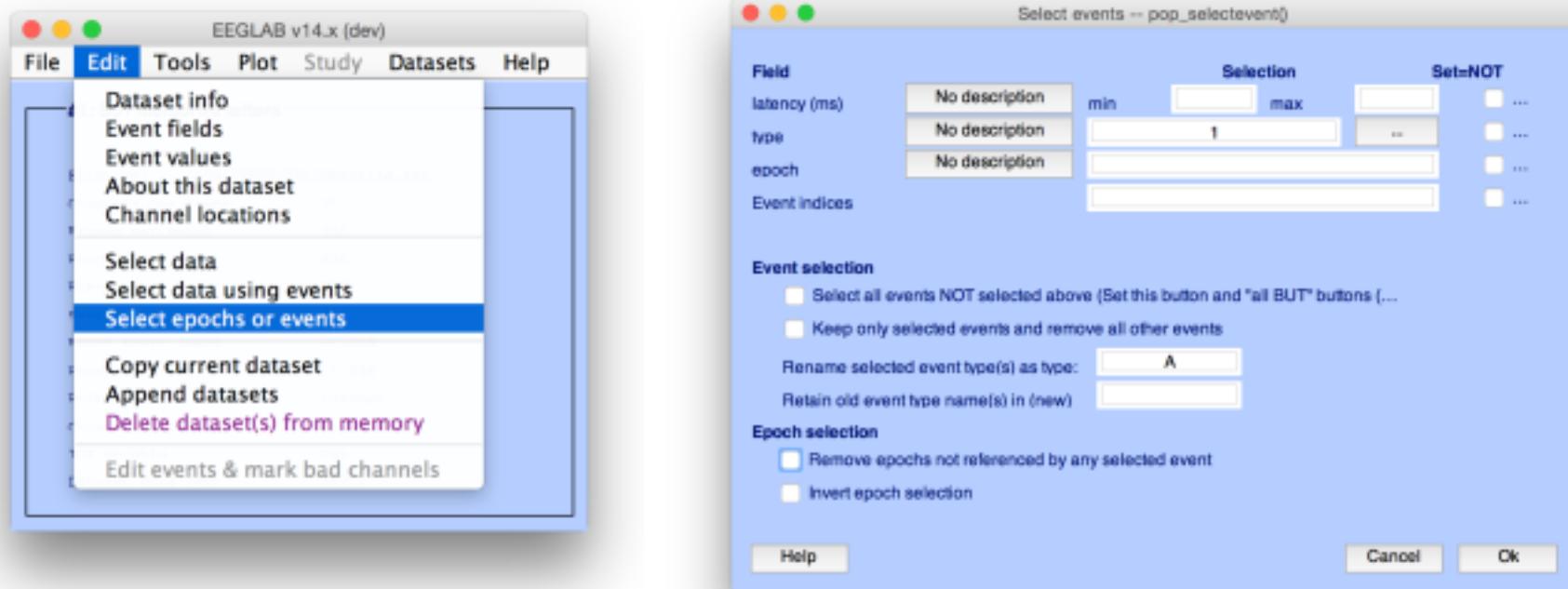
Redefining events



Adjusting latencies

```
for iEvent = 1:length(EEG.event)
    % shift by 16 samples (or 53.3ms at 200Hz) due to filter delay
    EEG.event(iEvent).latency = EEG.event(iEvent).latency + 16;
end;
EEG.saved = 'no';
[ALLEEG EEG CURRENTSET] = eeg_store(ALLEEG, EEG);
eeglab redraw
```



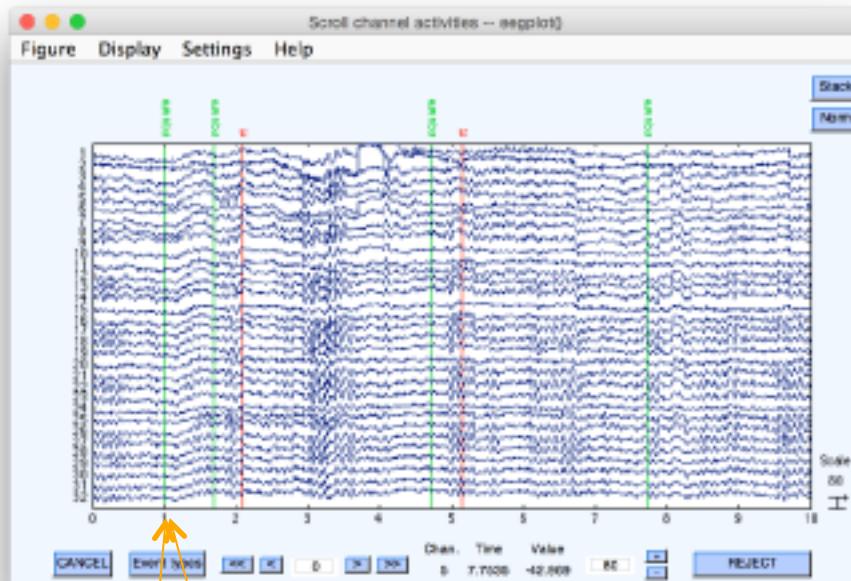


Renaming events

```

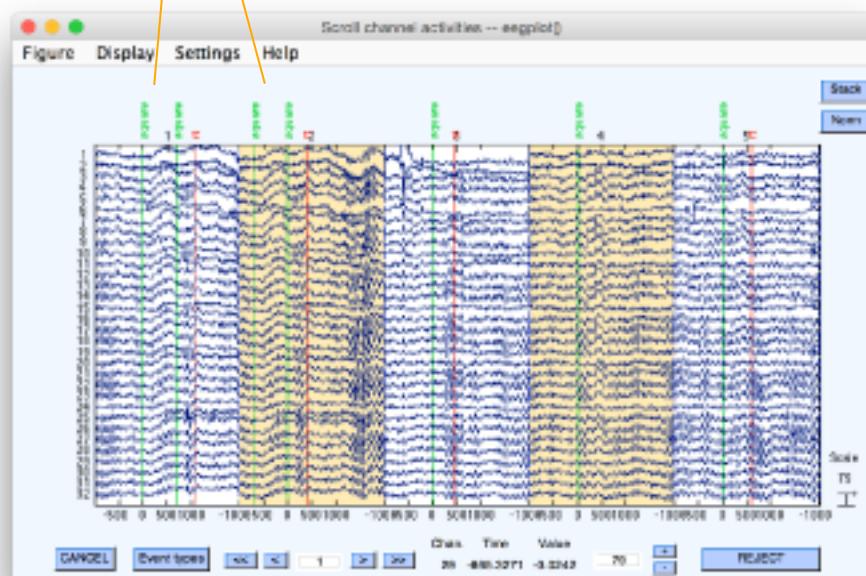
for iDat = 1:length(ALLEEG)
    EEG = ALLEEG(iDat);
    % rename events
    EEG = pop_selectevent( EEG, 'type',1,'renametype','A','deleteevents','off');
    EEG = pop_selectevent( EEG, 'type',2,'renametype','B','deleteevents','off');
    EEG = pop_selectevent( EEG, 'type',3,'renametype','C','deleteevents','off');
    EEG = pop_selectevent( EEG, 'type',4,'renametype','D','deleteevents','off');
    EEG = pop_selectevent( EEG, 'type',8,'renametype','rt','deleteevents','off');
    EEG.saved = 'no';
    [ALLEEG EEG CURRENTSET] = eeg_store(ALLEEG, EEG, iDat);
end;
eeglab redraw

```



EEG.event(4)

```
    type: 'square'  
position: 2  
latency: 424  
urevent: 1  
epoch: 2
```



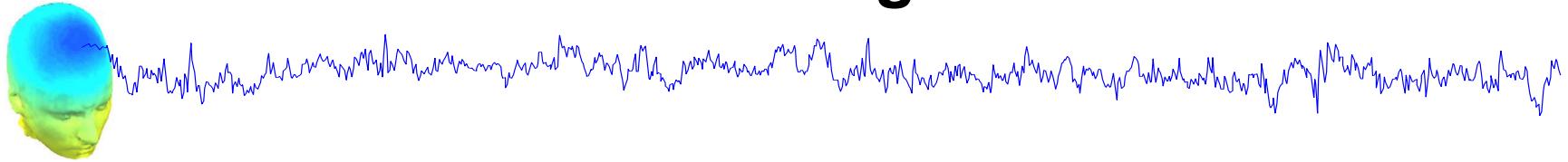
EEG.event(1)

```
    type: 'square'  
position: 2  
latency: 129  
urevent: 1  
epoch: 1
```

EEG.urevent(1)

```
    type: 'square'  
position: 2  
latency: 129
```

Redefining events



```
for iDat = 1:length(ALLEEG)
    TMPEEG = eeg_checkset(ALLEEG(iDat), 'loaddata'); % load data

    % 'B' 'C' 'D' 'F' 'G' 'H' 'J' 'K' 'L' ... -> Memorize
    % 'gB' 'gC' 'gD' 'gF' 'gG' 'gH' 'gJ' 'gK' 'gL' ... -> Ignore
    for iEvent = 1:length(TMPEEG.event)
        prevEvent = TMPEEG.event(iEvent).urevent-2;
        if prevEvent > 2 && TMPEEG.urevent(prevEvent).type(1) == 'g'
            TMPEEG.event(iEvent).prevEvent = 'ignore';
        else TMPEEG.event(iEvent).prevEvent = 'memorize';
        end;
    end;

    TMPEEG.saved = 'no'; % tag as not saved
    ALLEEG(iDat) = pop_saveset(TMPEEG, 'savemode', 'resave'); % resave data
end;

STUDY = std_maketrialinfo(STUDY, ALLEEG); % update STUDY
STUDY.saved = 'no';
[STUDY EEG] = pop_savestudy( STUDY, EEG, 'savemode', 'resave'); % resave STUDY
```

Precomputed files need to be recomputed after changing events.

Edit STUDY design -- pop_studydesign()

Select STUDY design New Rename Delete Design Matrix

memorize vs ignore
Design 2
Design 3
Design 4

Resave STUDY

Edit selected design

Independent variables New Import Edit Delete

Categorical variable: condition - Values (probe - ignore .
Categorical variable: prevEvent - Values (ignore - memo

Subjects

S07
S08
S09
S10
S11
S12
S13

Delete all pre-computed datafiles for this STUDY design

Web help Cancel Ok

Add variable

Select independent variable

prevEvent
regtime
rt
stimulus
time
type
uncertainty1

Categorical variable

Select variable values

ignore
memorize

Combine selected values

Cancel Ok

Design Matrix: Design 4

Row number: S01

Column headers: condition-probe, condition-ignore & memorize, prevEvent-ignore, prevEvent-memorize, constant

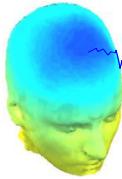
Trails	1	2	3	4	5
100	1	1	1	1	1
200	1	1	1	1	1
300	1	1	1	1	1
400	1	1	1	1	1
500	1	1	1	1	1
600	1	1	1	1	1
700	1	1	1	1	1

Design Matrix

Value on Click

24

Load dataset info from commandline



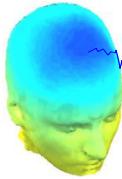
```
% Create Stern STUDY
[ALLEEG EEG CURRENTSET ALLCOM] = eeglab;
pop_editoptions( 'option_storedisk', 1);
subjects = {'S01' 'S02' 'S03' 'S04' 'S05' 'S06' 'S07' 'S08' 'S09' 'S10' 'S11' 'S12'};
filepath = '/Users/arno/temp/STUDY'; % XXXXX Change path here XXXXX
if ~exist(filepath), error('You need to change the path to the STUDY'); end;
commands = {};% initialize STUDY dataset list

% Loop through all of the subjects in the study to create the dataset
for loopnum = 1:length(subjects) %for each subject
    IgnoreFile = fullfile(filepath, subjects{loopnum}, 'Ignore.set');
    MemorizeFile = fullfile(filepath, subjects{loopnum}, 'Memorize.set');
    ProbeFile = fullfile(filepath, subjects{loopnum}, 'Probe.set');
    commands = {commands{:} ...
        {'index' 3*loopnum-2 'load' IgnoreFile 'subject' subjects{loopnum} 'condition' 'Ignore'} ...
        {'index' 3*loopnum-1 'load' MemorizeFile 'subject' subjects{loopnum} 'condition' 'Memorize'} ...
        {'index' 3*loopnum 'load' ProbeFile 'subject' subjects{loopnum} 'condition' 'Probe'}};
end;
% Uncomment the line below to select ICA components with less than 15% residual variance
% commands = {commands{:} {'dipselect', 0.15}};
[STUDY, ALLEEG] = std_editset(STUDY, ALLEEG, 'name','Sternberg', 'commands', commands, 'updatedat', 'on');

% Update workspace variables and redraw EEGLAB
CURRENTSTUDY = 1; EEG = ALLEEG; CURRENTSET = [1:length(EEG)];
[STUDY, ALLEEG] = std_checkset(STUDY, ALLEEG);
eeglab redraw
```

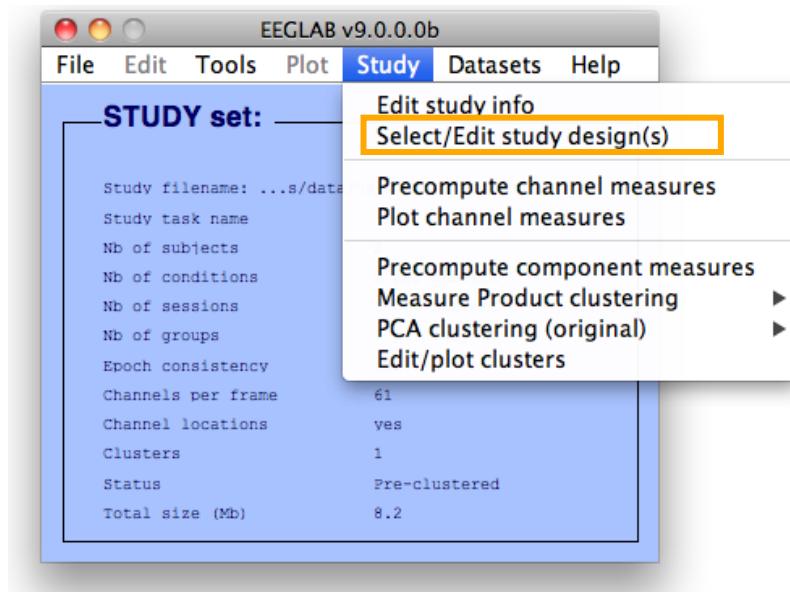
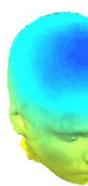


STUDY structure



```
STUDY =
    name: 'Sternberg'
    task: 'Sternberg'
datasetinfo: [1x39 struct]
    notes: ''
    filename: 'stern.study'
    filepath: 'C:\Users\julie\Documents\Workshops\Finland\STUDY'
    history: [1x7332 char]
    subject: {1x13 cell}
    group: {''}
    session: []
    condition: {'ignore' 'memorize' 'probe'}
design: [1x1 struct]
    etc: [1x1 struct]
    preclust: [1x1 struct]
cluster: [1x1 struct]
changrp: [1x71 struct]
    saved: 'yes'
```



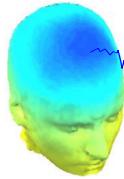


```
STUDY = std_makedesign(STUDY, ALLEEG, 3,
'variable1','condition',
'variable2','','',
'name','Design 3',
'velues1',{'ignore' 'memorize' 'probe'},
'subjselect',{'S02' 'S03'},
'dataselect',{'condition' {'probe'}});
```

Select subjects



STUDY design structure



```
STUDY.design(1)

ans =

    name: 'Design 1 - compare letter types'
    variable: [1x2 struct]
    cases: [1x1 struct]
    include: {}
    cells: [1x39 struct]
```

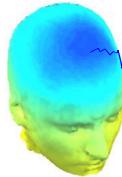
Exploding the contents of each of these sub-structures, we obtain

```
name: 'Design 1 - light and audio all subjects'

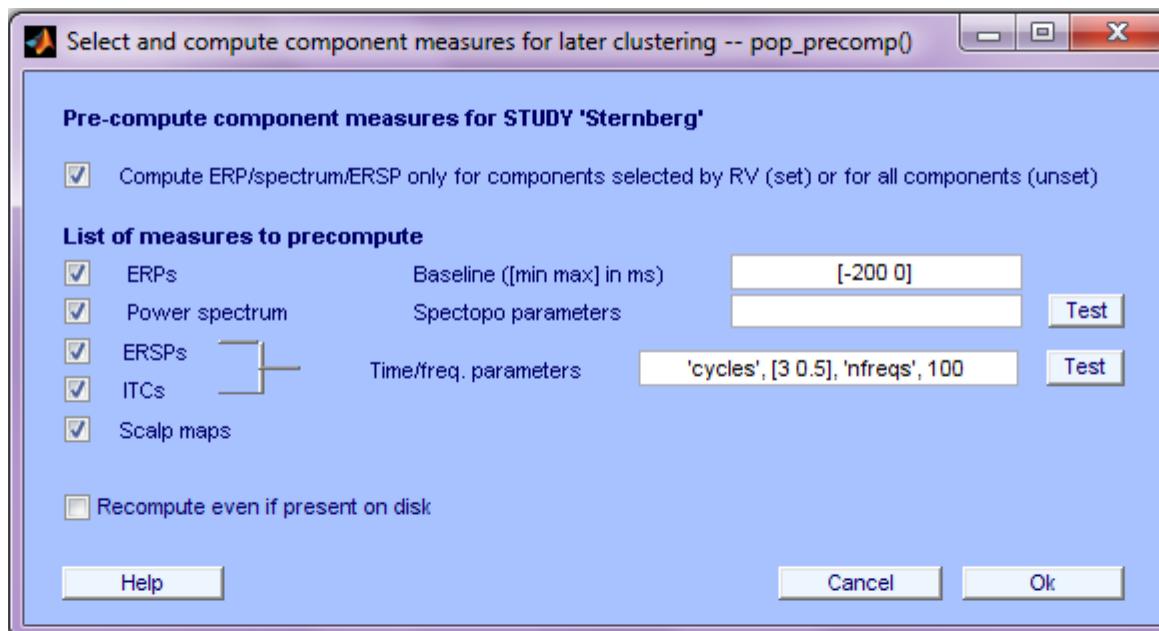
variable: [1x2 struct]
    (1).label : 'condition'
    (1).pairing: 'on'
    (1).value : {'ignore' 'memorize' 'probe'}
    (2).label : ''
    (2).pairing: 'off'
    (2).value : {}

cases: [1x1 struct]
    label: 'subject'
    value: {'S01' 'S02' 'S03' 'S04' 'S05' 'S06' }
```

Precompute data measures

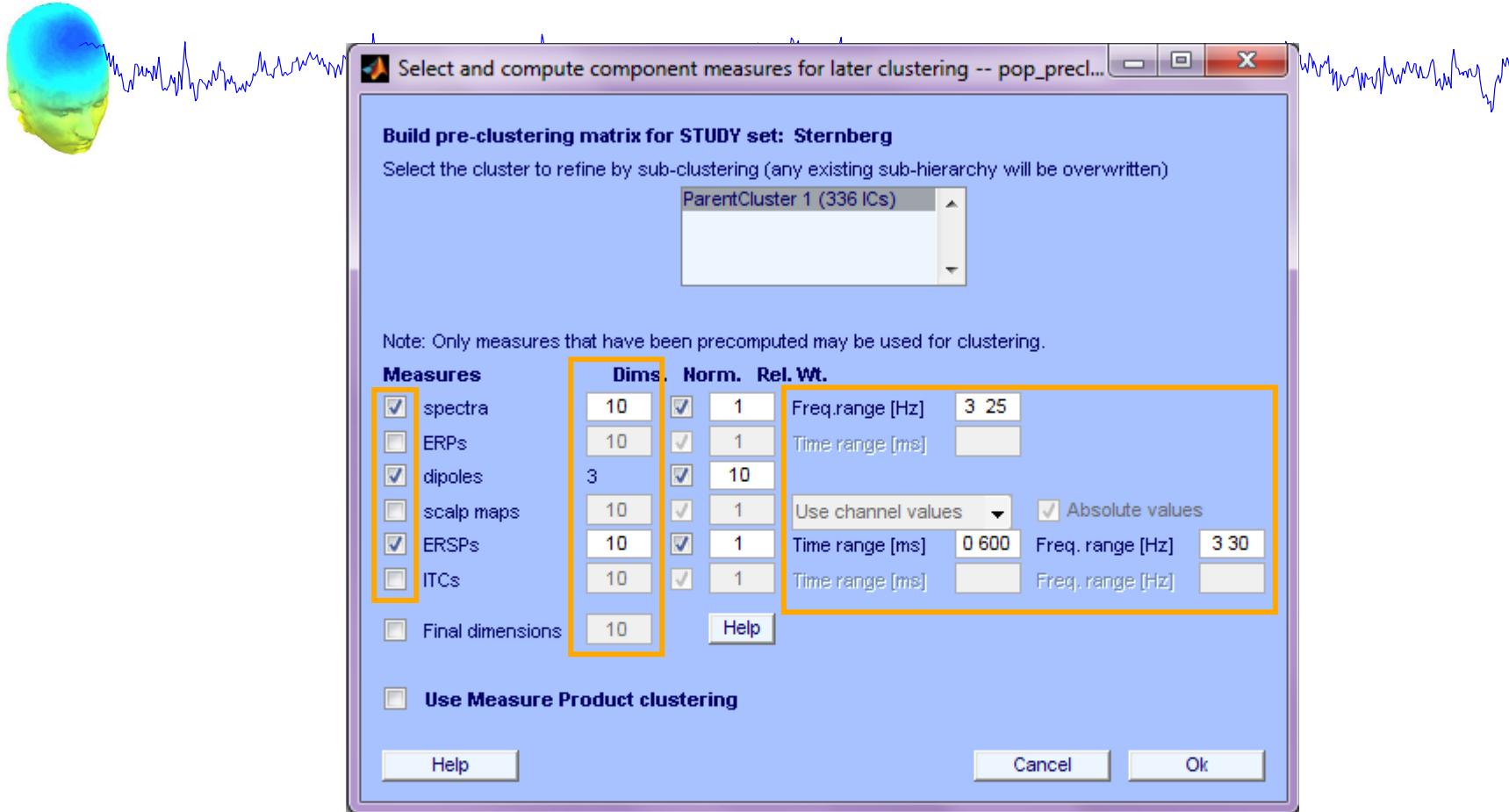


TIP: Compute all measures so you can test different combinations for clustering



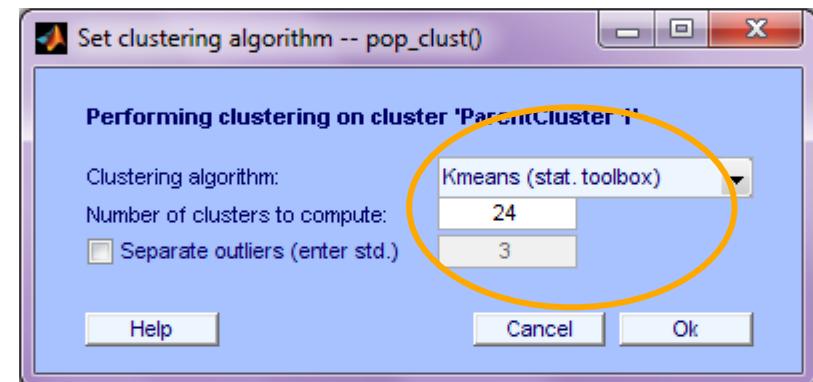
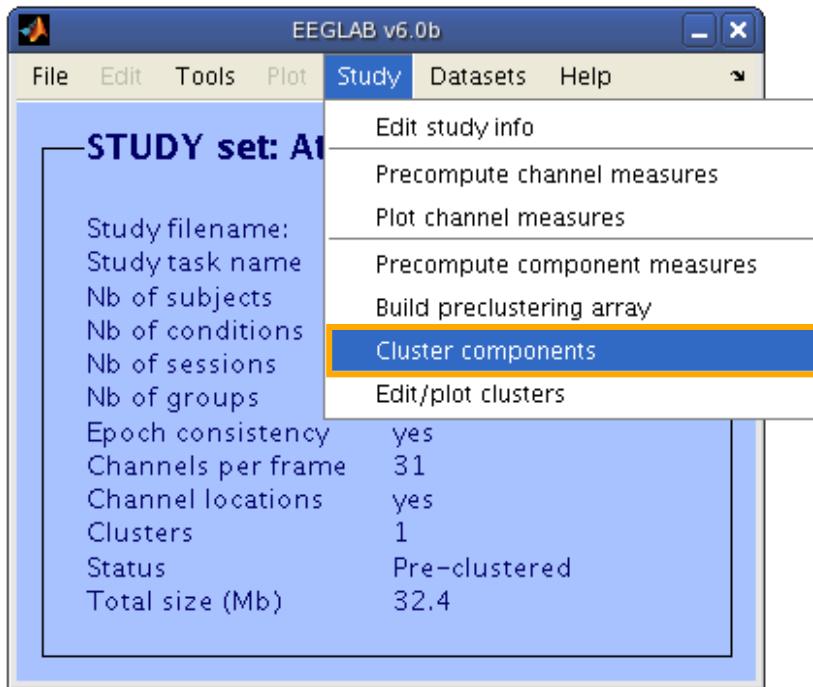
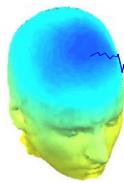
```
[STUDY ALLEEG] = std_precomp(STUDY, ALLEEG, 'components', 'erp', ...
'on', 'rmbase', [-200 0], 'scalp', 'on', 'spec', 'on', ...
'specparams', {}, 'ersp', 'on', 'erspparams', {'cycles' [3 0.5] ...
'nfreqs', 100, 'freqs', [3 70]}, 'itc', 'on');
```

Precluster the data



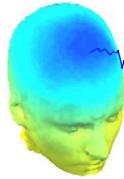
```
[STUDY ALLEEG] = std_preclust(STUDY, ALLEEG, 1, {'spec','npca',5,...  
'norm',1,'weight',1,'freqrange',[3 25]},{'erp','npca',6,'norm',1,...  
'weight',1, 'timewindow',[0 400]},{'scalp','npca',10,'norm',1,'weight',1,...  
'abso',1},{'dipoles','norm',1,'weight',10},{'ersp','npca',20,...  
'freqrange',[3 30] , 'timewindow',[0 600], 'norm',1,'weight',1},{'itc',...  
'npca',6,'freqrange',[3 30], 'timewindow',[0 400] , 'norm',1, 'weight',1});
```

Cluster components



```
[STUDY] = pop_clust(STUDY, ALLEEG, 'algorithm', 'kmeans', 'clus_num', 24);
```

Understanding STUDY structure



26 = # of clusters

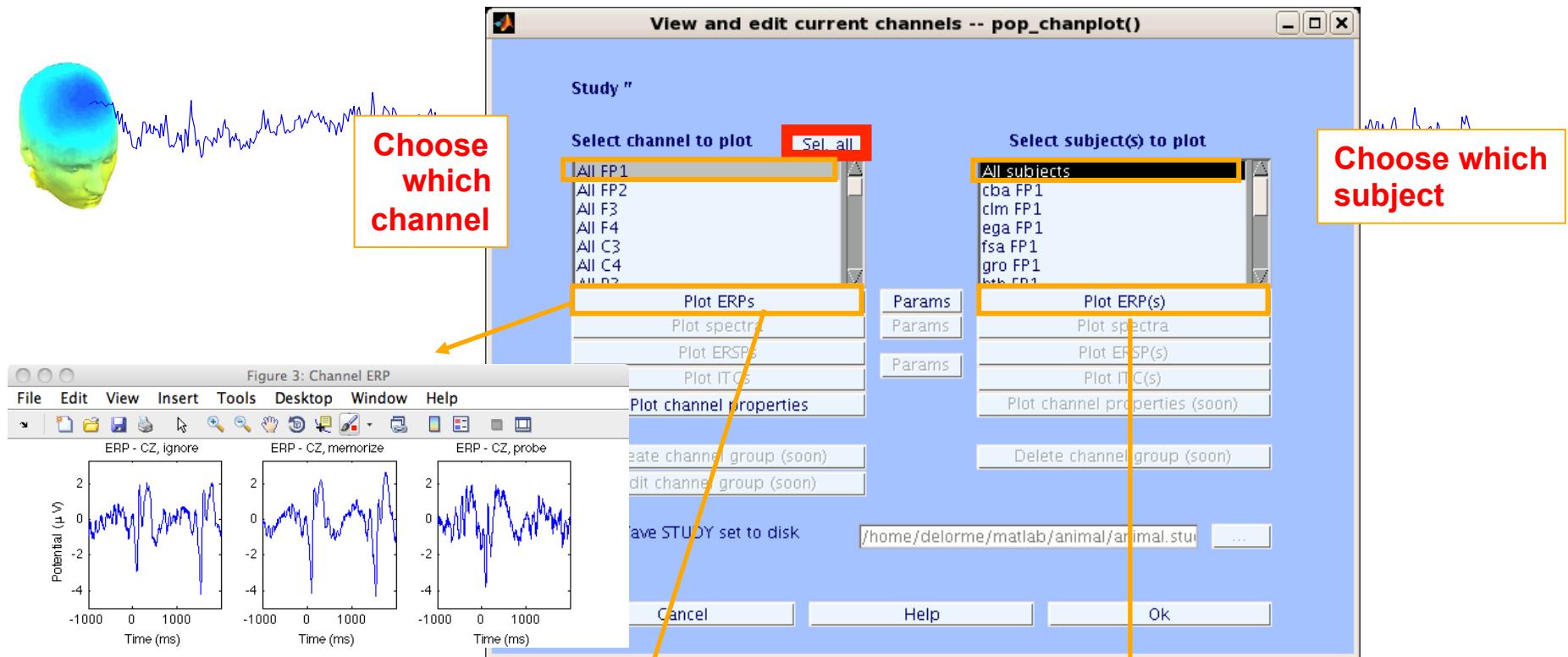
```
>> STUDY.cluster  
1x26 struct array with fields:  
  
parent  
name  
child  
comps  
sets  
algorithm  
preclust  
dipole  
allinds  
setinds
```

One cluster:

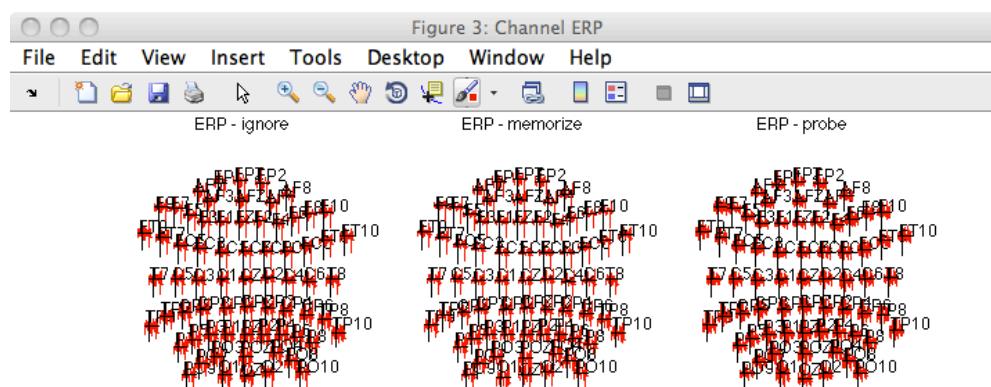
```
>> STUDY.cluster(6) → 6 = cluster index  
ans =  
  
parent: {'ParentCluster 1'}  
name: 'Cls 6'  
child: []  
comps: [35 7 12 35 10 23 7 30 4 ...]  
sets: [1 2 3 4 5 6 7 8 9 10 1 2 ...]  
algorithm: {'Kmeans' [24]}  
preclust: [1x1 struct]  
dipole: [1x1 struct]
```

IC indices

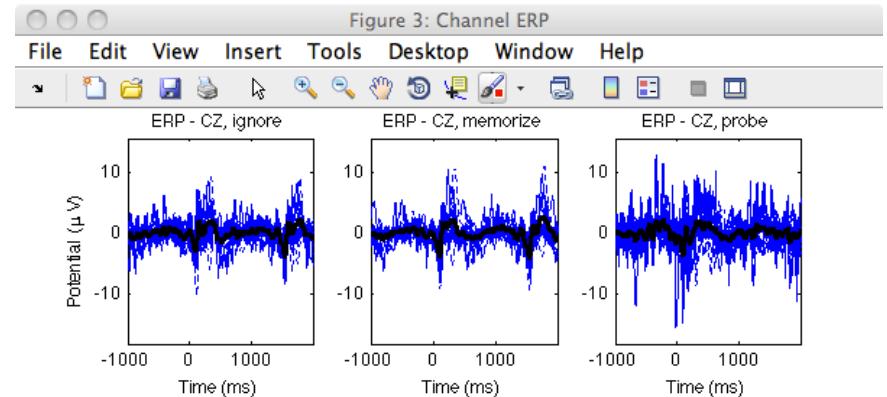
dataset indices for ICs



```
STUDY = std_erpplot(STUDY, ALLEEG, 'channels', {'FP1'});
```

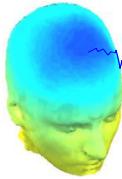


```
STUDY = std_erpplot(STUDY, ALLEEG, 'channels', {'FP1'} ... );
```

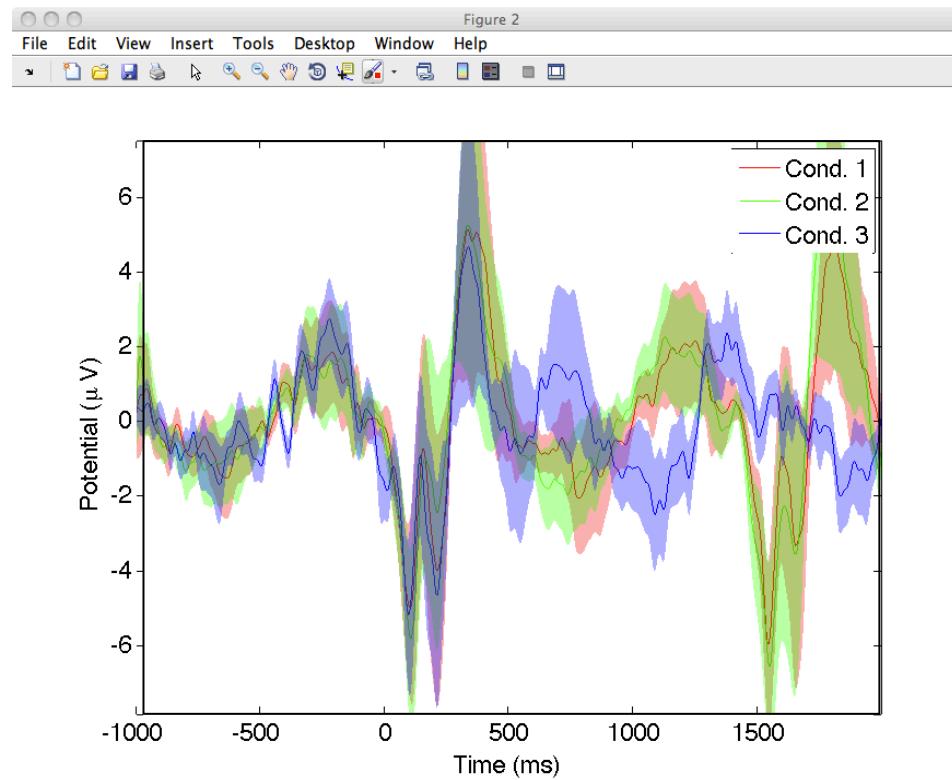


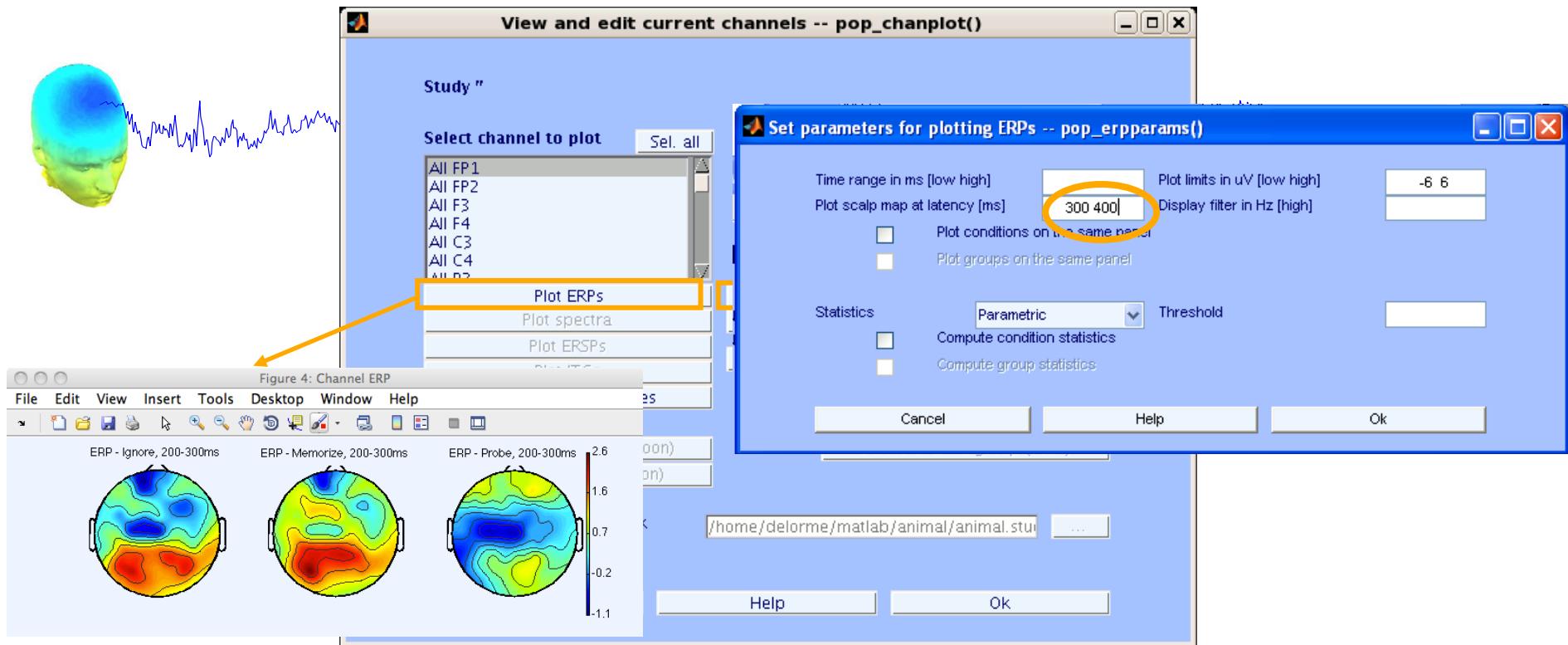
```
STUDY = std_erpplot(STUDY, ALLEEG, 'channels', {'FP1'}, 'plotsubjects', 'on' );
```

Advanced plotting features



```
STUDY = std_erpplot(STUDY, ALLEEG, 'channels', {'CZ'});  
std_plotcurve(STUDY.changrp(39).erptimes,  
STUDY.changrp(39).erpdata, 'plotconditions',  
'together', 'plotstderr', 'on', 'filter', 30);
```





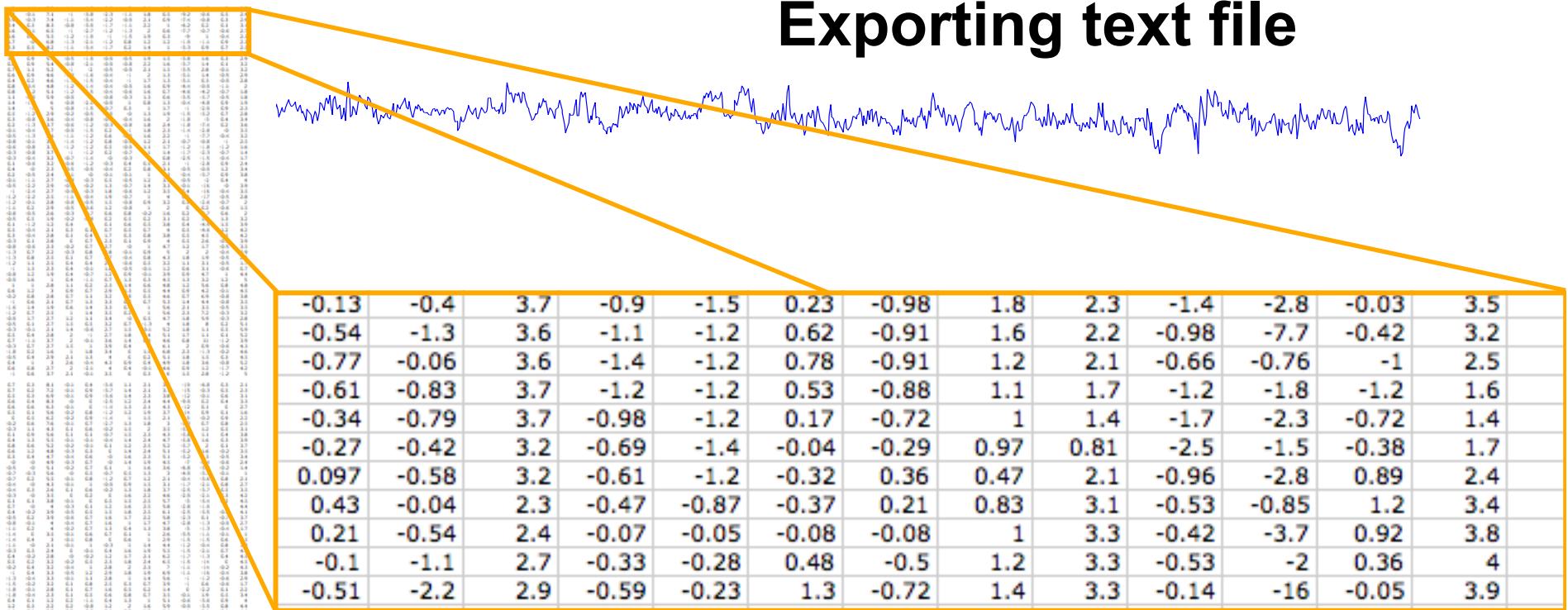
```
STUDY = std_erpplot(STUDY, ALLEEG, 'topotime', [200 300], 'channels', {'OZ' 'O2' 'FP1' 'FPZ' 'FP2'});
[STUDY erpdata] = std_erpplot(STUDY, ALLEEG, , 'topotime', [200 300], 'channels', {'OZ' 'O2'});
```

Exporting to excell file

[1x67x13 double]
 [1x67x13 double]
 [1x67x13 double]

```
xlswrite('myxlsfile', squeeze(erpdata{1}), 1);
xlswrite('myxlsfile', squeeze(erpdata{2}), 2);
xlswrite('myxlsfile', squeeze(erpdata{3}), 3);
```

Exporting text file



```

dlmwrite('erpfile.txt',squeeze(erpdata{1}),'delimiter', '\t', 'precision', 2);

dlmwrite('erpfile.txt',squeeze(erpdata{2}),'-append', 'roffset', 1,
         'delimiter', '\t', 'precision', 2);

dlmwrite('erpfile.txt',squeeze(erpdata{2}),'-append', 'roffset', 1,
         'delimiter', '\t', 'precision', 2);

```

Exercice: run STUDY Script

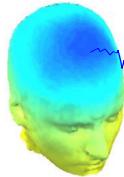
```
% Create Stern STUDY
[ALLEEG EEG CURRENTSET ALLCOM] = eeglab;
pop_editoptions('option_storedisk', 1);
subjects = {'S01' 'S02' 'S03' 'S04' 'S05' 'S06' 'S07' 'S08' 'S09' 'S10' 'S11' 'S12'};
filepath = '/Users/arno/temp/STUDY'; % XXXXX Change path here XXXXX
if ~exist(filepath), error('You need to change the path to the STUDY'); end;
commands = {};% initialize STUDY dataset list

% Loop through all of the subjects in the study to create the dataset
for loopnum = 1:length(subjects) %for each subject
    IgnoreFile = fullfile(filepath, subjects{loopnum}, 'Ignore.set');
    MemorizeFile = fullfile(filepath, subjects{loopnum}, 'Memorize.set');
    ProbeFile = fullfile(filepath, subjects{loopnum}, 'Probe.set');
    commands = {commands{:} ...
        {'index' 3*loopnum-2 'load' IgnoreFile 'subject' subjects{loopnum} 'condition' 'Ignore'} ...
        {'index' 3*loopnum-1 'load' MemorizeFile 'subject' subjects{loopnum} 'condition' 'Memorize'} ...
        {'index' 3*loopnum 'load' ProbeFile 'subject' subjects{loopnum} 'condition' 'Probe'}};
end;
% Uncomment the line below to select ICA components with less than 15% residual variance
% commands = {commands{:} {'dipselect', 0.15}};
[STUDY, ALLEEG] = std_editset(STUDY, ALLEEG, 'name', 'Sternberg', 'commands', commands, 'updatedat', 'on');

% Update workspace variables and redraw EEGLAB
CURRENTSTUDY = 1; EEG = ALLEEG; CURRENTSET = [1:length(EEG)];
[STUDY, ALLEEG] = std_checkset(STUDY, ALLEEG);
eeglab redraw

[STUDY ALLEEG] = std_precomp(STUDY, ALLEEG, {}, 'rmicacomp', 'on', 'interp', 'on', 'recompute', 'on', 'erp', 'on');
STUDY = pop_erpparams(STUDY, 'topotime', [200 300] );
[STUDY erpdata] = std_erpplot(STUDY, ALLEEG, 'channels', {'LEYE' 'REYE' 'OZ' 'O2' 'FP1' 'FPZ' 'FP2' 'AF7' ...
    'AF3' 'AFZ' 'AF4' 'AF8' 'F9' 'F7' 'F5' 'F3' 'F1' 'FZ' 'F2' 'F4' 'F6' 'F8' 'F10' 'FT9' ...
    'FT7' 'FC5' 'FC3' 'FC1' 'FCZ' 'FC2' 'FC4' 'FC6' 'FT8' 'FT10' 'T7' 'C5' 'C3' 'C1' 'CZ' ...
    'C2' 'C4' 'C6' 'T8' 'TP9' 'TP7' 'CP5' 'CP3' 'CP1' 'CPZ' 'CP2' 'CP4' 'CP6' 'TP8' 'TP10' ...
    'P7' 'P5' 'P3' 'P1' 'PZ' 'P2' 'P4' 'P6' 'P8' 'PO9' 'PO7' 'PO3' 'POZ' 'PO4' 'PO8' 'PO10' 'O1'});
dlmwrite('erpfile.txt', squeeze(erpdata{1}), 'delimiter', '\t', 'precision', 2);
dlmwrite('erpfile.txt', squeeze(erpdata{2}), '-append', 'roffset', 1, 'delimiter', '\t', 'precision', 2);
dlmwrite('erpfile.txt', squeeze(erpdata{2}), '-append', 'roffset', 1, 'delimiter', '\t', 'precision', 2);
```

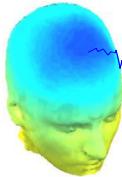
Advanced scripting: EEG pipeline



```
sInfo = [];
sInfo(end+1).file = 'rawdata/S01.raw'; ← Raw data file
sInfo(end).name = 'S01'; ← Subject name
sInfo(end).bad_channels = { 'E1' }; ← Subject name
```



Advanced scripting: EEG pipeline

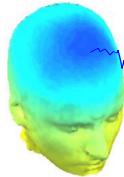


```
sInfo = [];
sInfo(end+1).file = 'rawdata/S01.raw'; ← Raw data file
sInfo(end).name = 'S01'; ← Subject name
sInfo(end).bad_channels = { 'E1' }; ← Subject name
sInfo(end).bad_data = [726 1495;6098 6831;13245 14057;15715 16399;22756 2445'
```

Copy the output from the eeg_eegrej function in the history



Advanced scripting: EEG pipeline

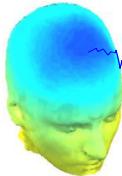


```
sInfo = [];
sInfo(end+1).file = 'rawdata/S01.raw'; ← Raw data file
sInfo(end).name = 'S01'; ← Subject name
sInfo(end).bad_channels = { 'E1' }; ← Subject name
sInfo(end).bad_data = [726 1495;6098 6831;13245 14057;15715 16399;22756 2445];
sInfo(end).bad_comps = [1.6681 1.9870 0.3979 0.4444 -0.2274 -0.1433 -0.2626 ·
                        1.1917 -1.4838 0.7469 -1.1599 0.4773 -0.3257 0.3074 ·
```

Copy transposed columns of the inverse weight matrix
EEG.icawinv for your selected artifact components



Advanced scripting: EEG pipeline



EEG signal visualization: A blue waveform representing raw EEG data over time.

```
sInfo = [];
sInfo(end+1).file = 'rawdata/S01.raw';
sInfo(end).name = 'S01';
sInfo(end).bad_channels = { 'E1' };
sInfo(end).bad_data = [726 1495; 6098 6831; 13245 14057; 15715 16399; 22756 2445];
sInfo(end).bad_comps = [1.6681 1.9870 0.3979 0.4444 -0.2274 -0.1433 -0.2626 -
1.1917 -1.4838 0.7469 -1.1599 0.4773 -0.3257 0.3074 -
```

datainfo.m file

```
sInfo(end+1).file = 'rawdata/S02.raw';
sInfo(end).name = 'S02';
sInfo(end).bad_channels = { };
sInfo(end).bad_data = [41661 43713; 24000 24833; 44878 46501; 48706 49210; 51190];
sInfo(end).bad_comps = [0.6960 -0.8637 0.9087 -0.8028 0.4873 -0.2142 0.273 -
-0.0875 -0.4056 -0.0287 -0.3870 0.0600 -0.3716 0.342!
2.1928 1.5712 0.8622 0.3215 -0.0357 -0.3125 -0.226!
```

```
sInfo(end+1).file = 'rawdata/S03.raw';
sInfo(end).name = 'S03';
sInfo(end).bad_channels = { 'E10' 'E19' 'E20' 'E29' };
sInfo(end).bad_data = [1 10449; 19808 21815; 25678 27254; 29257 30010; 34023 360;
sInfo(end).bad_comps = [ 1.8583 2.0468 -0.0516 0.3159 -0.4256 -0.2770 -0.36 -
1.2189 -0.7385 1.2464 -0.8913 0.5475 -0.3971 0.29!
-0.1248 -0.1358 -0.1954 -0.2533 -0.1555 -0.2313 -0.03!
```

EEG signal visualization: A red waveform representing raw EEG data over time.

```

datainfo;
pop_editoptions( 'option_storedisk' , 1 );
Folder = 'ds_eeg_preprocessed_data';
if ~exist(outputfolder), mkdir(outputfolder); end;

for iSubj = 1:length(sInfo)

    % load dataset
    EEG = pop_biosig( sInfo(iSubj).file );
    EEG.setname = sInfo(iSubj).name;

    % preprocess data
    EEG = pop_iirfilt( EEG, 0.5, 0, [], 0, 0); % high pass filtering
    EEG = pop_iirfilt( EEG, 0, 55, [], 0, 0); % low pass filtering
    EEG = pop_select(EEG, 'nochannel', sInfo(iSubj).bad_channels); % remove bad channels
    EEG = pop_reref( EEG, []); % average reference (optional)
    EEG = eeg_eegrej( EEG, sInfo(iSubj).bad_data); % remove bad portions of data

    % run ICA
    EEG = pop_runica(EEG, 'pca', EEG.nbchan-1);

    % tag bad components
    % EEG = pop_findmatchingrejcomps(EEG, 'matchcomps',sInfo(iSubj).bad_comps,'corrthresh',0.92);
    if ~isempty(sInfo(iSubj).bad_comps)
        [corr,indx,indy] = matcorr(EEG.icawinv', sInfo(iSubj).bad_comps);
        if ~all(abs(corr(1:size(sInfo(iSubj).bad_comps,1))) > 0.92),
            error('Correlation too low'); end;
        EEG.reject.gcompreject(indx(1:size(sInfo(iSubj).bad_comps,1))) = 1;
    end;

    % extract data epochs
    EEG = pop_epoch(EEG, { 'S1' 'S2' });

    % save dataset
    EEG.saved = 'no';
    EEG = pop_saveset( EEG, 'filepath', folder, 'filename', [ sInfo(iSubj).name '.set' ]);
end;

```

Create STUDY



```
datainfo;
pop_editoptions( 'option_storedisk', 1);
outputEEGFolder = 'preprocessed_data';
studyCommand    = {};

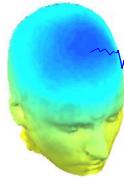
% generate STUDY commands
for iSubject = 1:length(sInfo)
    fileName = fullfile(outputEEGFolder, [ sInfo(iSubject).name '.set' ]);
    studyCommand = [ studyCommand { 'index' iSubject 'load' fileName 'subject' ...
        sInfo(iSubject).name } ];
end;

% create data
[STUDY ALLEEG] = std_editset( [], [], 'name', 'test', 'commands', studyCommand, ...
    'updatedat','off', 'filename', 'test.study', 'resave', 'on');
STUDY = std_makedesign(STUDY, ALLEEG, 1, 'variable1','','variable2','','name', 'Design 1', ...
    'pairing1','on','pairing2','on','delfiles','off','defaultdesign','off','subjselect', ...
    {sInfo.name}, 'filepath', 'studyfiles');

% update workspace variables and redraw EEGLAB
CURRENTSTUDY = 1; EEG = ALLEEG; CURRENTSET = [1:length(EEG)];
[STUDY, ALLEEG] = std_checkset(STUDY, ALLEEG);
eeglab redraw

% precompute and plot data
[STUDY ALLEEG] = std_precomp(STUDY, ALLEEG, {},'interp','on','recompute','on','erp', 'on');
[STUDY erp] = std_erpplot(STUDY,ALLEEG, 'channels',{allchanlocs.labels}, 'topotime',[200 300]);
```

Exercice: build your own pipeline



Suggestion for exercise

1. Load SimpleOddball.set dataset
2. High pass filter at 1Hz (menu Tools > Filter)
3. Reject bad portion of data by hand (menu Tools > Reject continuous...)
4. Re-reference to average ref. (optional) (menu Tools > Re-reference)
5. Build your *datainfo.m* file using EEGLAB history (see scripting lecture)
6. Run ICA (use PCA if necessary); select bad ICA components
7. Epoch data on Oddball (type 2) and Standard (type 1) – save dataset
8. Add components to you *datainfo.m* file
9. Create a STUDY with this single file
10. Compare the ERP for Oddball (type 2) and Standard (type 1) and use single-trial statistics with cluster correction for multiple comparisons
11. Build a script that creates the STUDY and perform the same analysis
12. Save the figure at the end of the script in eps or jpg format (“print –deps file” command or “print –djpeg file” command).
13. Run the full pipeline (dataset processing and STUDY processing)
14. Change the filtering in the pipeline (step 2) and observe effects