

# **STUDY plotting and statistics**

## **STEP 1**

Build a STUDY

## **STEP 2**

Build design(s)

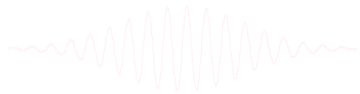
## **STEP 3**

Precompute the data

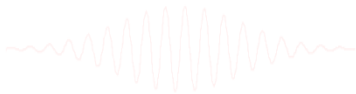
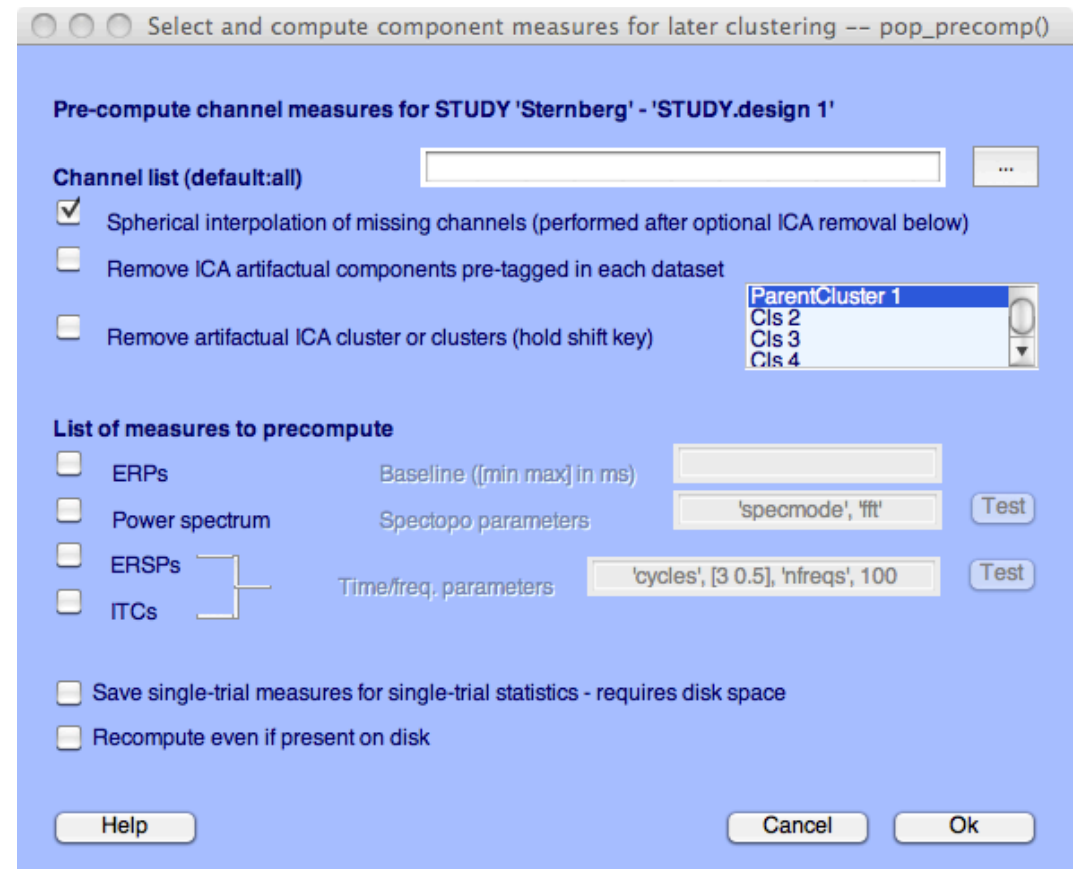
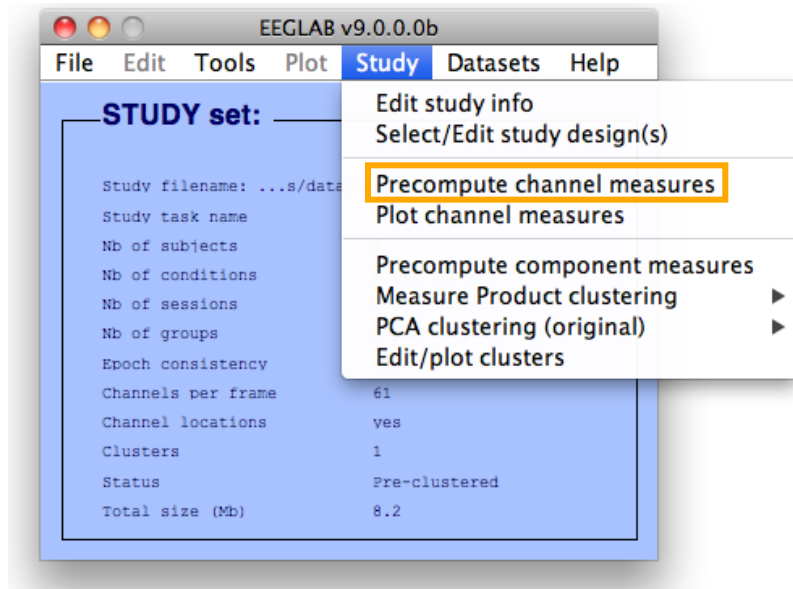
## **STEP 4**

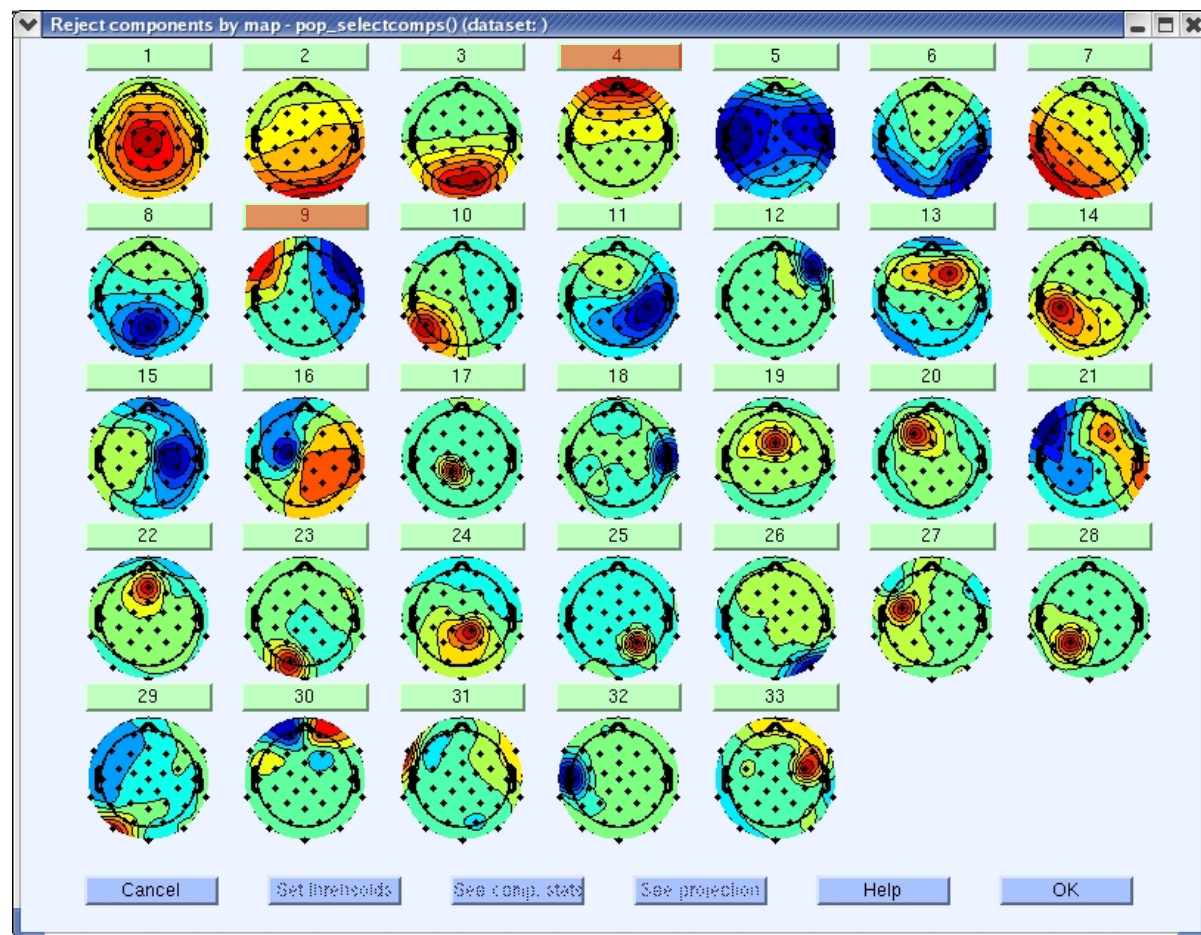
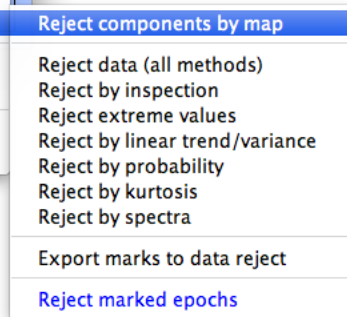
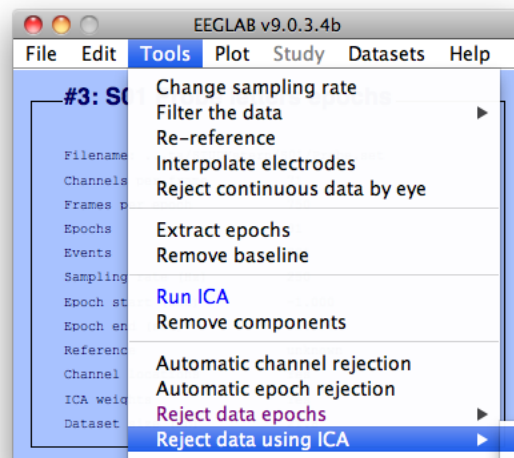
Plot the data

Exercise...

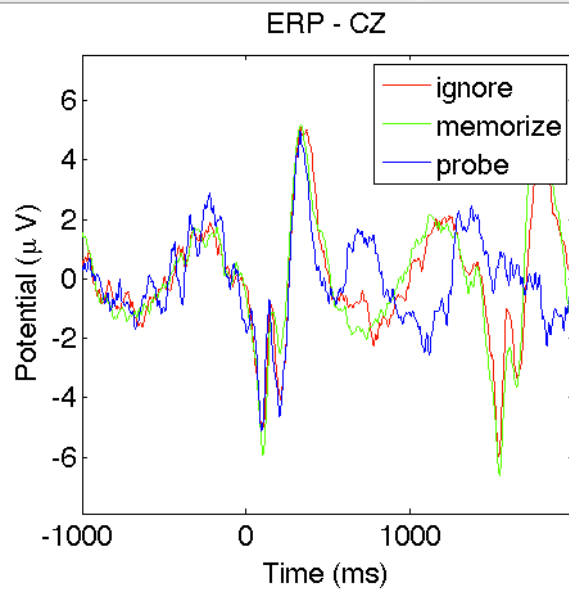
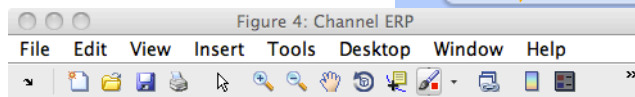
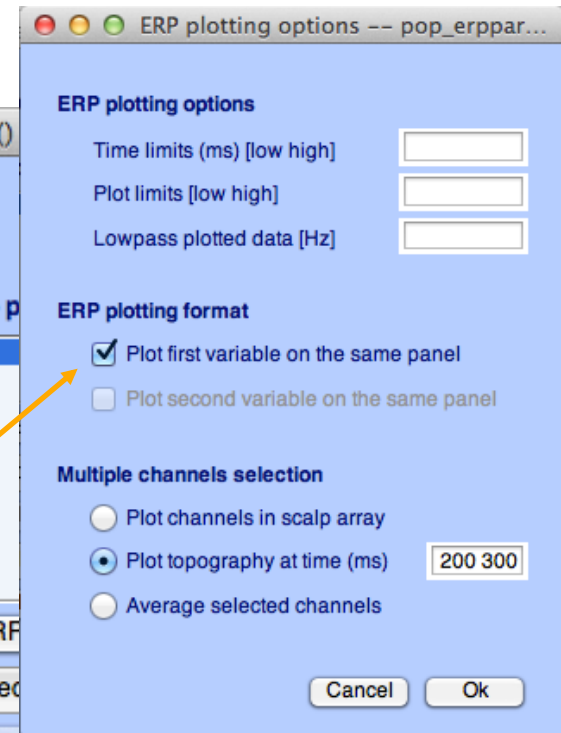
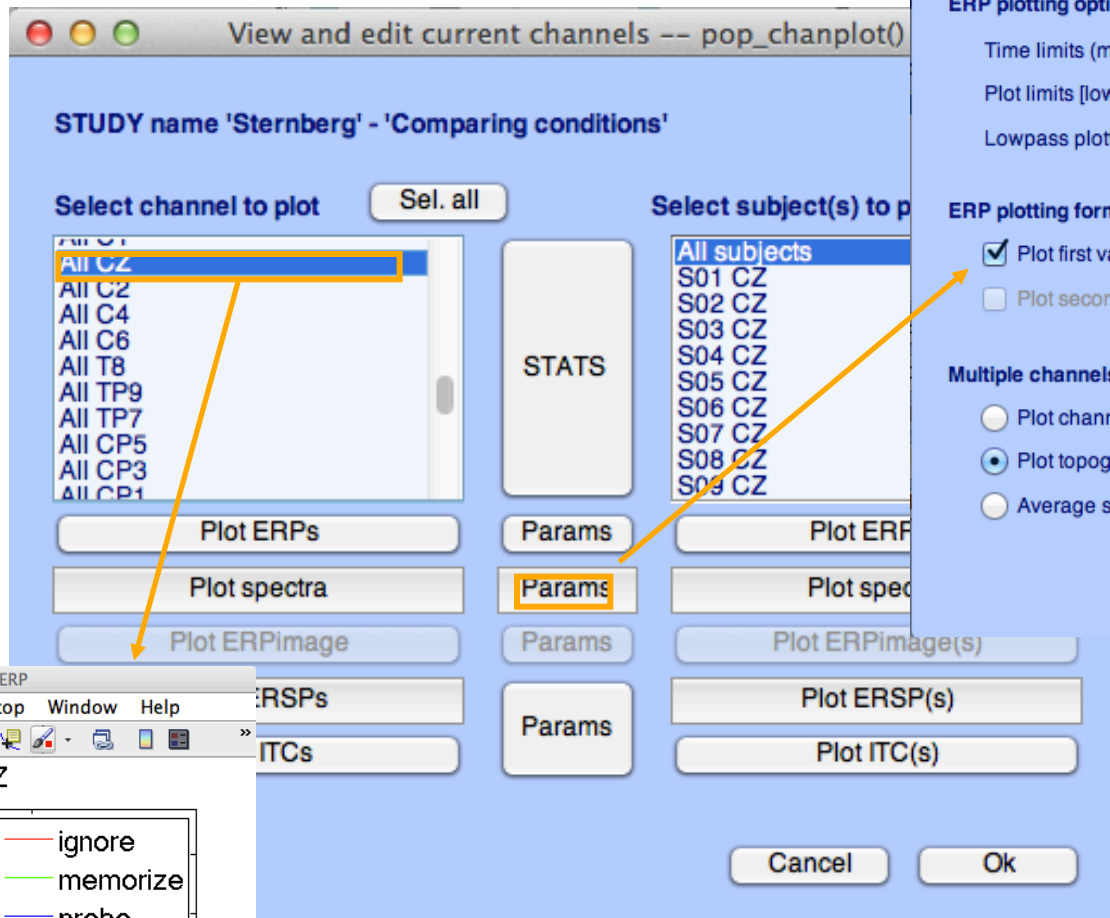


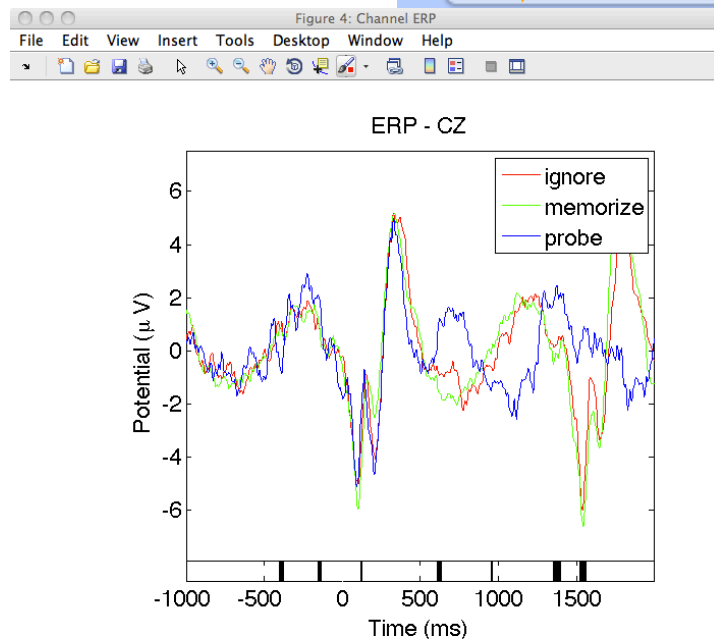
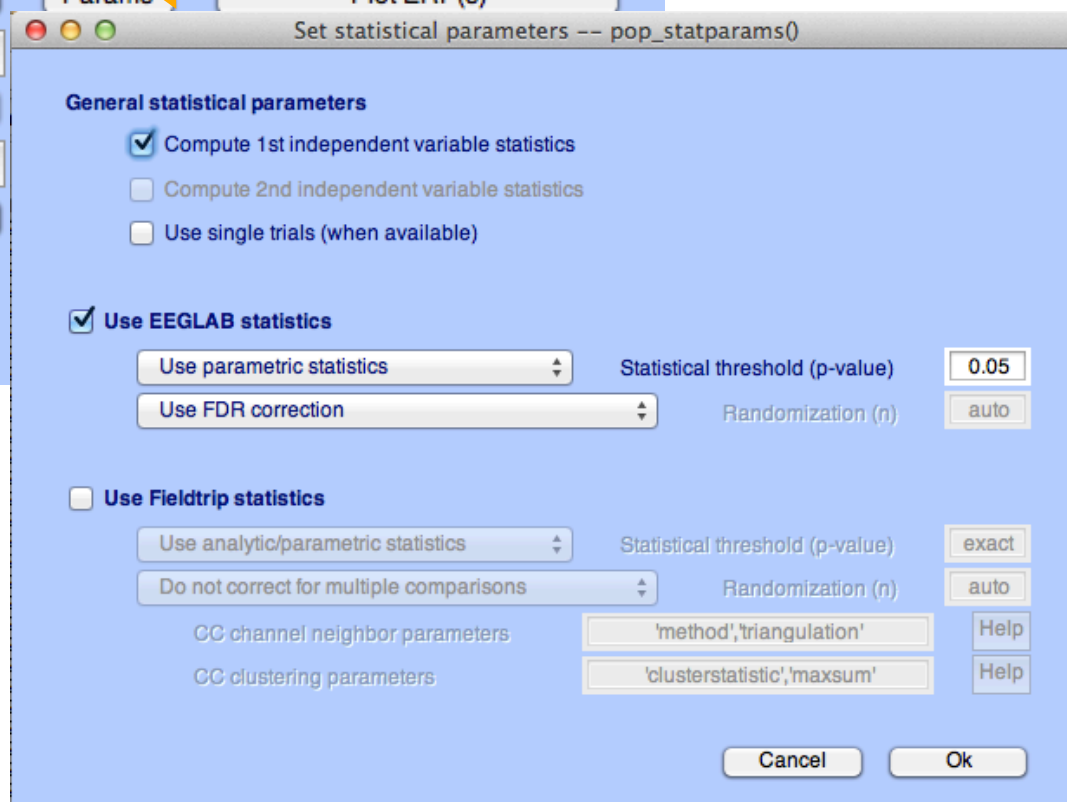
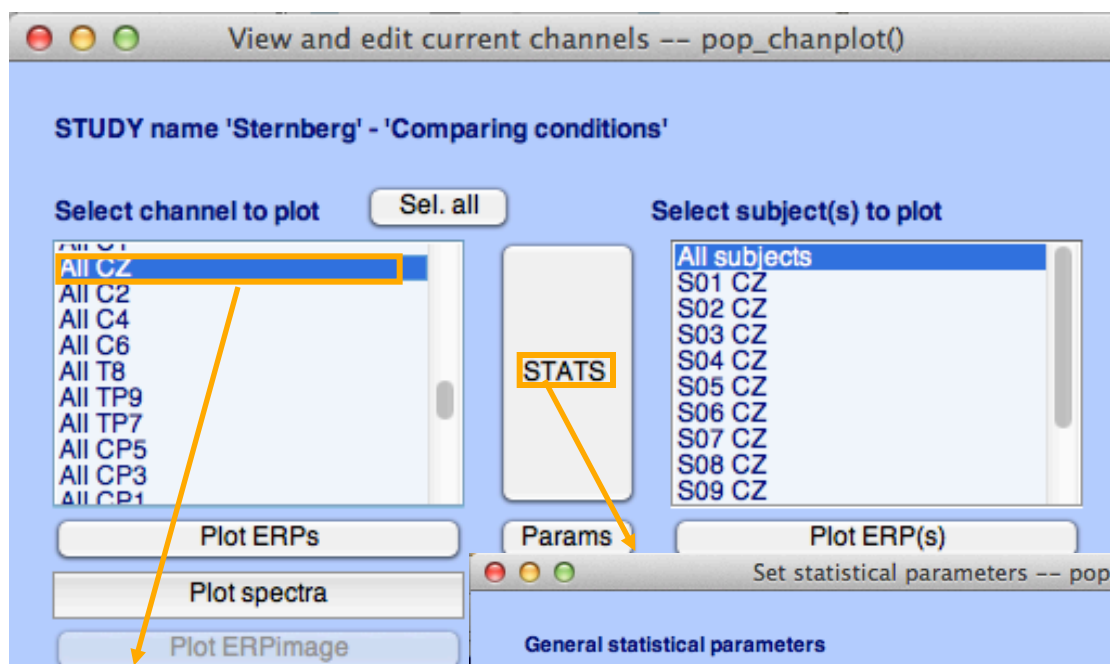
# Precompute data measures

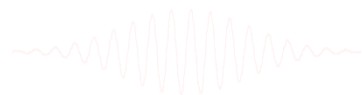
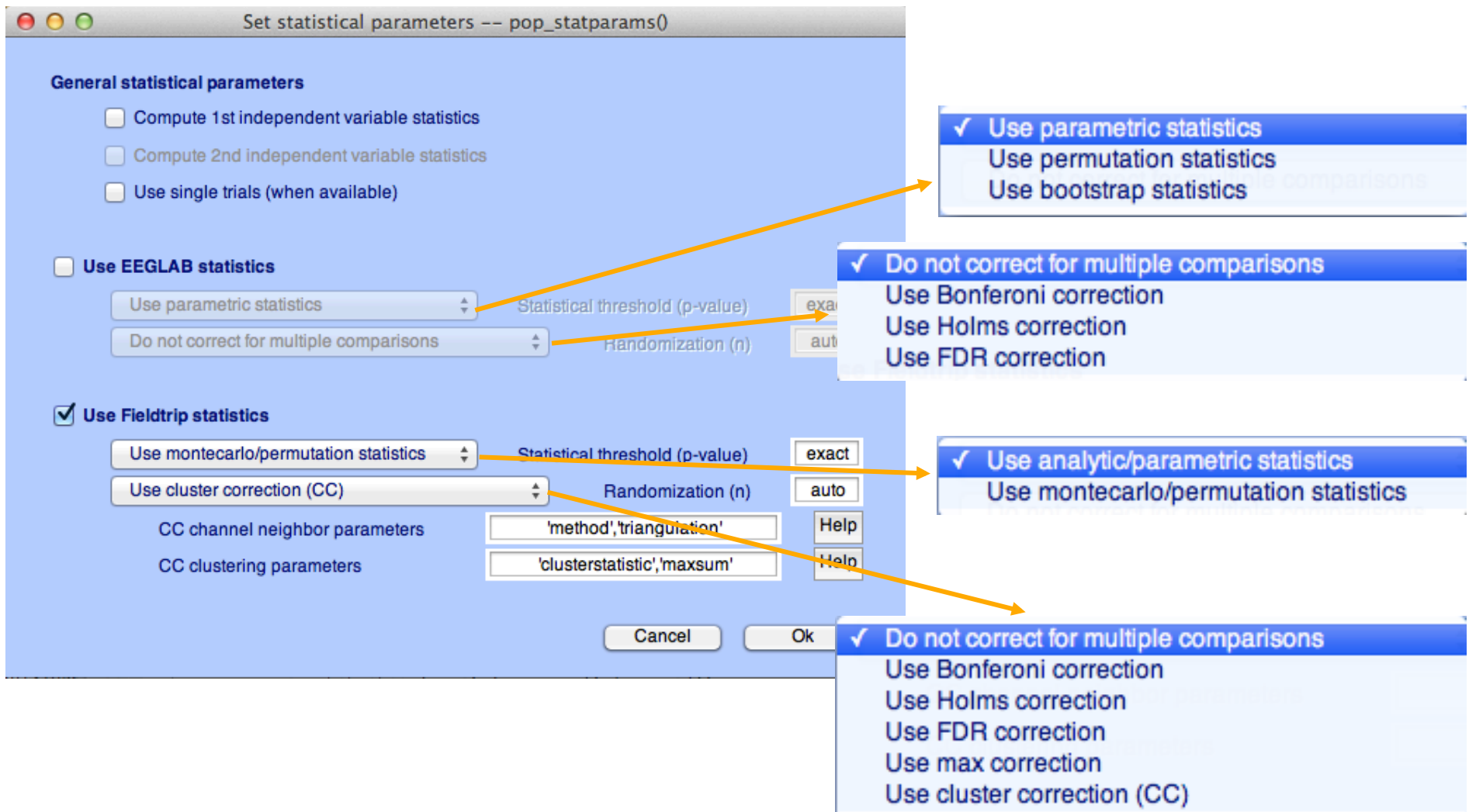






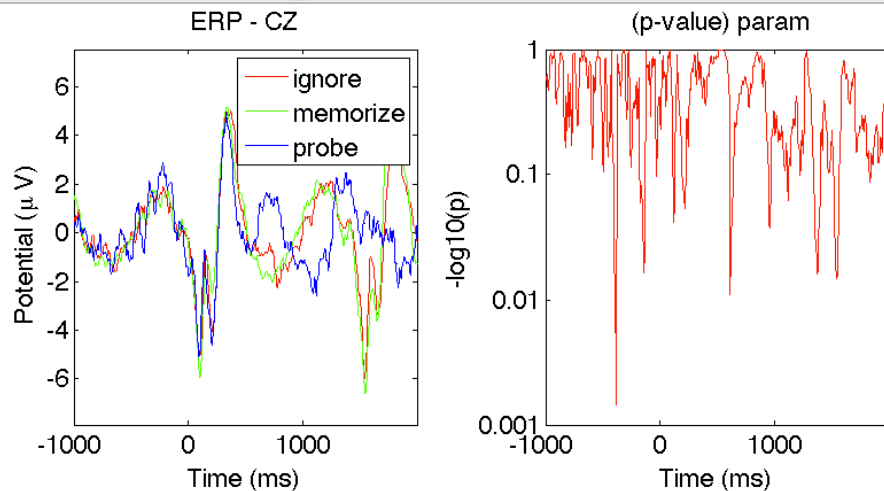
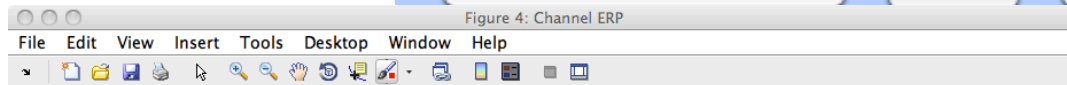
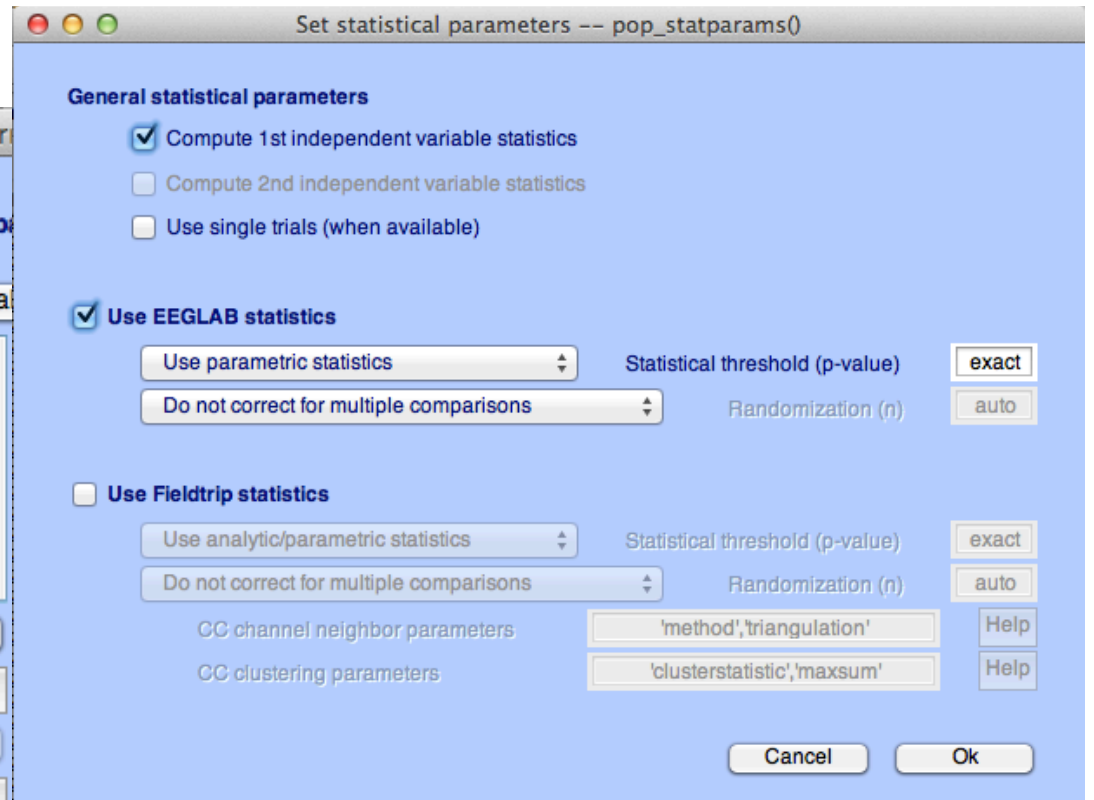
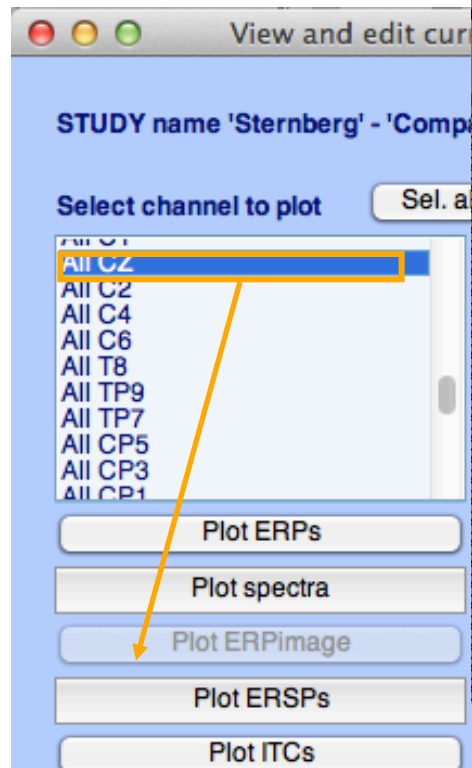




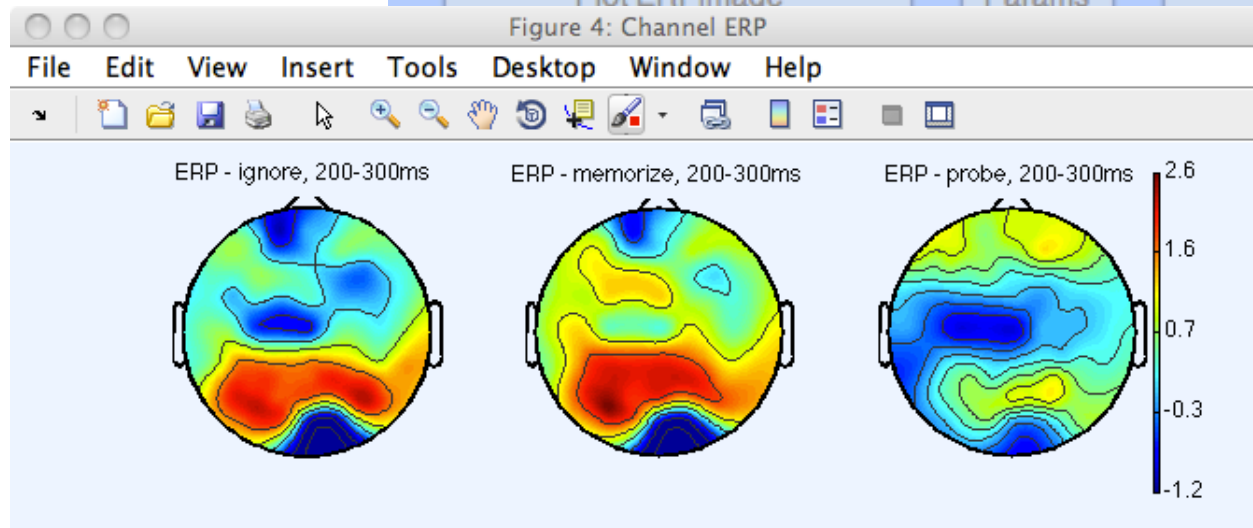
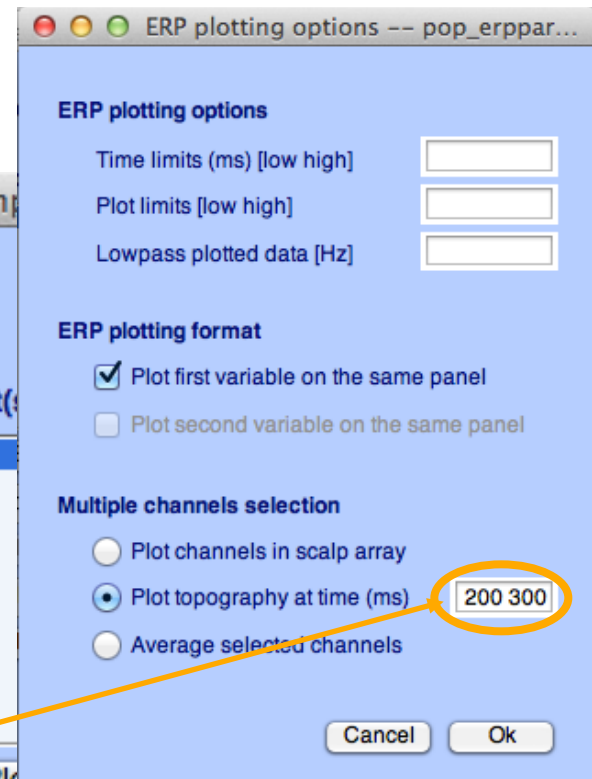
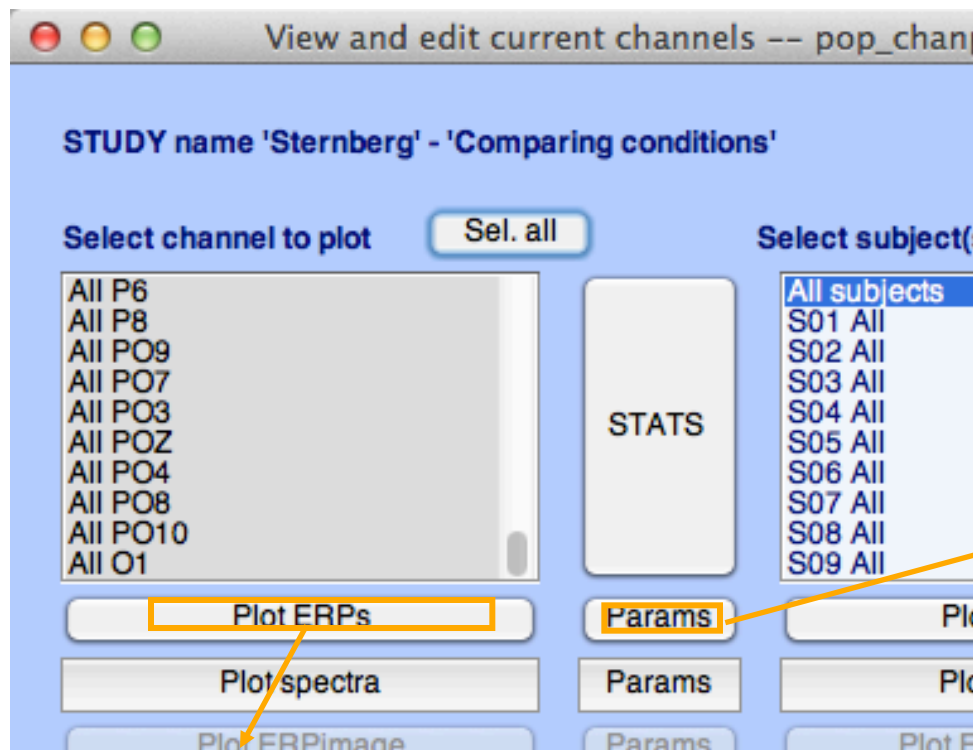


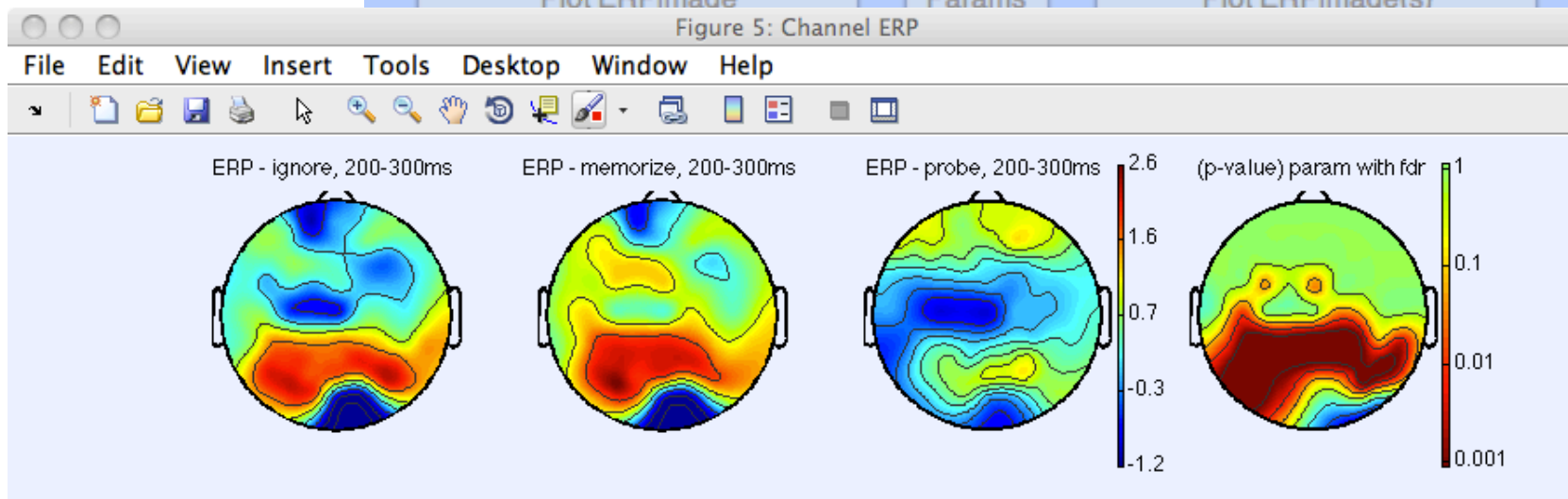
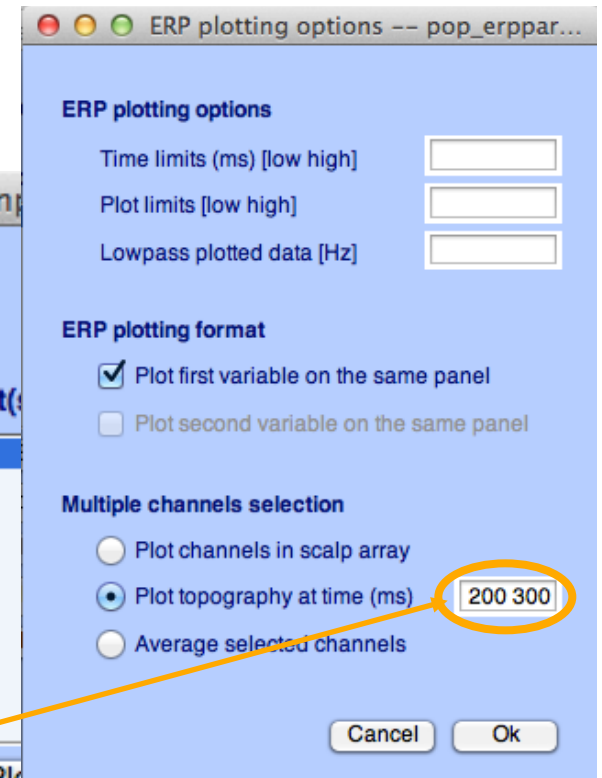
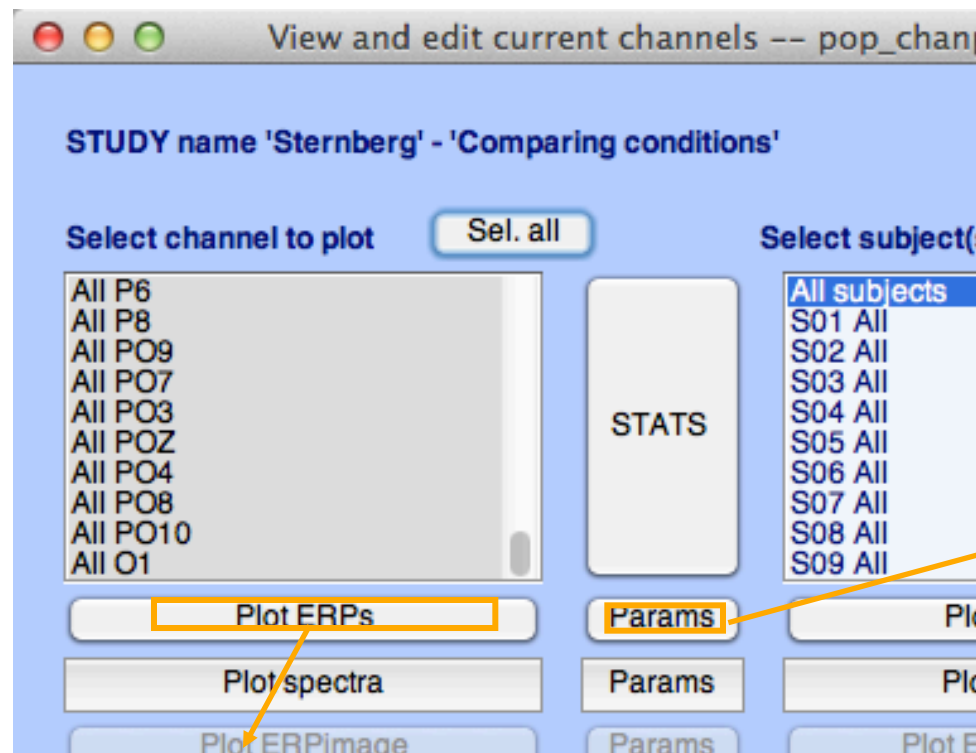
**std\_stat() function in EEGLAB**











# Computing Spectrum

Select and compute component measures for later clustering -- pop\_precomp()

**Pre-compute channel measures for STUDY 'Sternberg' - 'STUDY.design 1'**

**Channel list (default:all)**  ...

☒ Spherical interpolation of missing channels (performed after optional ICA removal below)

☐ Remove ICA artifactual components pre-tagged in each dataset

☐ Remove artifactual ICA cluster or clusters (hold shift key)

ParentCluster 1  
Cls 2  
Cls 3  
Cls 4

**List of measures to precompute**

☐ ERPs Baseline ([min max] in ms)

☒ Power spectrum Spectopo parameters

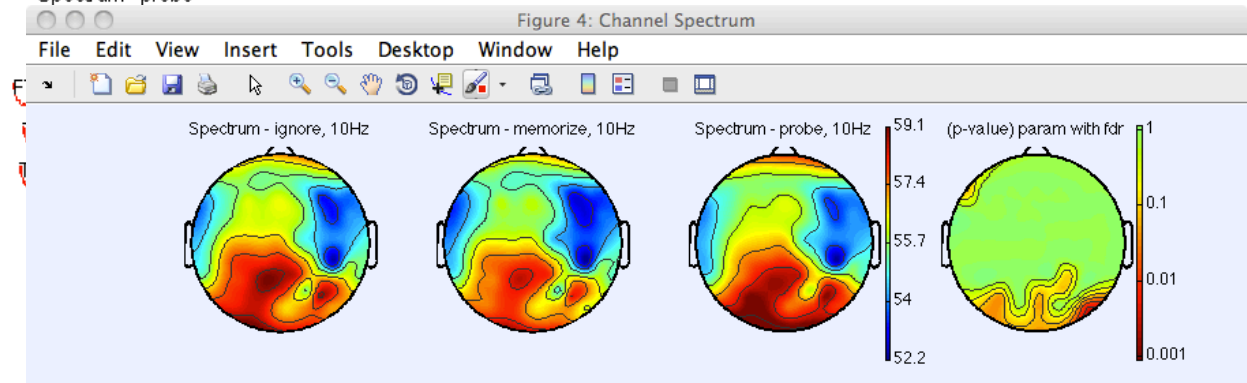
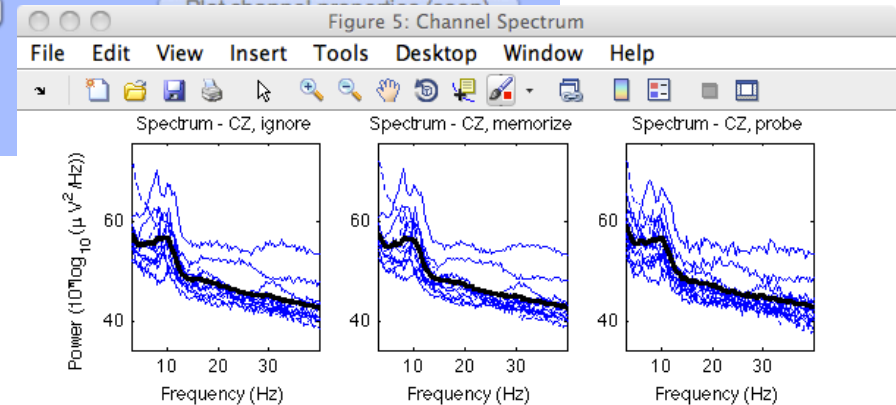
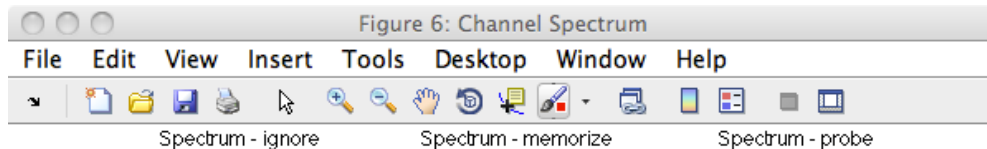
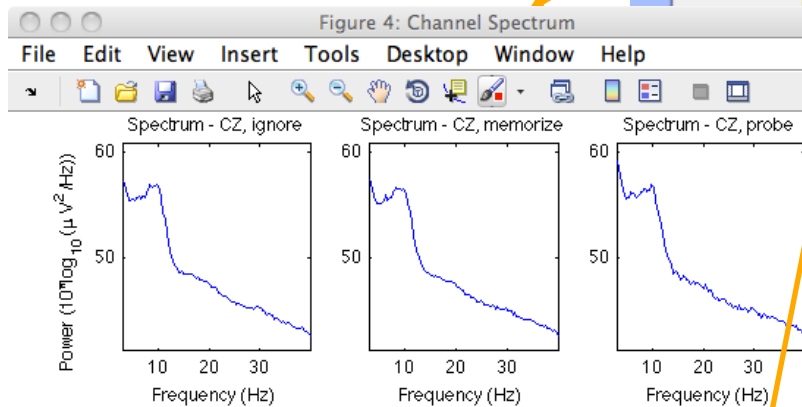
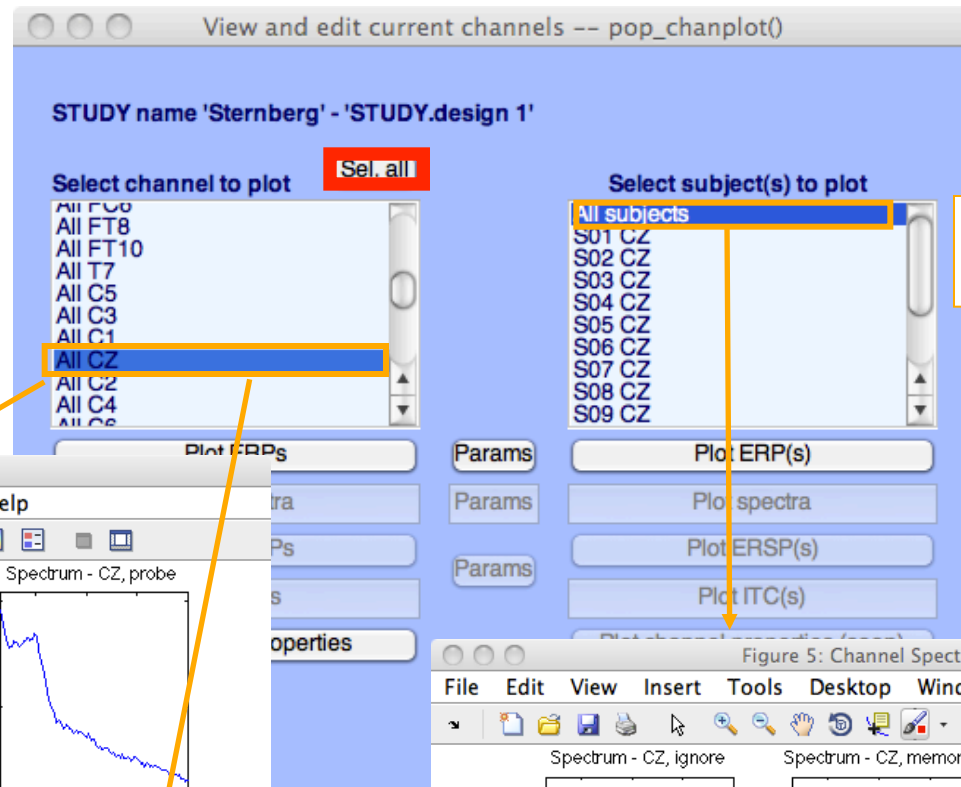
☐ ERSPs

☐ ITCs

☐ Save single-trial measures for single-trial statistics - requires disk space

☐ Recompute even if present on disk

Use 'timerange' option  
to select time range,  
see "help std\_spec"



# Computing ERSP

'cycles', [3 0.8], 'nfreqs', 50, 'ntimesout', 100

Select and compute component measures for later clustering -- pop\_precomp()

Pre-compute channel measures for STUDY 'Sternberg' - 'Design 2'

Channel list (default:all)  ...

☒ Spherical interpolation of missing channels (performed after optional ICA removal below)

☐ Remove ICA artifactual components pre-tagged in each dataset

☐ Remove artifactual ICA cluster or clusters (hold shift key)

ParentCluster 1  
Cls 2  
Cls 3  
Cls 4

List of measures to precompute

☐ ERPs Baseline ([min max] in ms)

☐ Power spectrum Spectopo parameters  Test

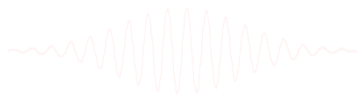
☒ ERSPs  Time/req. parameters Test

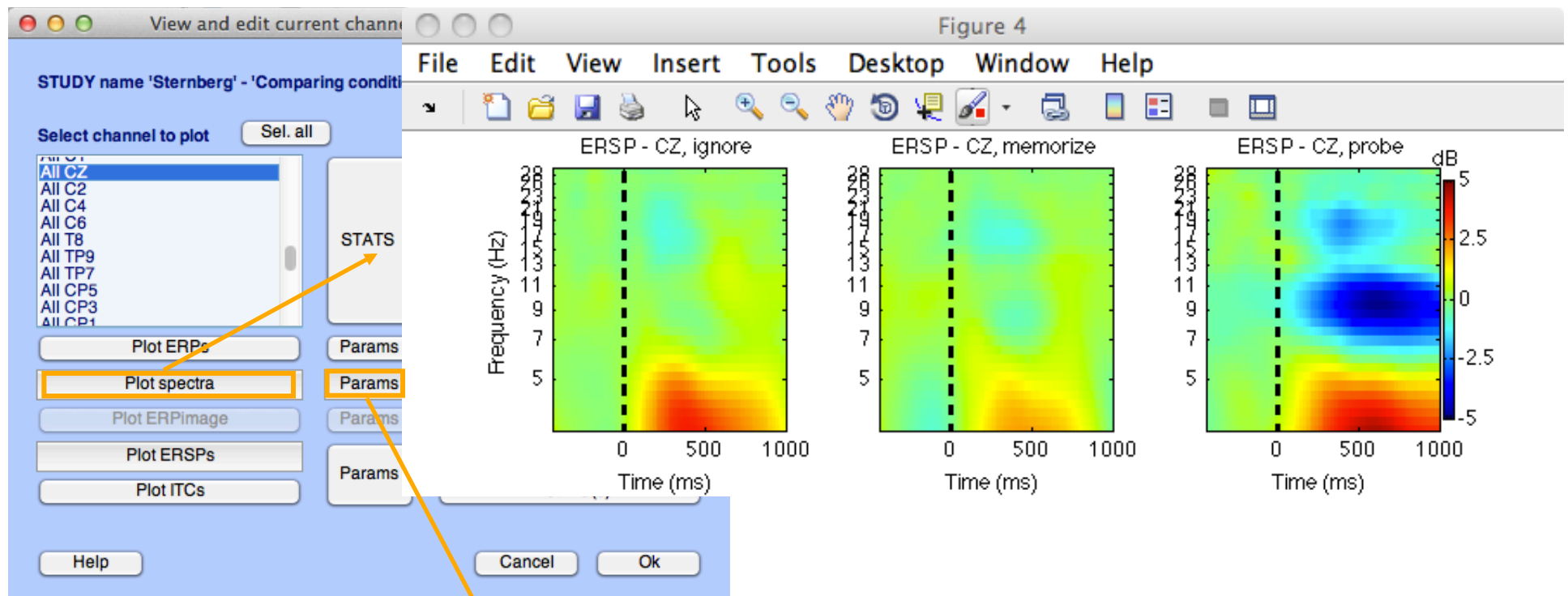
☐ ITCs

☐ Save single-trial measures for single-trial statistics - requires disk space

☐ Recompute even if present on disk

Help Cancel Ok





Set ERSP/ITC plotting parameters -- pop\_erspparams()

**ERSP/ITC plotting options**

Time range in ms [Low High] -500 1000

Freq. range in Hz [Low High] 3 30

Power limits in dB [Low High]

☐ Compute common ERSP baseline (assumes additive baseline)

Plot scalp map at time [ms]

Plot scalp map at freq. [Hz]

ITC limit (0-1) [High]

Cancel

Ok



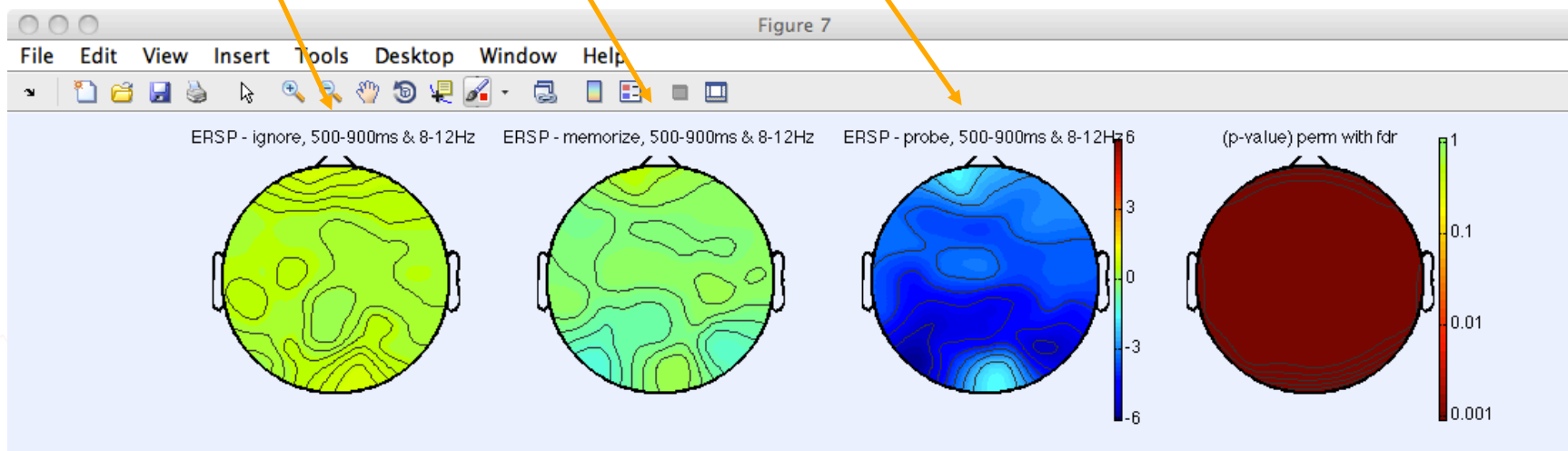
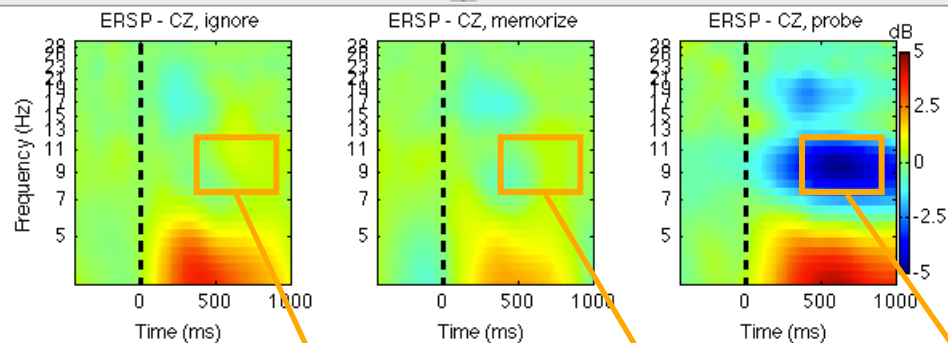
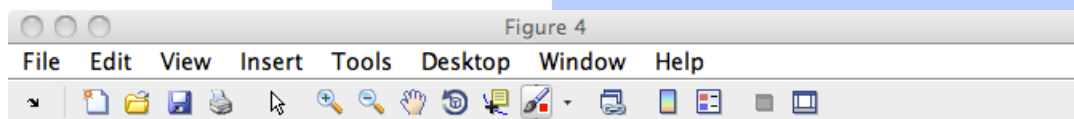
Set ERSP/ITC plotting parameters -- pop\_erspparams()

**ERSP/ITC plotting options**

Time range in ms [Low High]	-500 1000	Plot scalp map at time [ms]	500 900
Freq. range in Hz [Low High]	3 30	Plot scalp map at freq. [Hz]	8 12
Power limits in dB [Low High]		ITC limit (0-1) [High]	

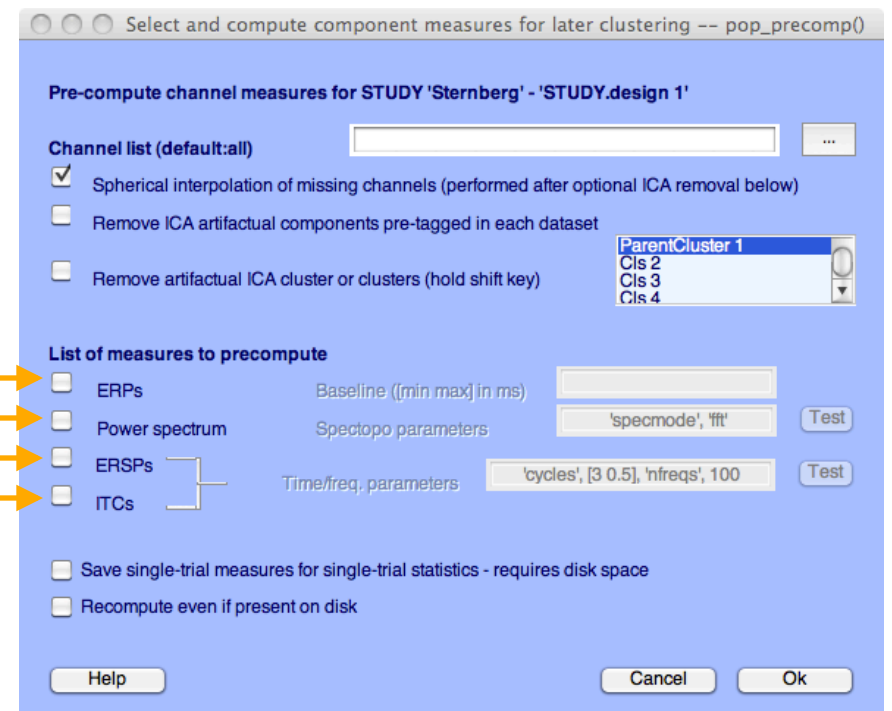
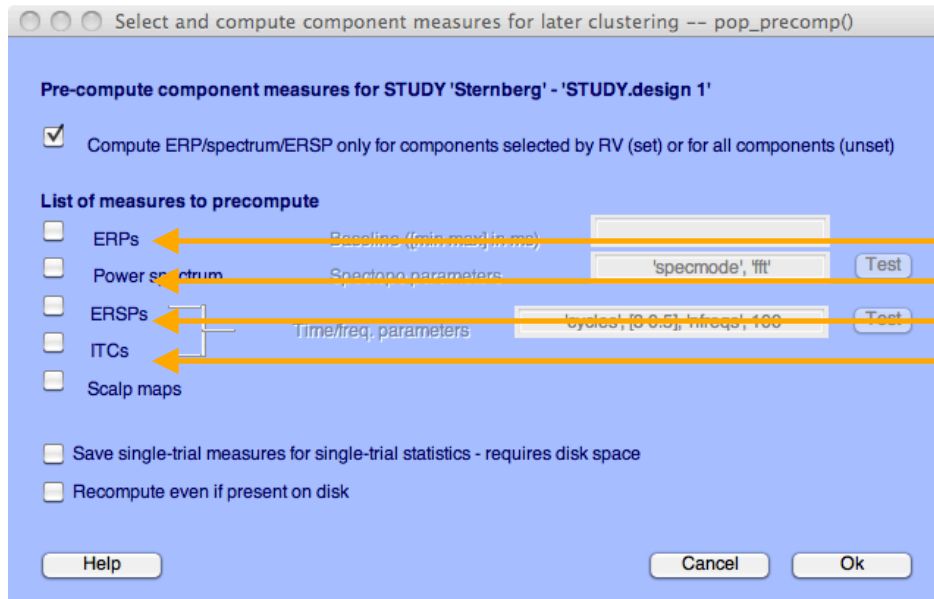
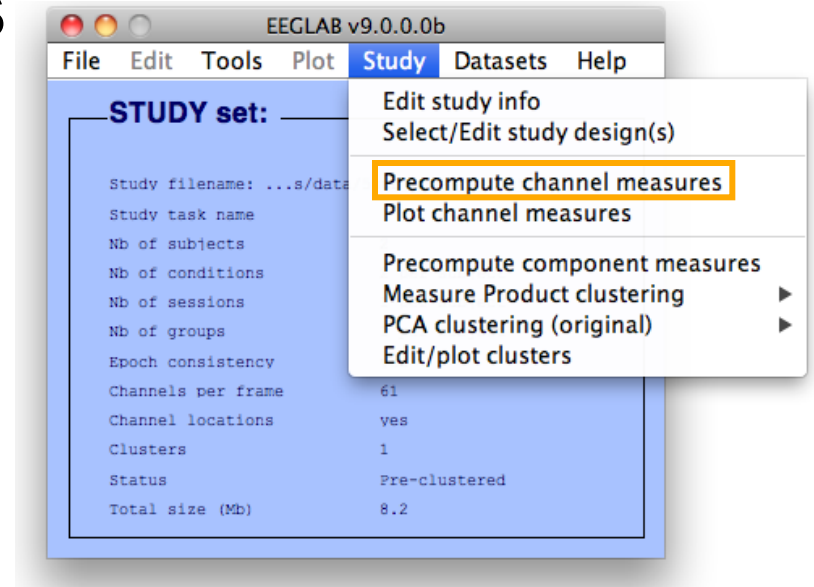
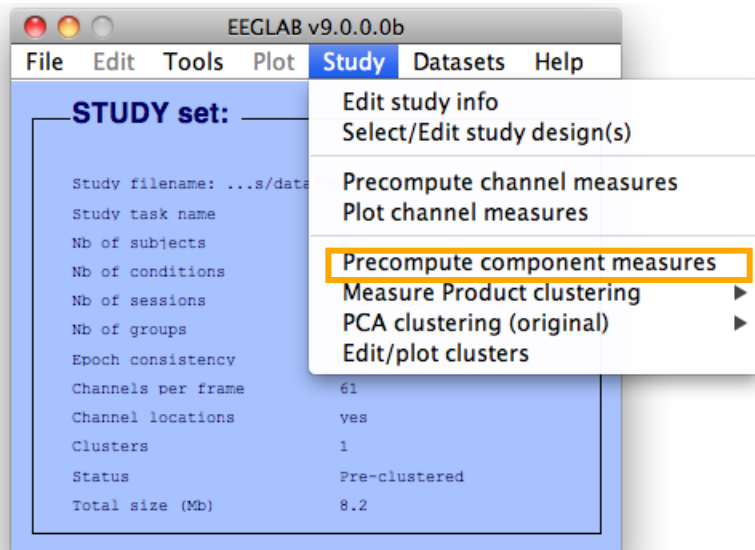
☒ Compute common ERSP baseline (assumes additive baseline)

Cancel Ok

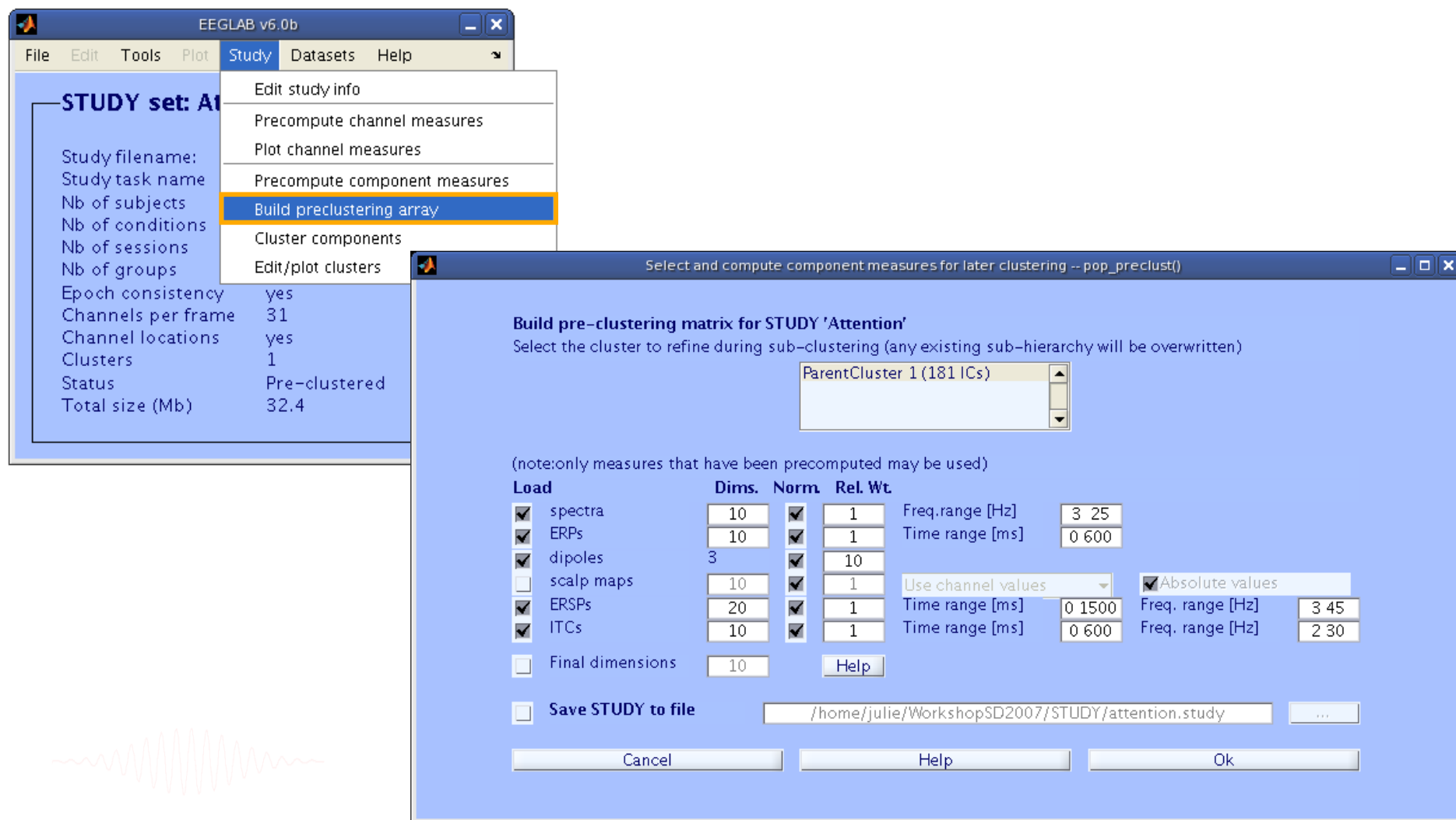




## 2. Pre-compute measures



# 3. Cluster components



The image displays the EEGLAB v6.0b software interface. The 'Study' menu is open, showing options like 'Edit study info', 'Precompute channel measures', 'Plot channel measures', 'Precompute component measures', 'Build preclustering array' (highlighted), 'Cluster components', and 'Edit/plot clusters'. The 'STUDY set: Attention' panel shows study details: Study filename, Study task name, Nb of subjects, Nb of conditions, Nb of sessions, Nb of groups, Epoch consistency (yes), Channels per frame (31), Channel locations (yes), Clusters (1), Status (Pre-clustered), and Total size (Mb) (32.4).

The 'Select and compute component measures for later clustering -- pop\_preclust()' dialog box is open, titled 'Build pre-clustering matrix for STUDY 'Attention''. It prompts the user to 'Select the cluster to refine during sub-clustering (any existing sub-hierarchy will be overwritten)'. The selected cluster is 'ParentCluster 1 (181 ICs)'.

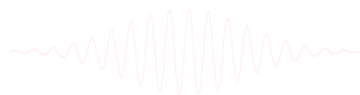
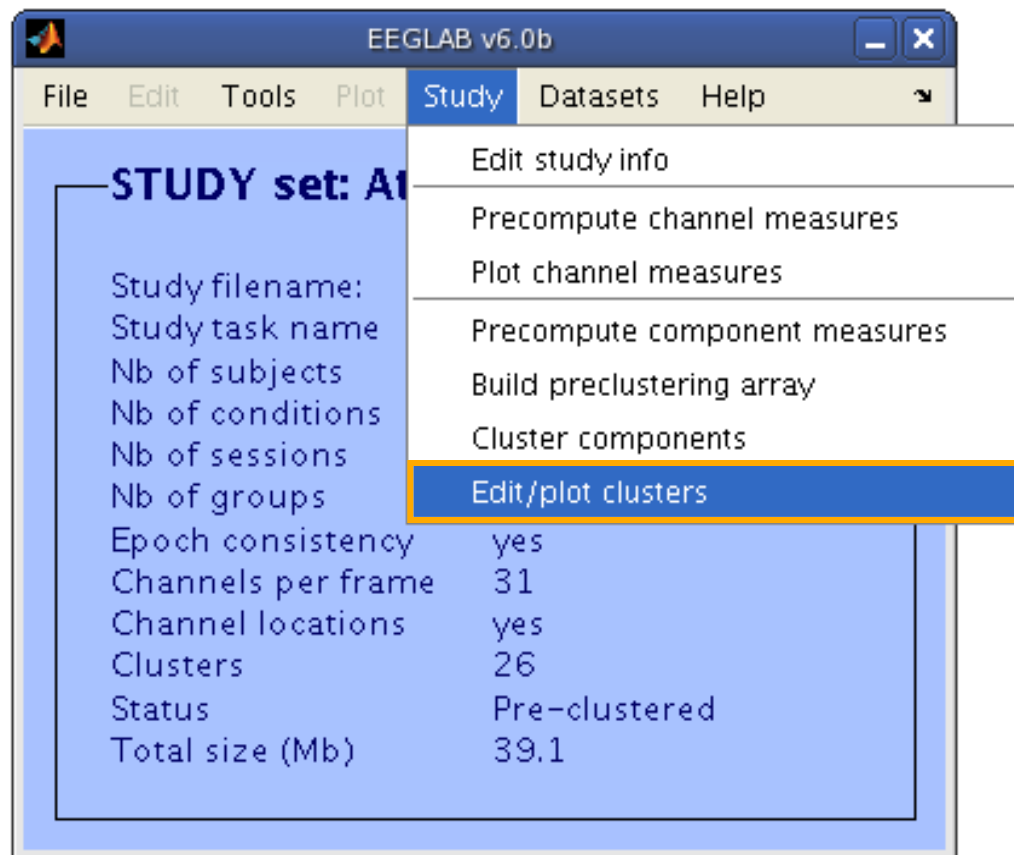
(note: only measures that have been precomputed may be used)

Load	Dims.	Norm.	Rel. Wt.	
<input checked="" type="checkbox"/> spectra	10	<input checked="" type="checkbox"/>	1	Freq. range [Hz] 3 25
<input checked="" type="checkbox"/> ERPs	10	<input checked="" type="checkbox"/>	1	Time range [ms] 0 600
<input checked="" type="checkbox"/> dipoles	3	<input checked="" type="checkbox"/>	10	
<input type="checkbox"/> scalp maps	10	<input checked="" type="checkbox"/>	1	Use channel values <input checked="" type="checkbox"/> Absolute values
<input checked="" type="checkbox"/> ERSPs	20	<input checked="" type="checkbox"/>	1	Time range [ms] 0 1500 Freq. range [Hz] 3 45
<input checked="" type="checkbox"/> ITCs	10	<input checked="" type="checkbox"/>	1	Time range [ms] 0 600 Freq. range [Hz] 2 30
<input type="checkbox"/> Final dimensions	10			

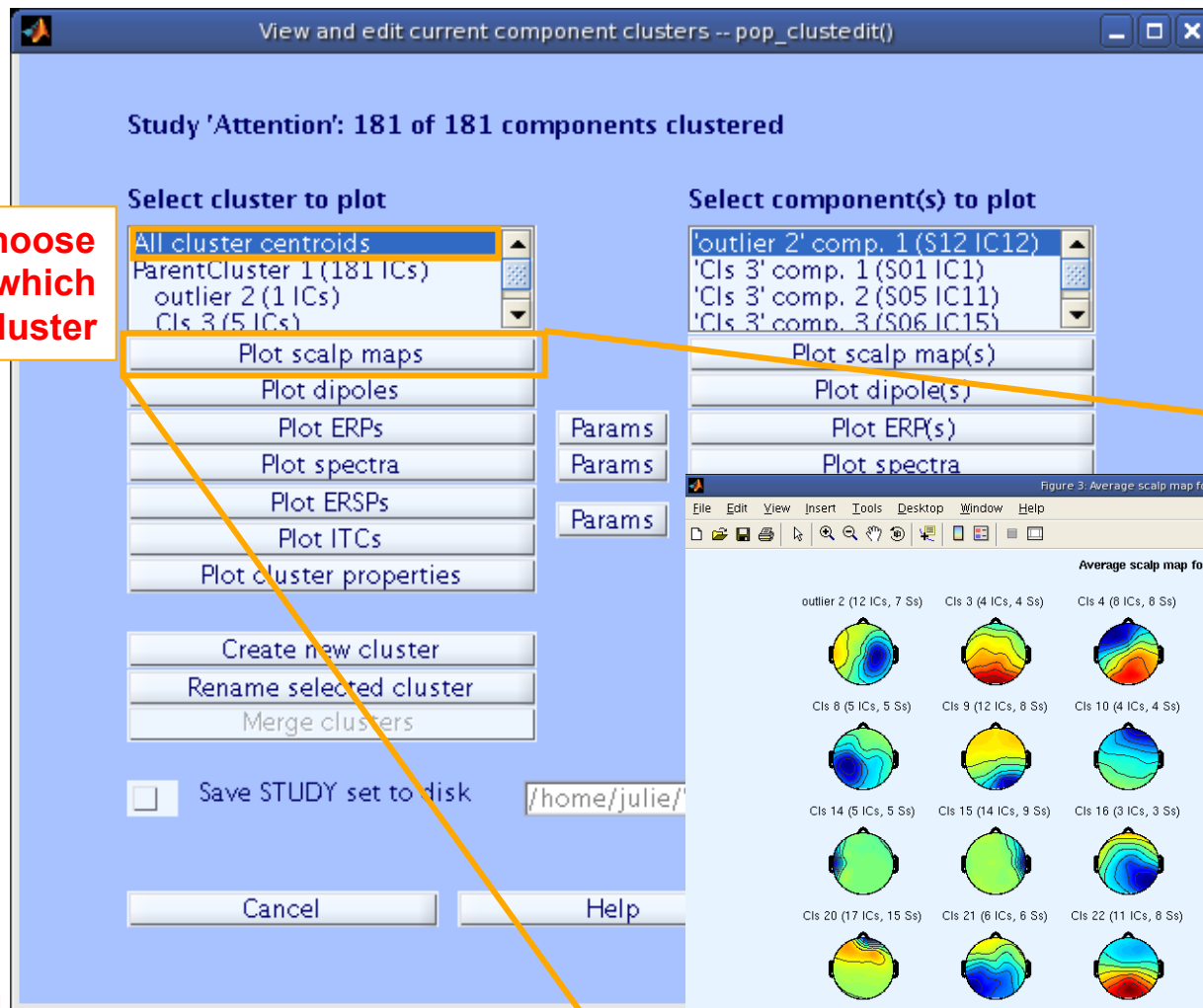
☐ Save STUDY to file /home/julie/WorkshopSD2007/STUDY/attention.study ...

Buttons: Cancel, Help, Ok

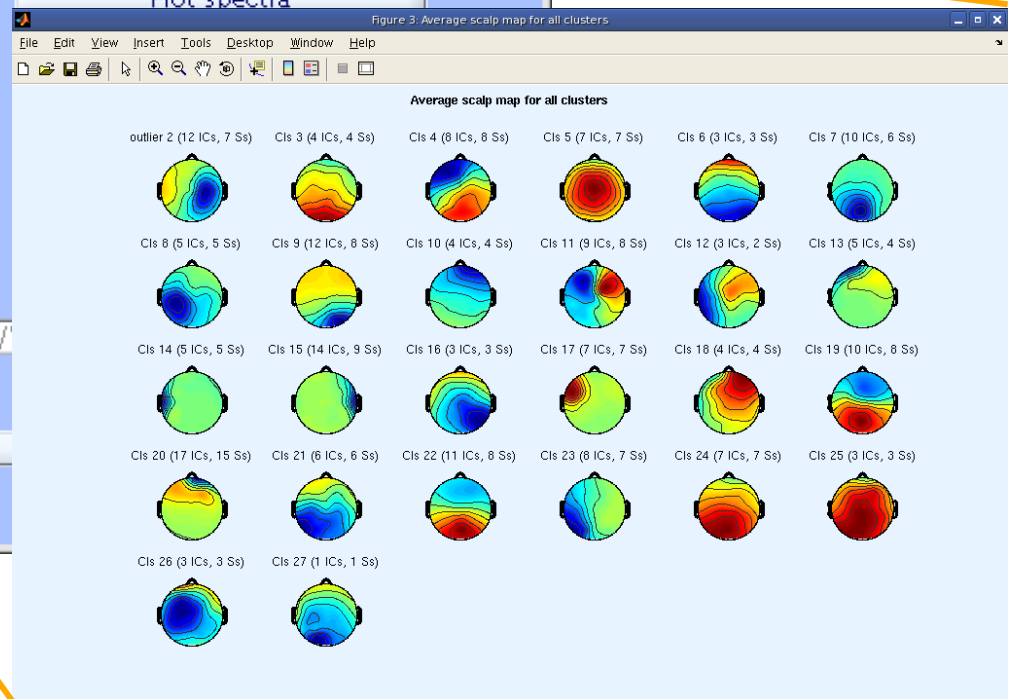
# View and edit clusters



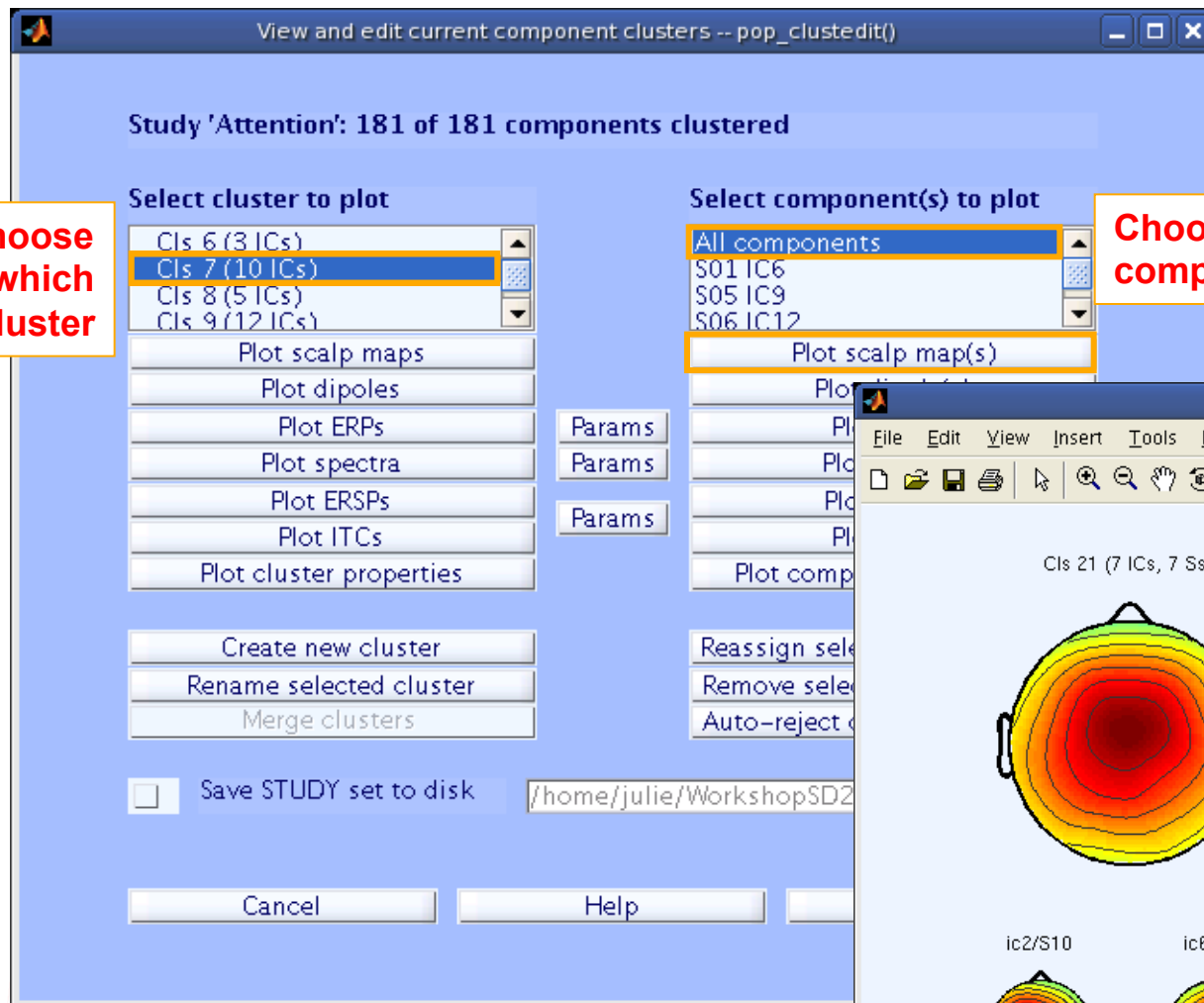
# Plot cluster data



Plot mean scalp maps for easy reference

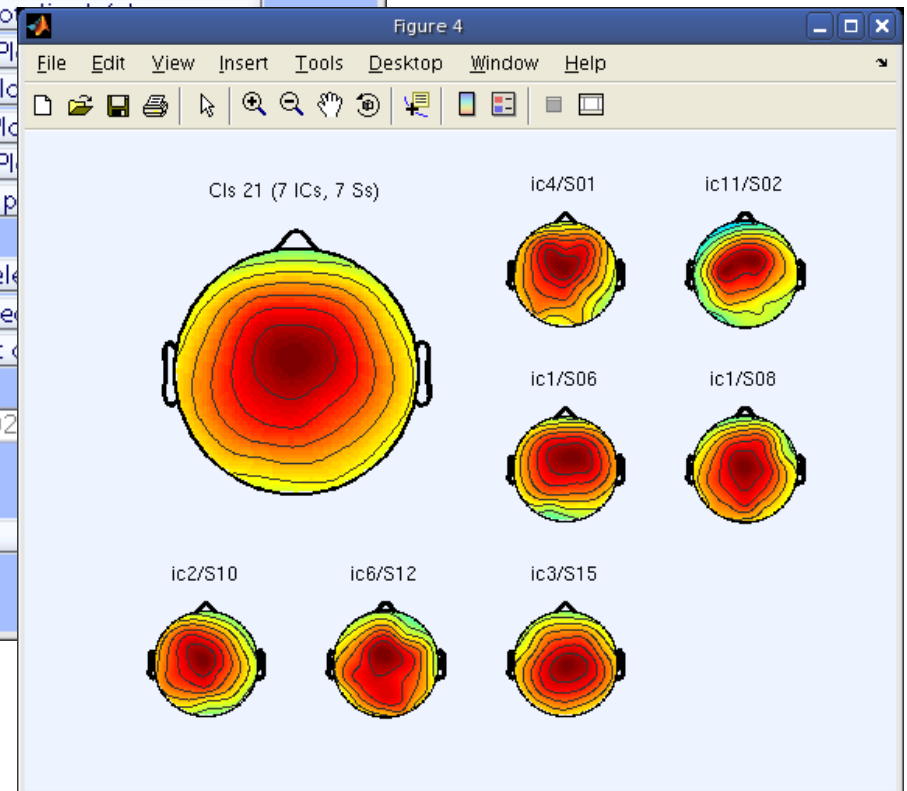


# Plot cluster data

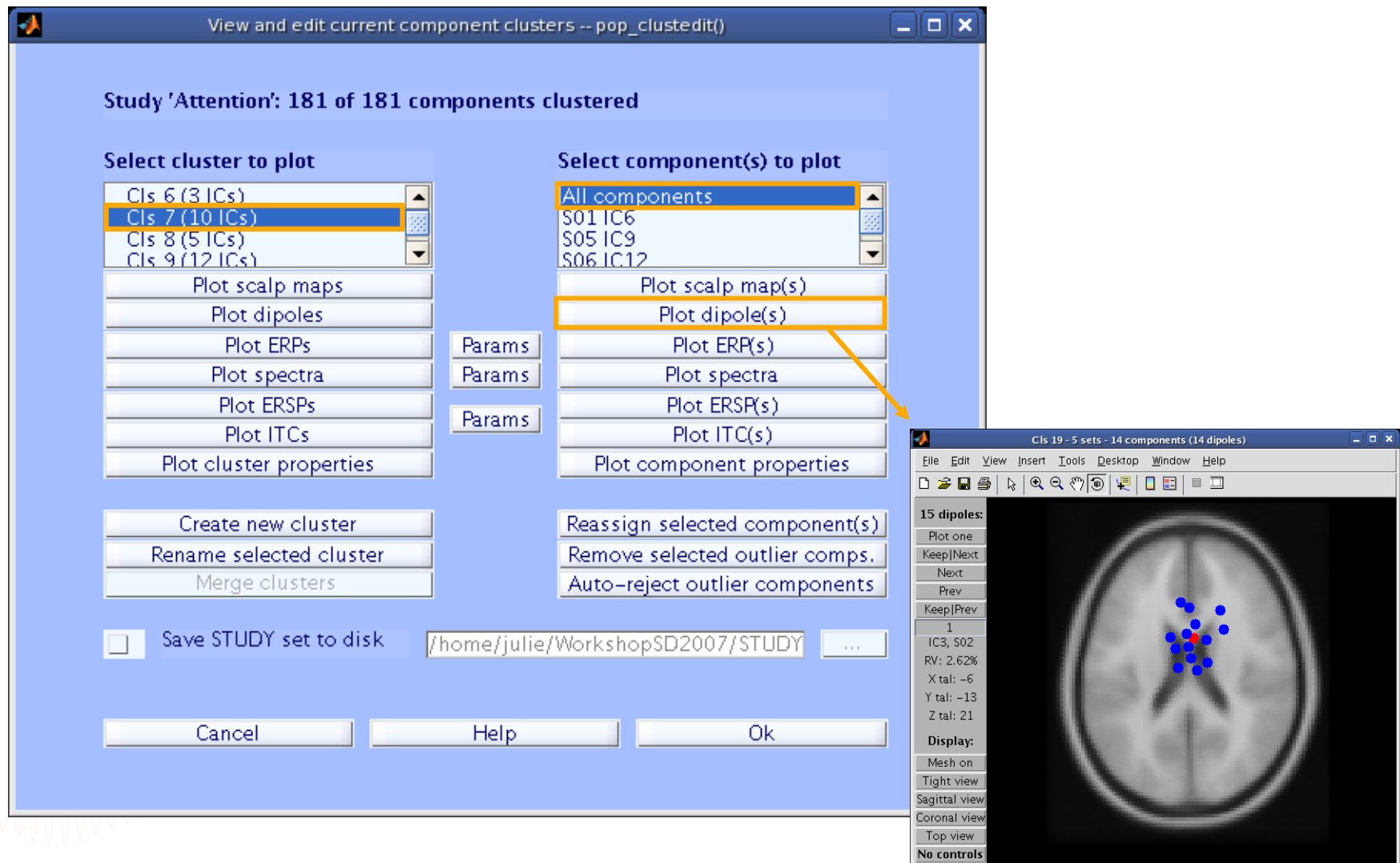


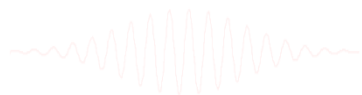
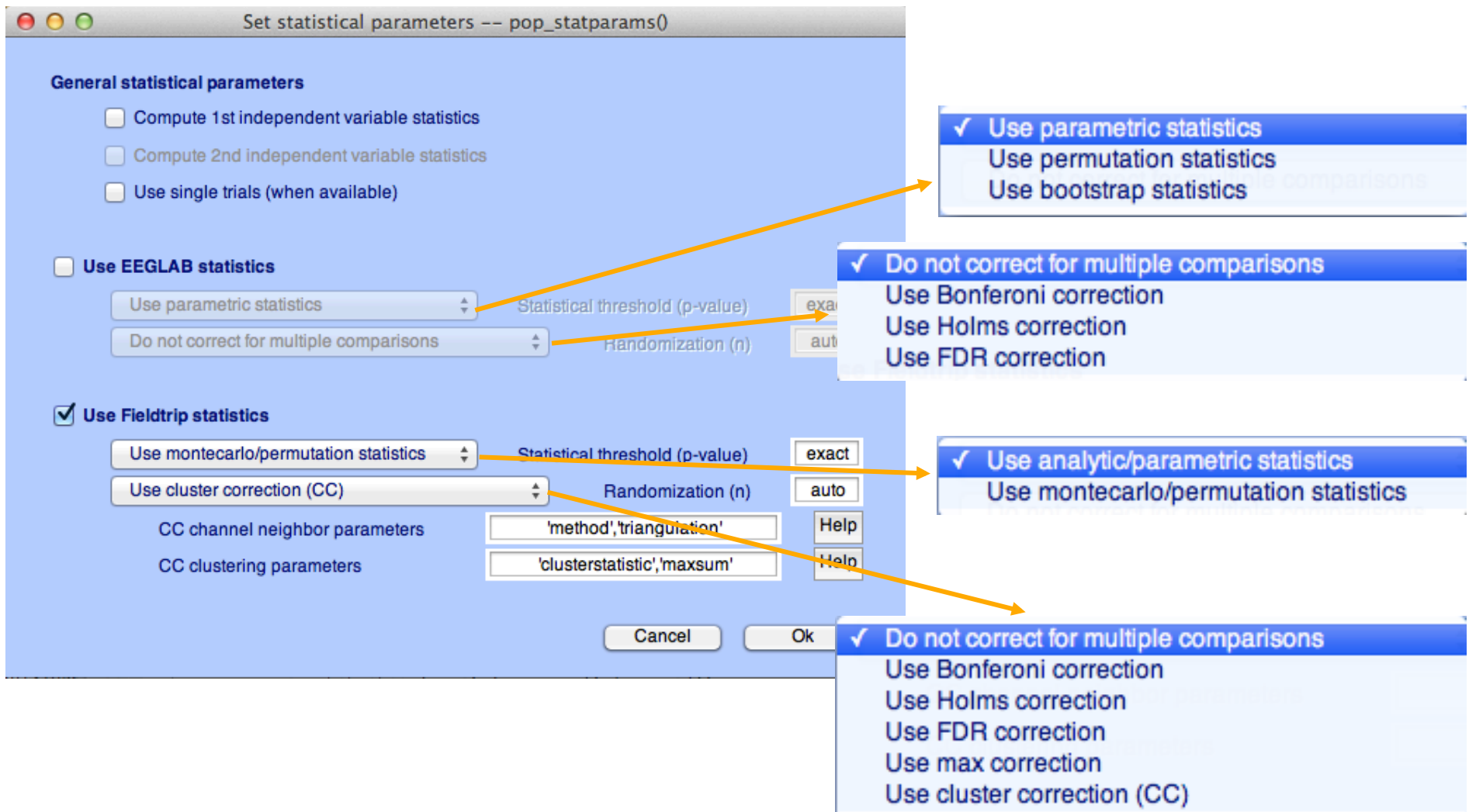
Choose  
which  
cluster

Choose which  
components



# Plot cluster data





**std\_stat() function in EEGLAB**



# statcond function in EEGLAB

```
a = { rand(1,10) rand(1,10)+0.5 }; % pseudo 'paired' data vectors
```

```
[t df pvals] = statcond(a , 'mode', 'perm'); % perform paired t-test  
pvals = 5.2807e-04 % standard t-test probability value
```

```
% Note: for different rand() outputs, results will differ.
```

```
[t df pvals surog] = statcond(a, 'mode', 'perm', 'naccu', 2000);  
pvals = 0.0065 % nonparametric t-test using 2000 permuted data sets
```

```
a = { rand(2,11) rand(2,10) rand(2,12)+0.5 };
```

```
[F df pvals] = statcond(a , 'mode', 'perm'); % perform an unpaired ANOVA
```

```
pvals =
```

```
0.00025 % p-values for difference between columns
```

```
0.00002 % for each data row
```

# statcond function in EEGLAB

```
a = { rand(3,4,10) rand(3,4,10) rand(3,4,10); ...  
      rand(3,4,10) rand(3,4,10) rand(3,4,10)+0.5 };
```

```
% pseudo (2,3)-condition data array, each entry containing  
% ten (3,4) data matrices
```

```
[F df pvals] = statcond(a , 'mode', 'perm');  
                % paired 2-way ANOVA
```

```
% Output:
```

```
pvals{1} % a (3,4) matrix of p-values; effects across columns
```

```
pvals{2} % a (3,4) matrix of p-values; effects across rows
```

```
pvals{3} % a (3,4) matrix of p-values; interaction effects across  
          rows and columns
```

# Exercise

- Experiment with the statcond function
  - Create 2 random vectors of values
  - Add “signal” to one of the variable
  - Use statcond and compare permutation and parametric results
  - Repeat 100 times and plot the histogram of p-values
- Experiment with STUDY statistics
  - Load the Stern STUDY
  - Look at significant difference between probe and memorize in component clusters (time-frequency plot, ERSP) using the cluster method (Fieldtrip – statistics)

