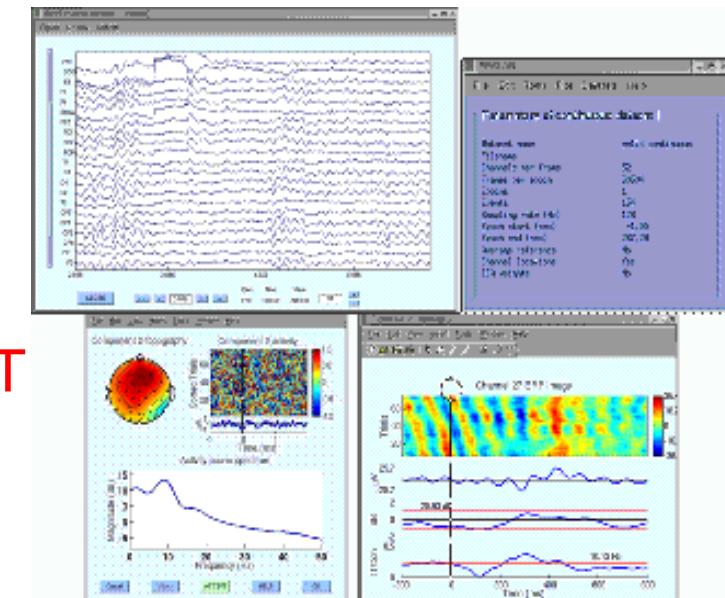


# The EEGLAB software



- Collection of over 300 functions (50000 lines of code)
- About 50000 download over the past 6 years
- About 3500 users on the discussion list and 6500 on the diffusion list
- NIH funding (2003-)



<http://sccn.ucsd.edu/wiki/EEGLAB09ASPET>



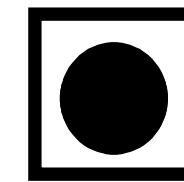
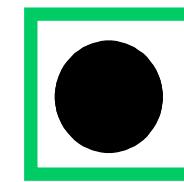
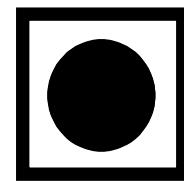
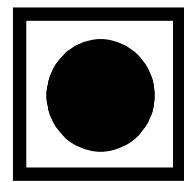
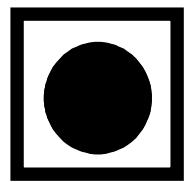
# Pros/Cons of Matlab based open source



- Pros
  - Easy to program, highly modular and extendable
  - Not dependent on any platform (64-bit)
  - Large community of users (latest development in signal processing research)
  - Cannot imagine more powerful scripting capabilities
- Cons
  - Matlab required
  - Large memory requirements
  - Matlab bugs, possible version differences, cross-platform compatibility problems



# Spatial visual attention task (STUDY)



+



# Publication using this data



- Delorme A, Westerfield M, Makeig S., **J Neurosci**. 2007 Oct 31;27(44):11949-59.
- Onton J, Westerfield M, Townsend J, Makeig S., **Neurosci Biobehav Rev**. 2006;30(6):808-22. Epub 2006 Aug 14. Review.
- Makeig S, Delorme A, Westerfield M, Jung TP, Townsend J, Courchesne E, Sejnowski TJ., **PLoS Biol**. 2004 Jun;2(6):e176. Epub 2004 Jun 15.
- Makeig S, Westerfield M, Jung TP, Enghoff S, Townsend J, Courchesne E, Sejnowski TJ. **Science**. 2002 Jan 25;295(5555):690-4. **Science** 2002 Feb 22;295(5559):1466.
- Jung TP, Makeig S, Westerfield M, Townsend J, Courchesne E, Sejnowski TJ., **Hum Brain Mapp**. 2001 Nov;14(3):166-85.
- Townsend J, Westerfield M, Leaver E, Makeig S, Jung T, Pierce K, Courchesne E., **Brain Res Cogn Brain Res**. 2001 Mar;11(1):127-45.
- Jung TP, Makeig S, Westerfield M, Townsend J, Courchesne E, Sejnowski TJ., **Clin Neurophysiol**. 2000 Oct;111(10):1745-58.
- Makeig S, Westerfield M, Townsend J, Jung TP, Courchesne E, Sejnowski TJ., **Philos Trans R Soc Lond B Biol Sci**. 1999 Jul 29;354(1387):1135-44.
- Makeig S, Westerfield M, Jung TP, Covington J, Townsend J, Sejnowski TJ, Courchesne E., **J Neurosci**. 1999 Apr 1;19(7):2665-80.

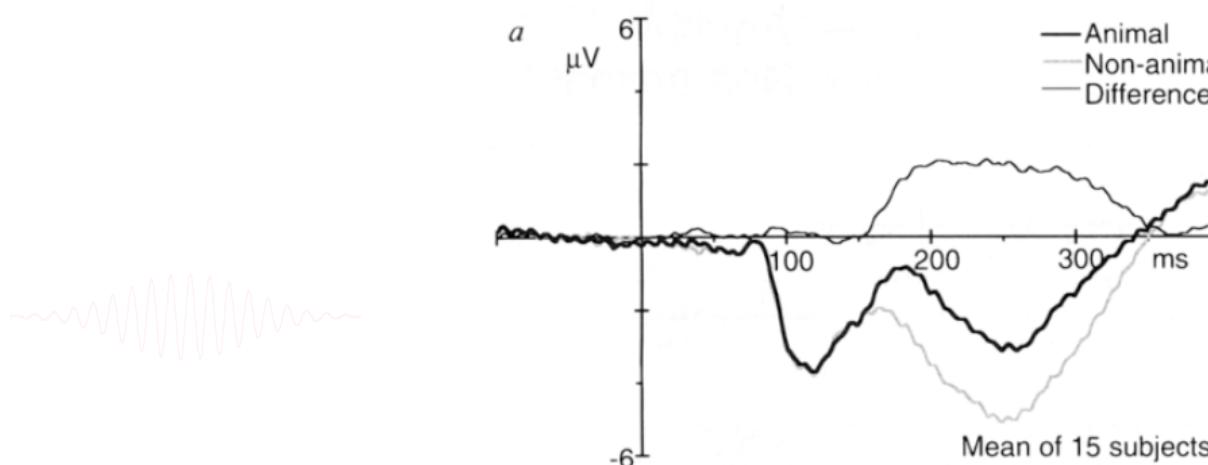


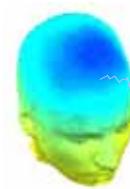


# Publication using this paradigm

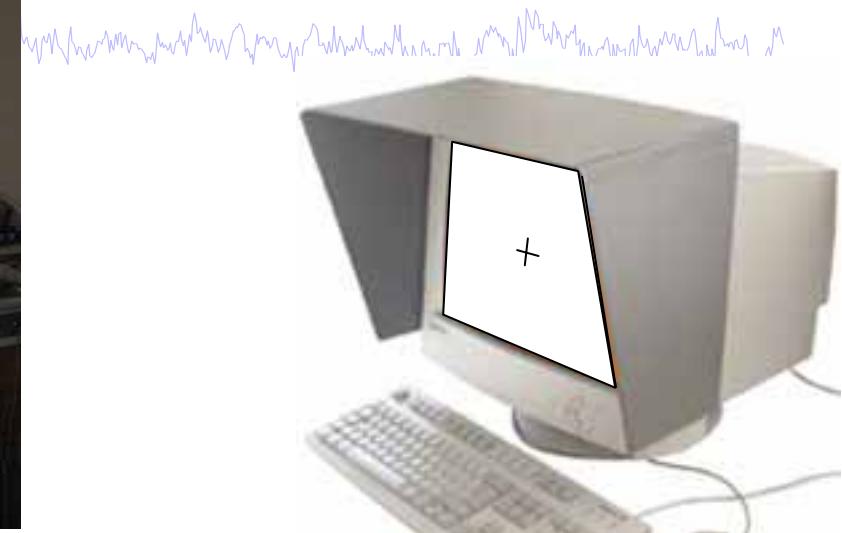
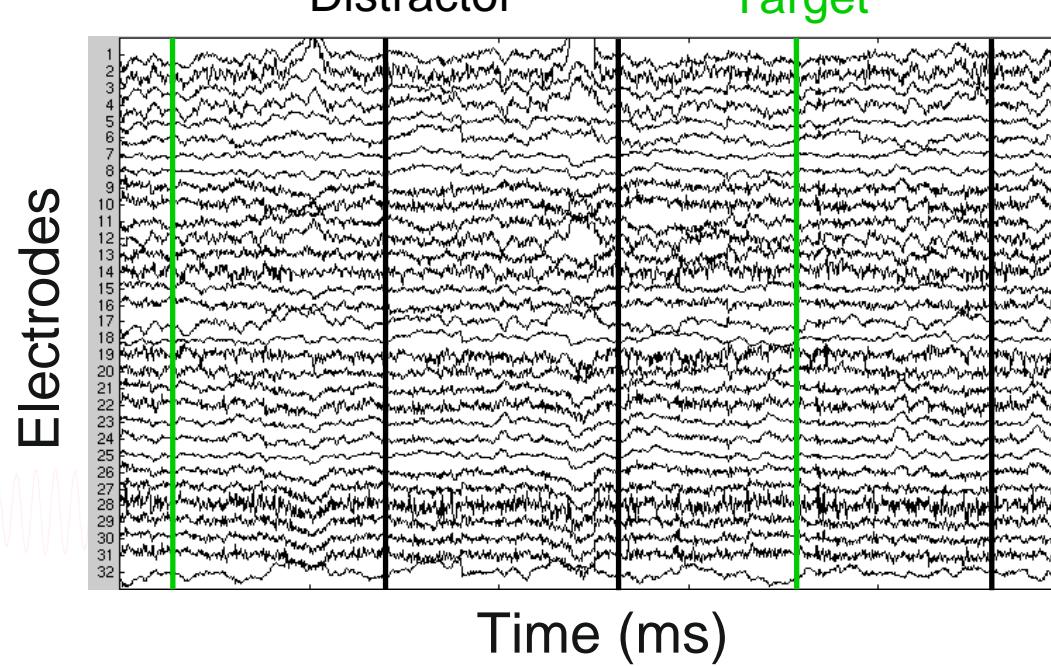


- Thorpe S, Fize D, Marlot C. 1996. Speed of processing in the human visual system. Nature 381: 520-2
- Fabre-Thorpe, M., Delorme, A., Marlot, C., & Thorpe, S.J. (2000). A limit to the speed of processing in Ultra-Rapid Visual Categorization of novel natural scenes. Journal of Cognitive Neuroscience. 13(2), 171-180.
- Delorme, A., Makeig, S., Fabre-Thorpe, M. Sejnowski, T. (2002) From Single-trials EEG to Brain Area Dynamics, Neurocomputing, 44-46, 1057-1064.
- Delorme, A., Rousselet, G., Mace, M., Fabre-Thorpe M. (2004) Interaction of Bottom-up and Top-down processing in the fast visual analysis of natural scenes. Cognitive Brain Research, 19(2), 103-113.
- ...



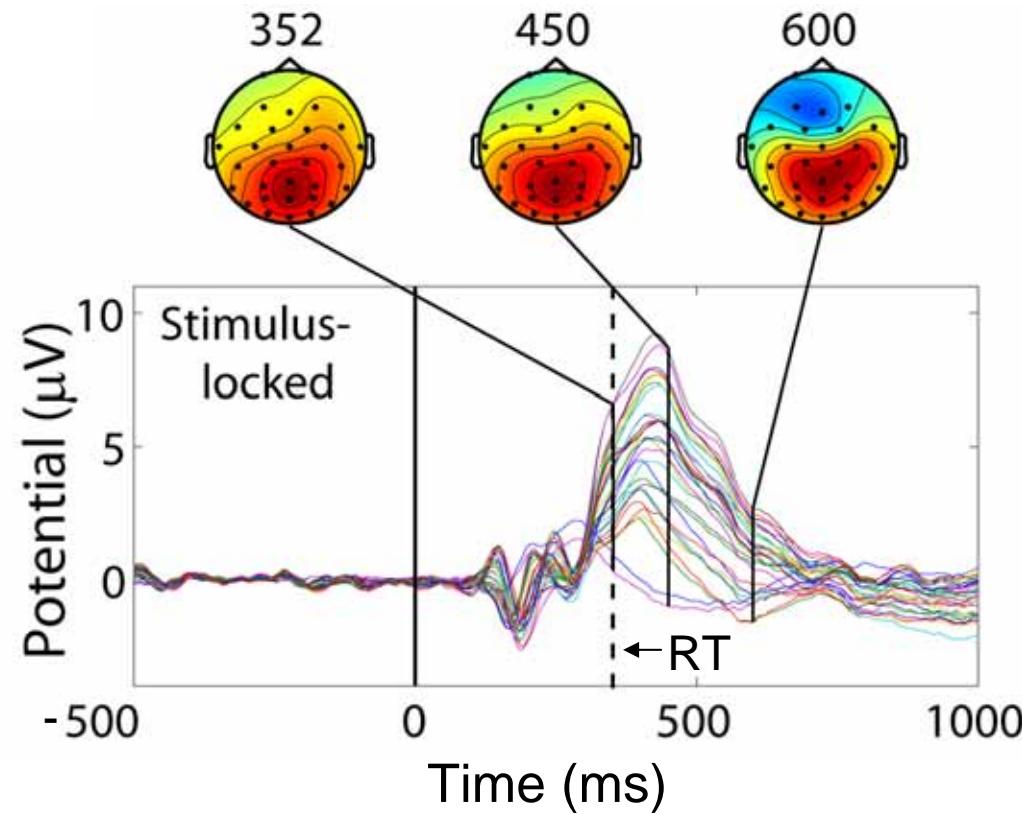


Recording

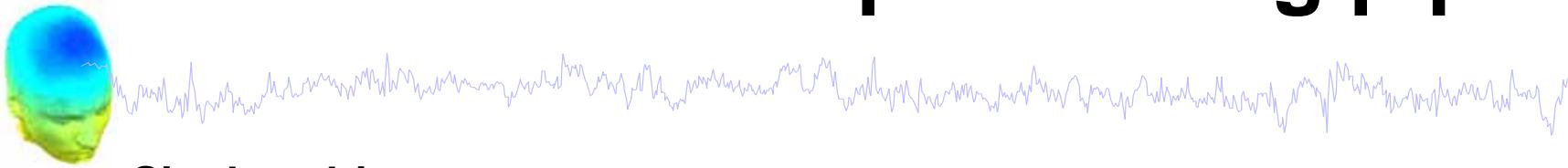


→ **Offline  
processing**

# Target trials



# EEGLAB standard processing pipeline



## Single subject

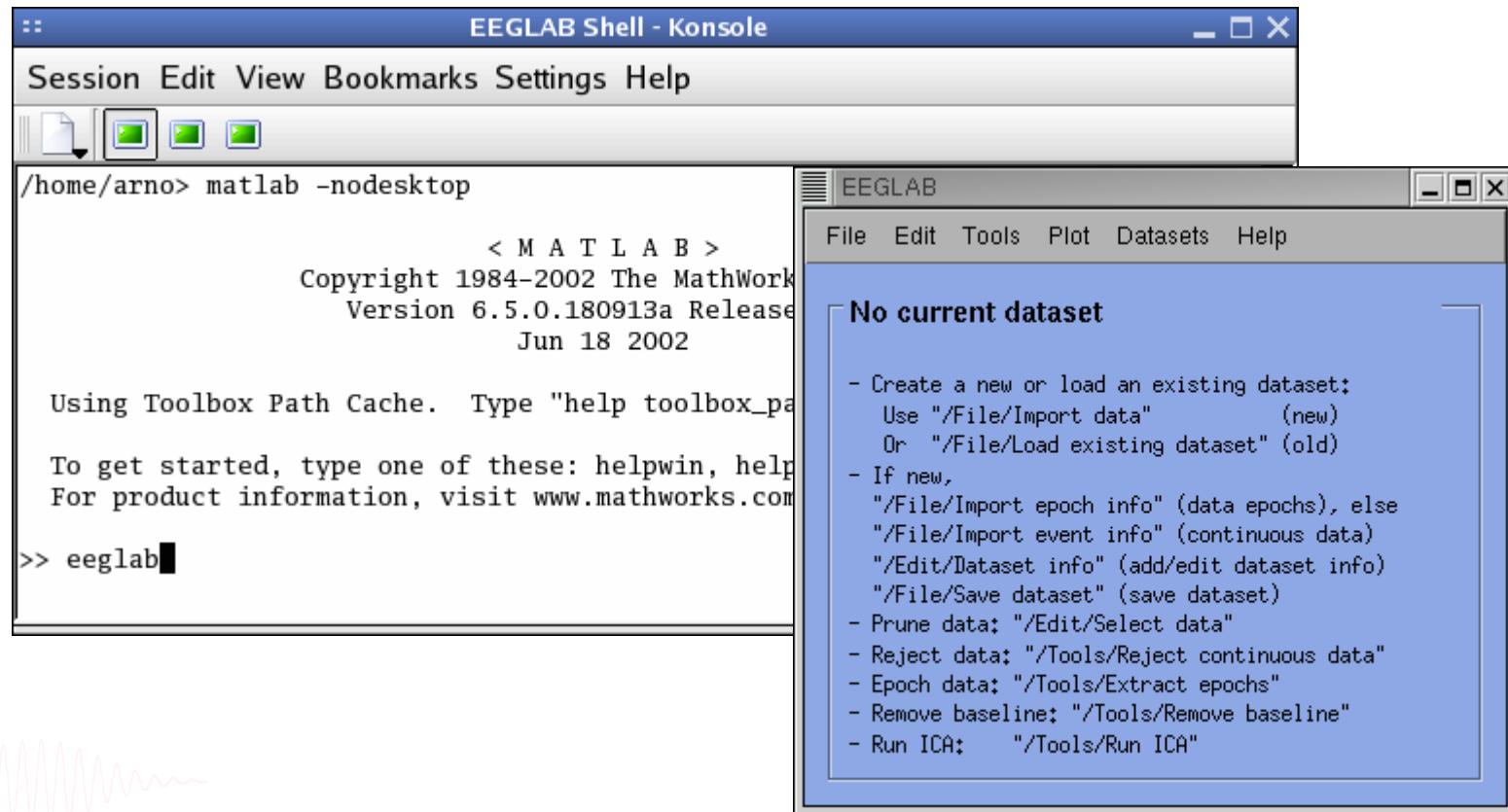
1. Import binary data, events and channel location
2. Edit, Re-reference, Resample, High pass filter data
3. Reject artifacts in continuous data by visual inspection
4. Extract epochs from data & reject artifactual epochs
5. Visualize data measures
6. Perform ICA decomposition
  - Perform source localization of components
  - Analyze components contribution to ERP
  - Analyze components contribution to spectrum

## Multi-subjects

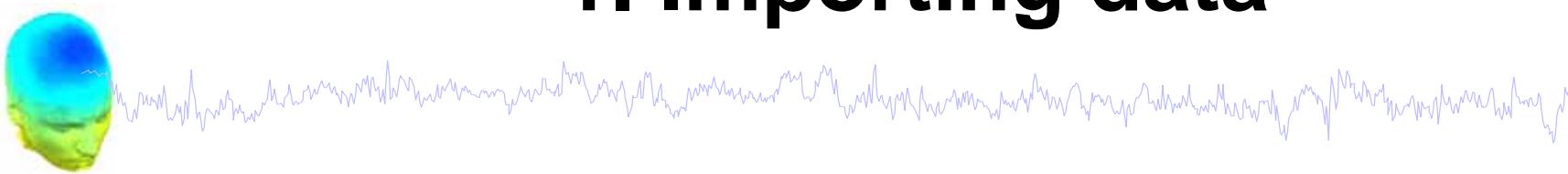
1. Build study
2. Pre-compute measures
3. Cluster components
4. Analyze clusters

**Advanced analysis using scripting and EEGLAB command line functions**

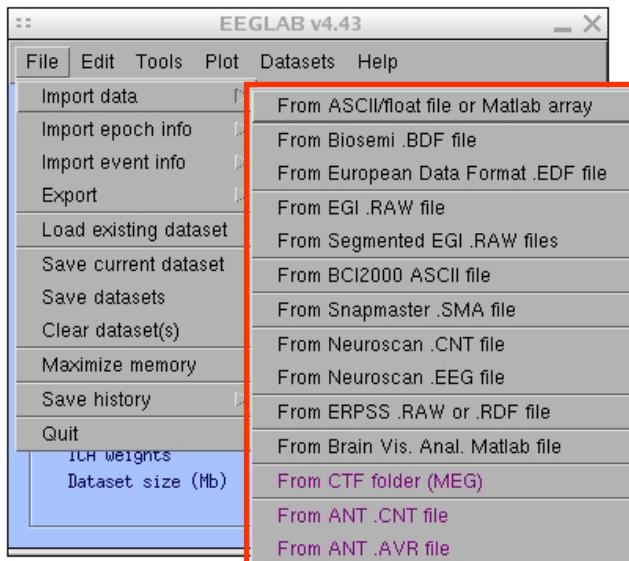
# The EEGLAB Matlab software



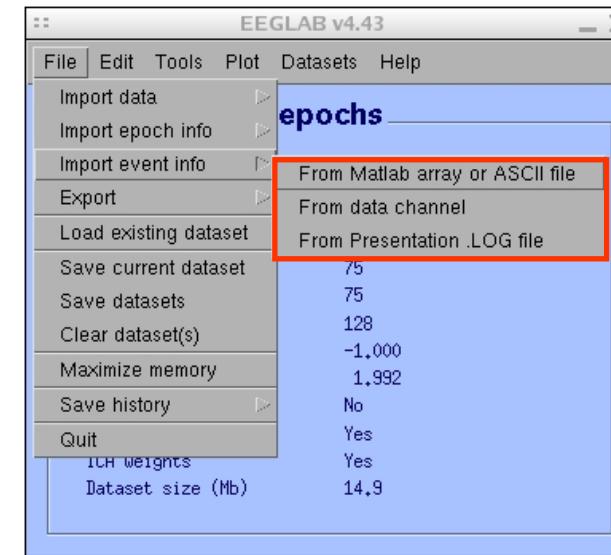
# 1. Importing data



## Import/load data



## Import events



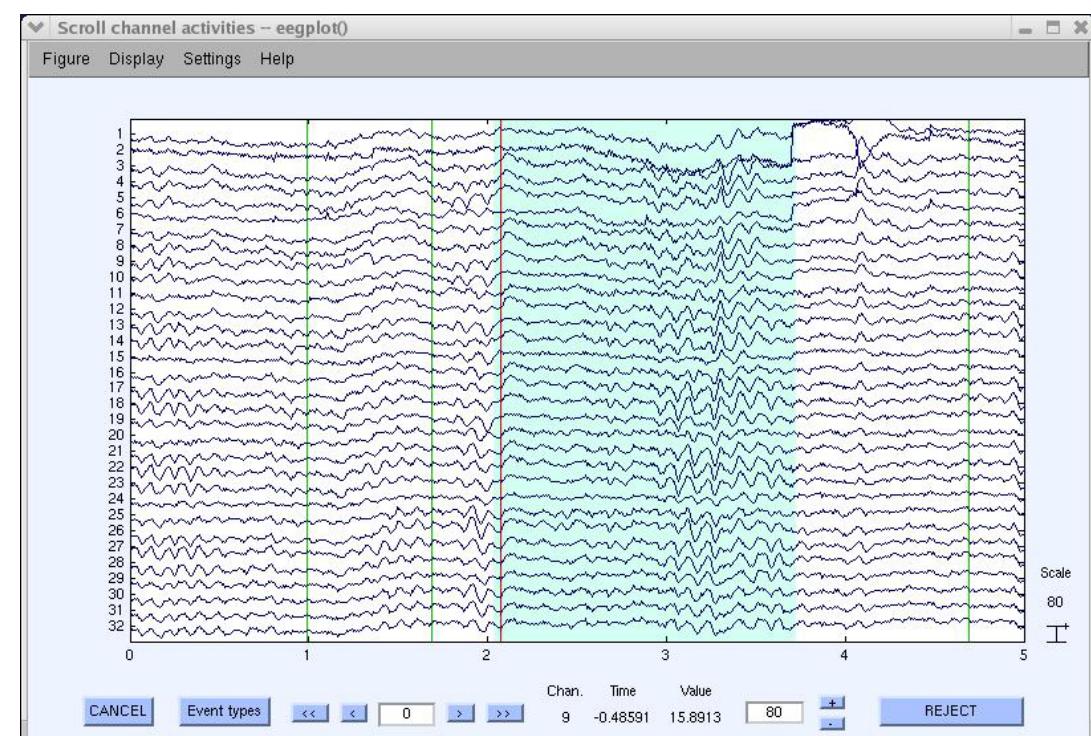
# 1. Importing data



## Data info



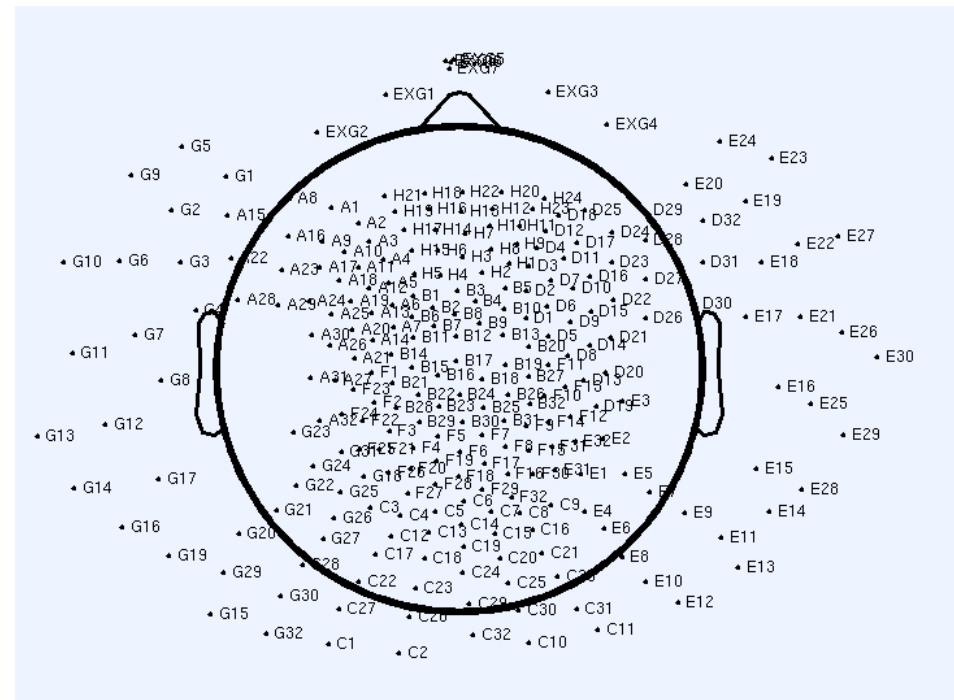
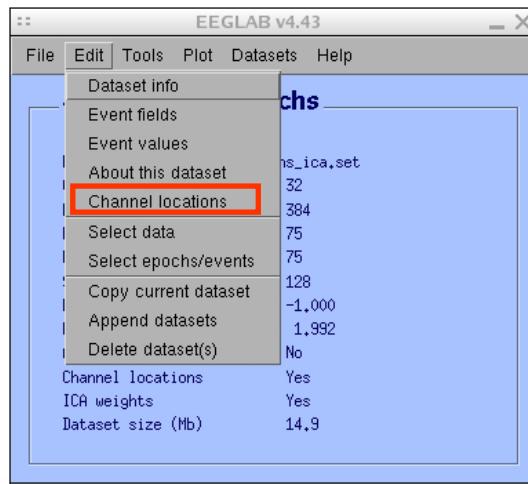
## Scrolling data



# 1. Importing channel location



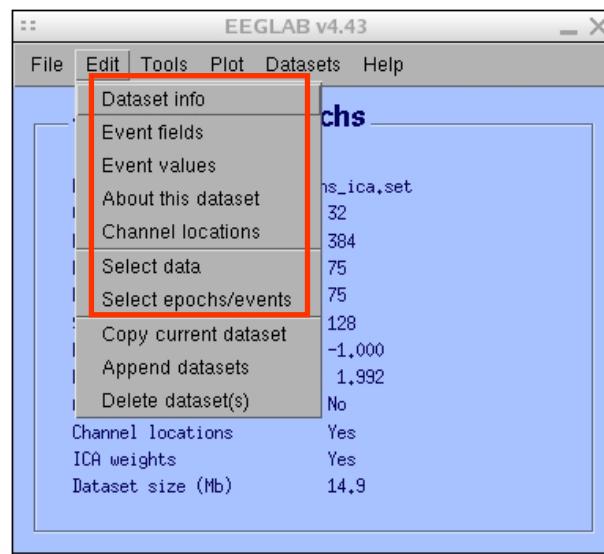
## Import channel location



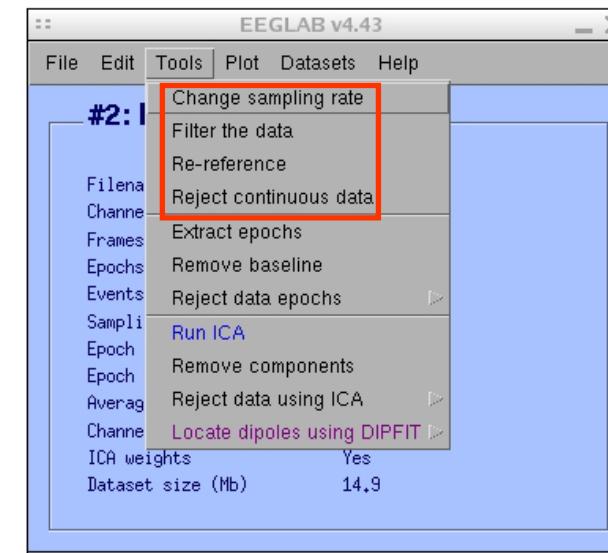
## 2. Edit, Re-reference, Resample, High pass filter data



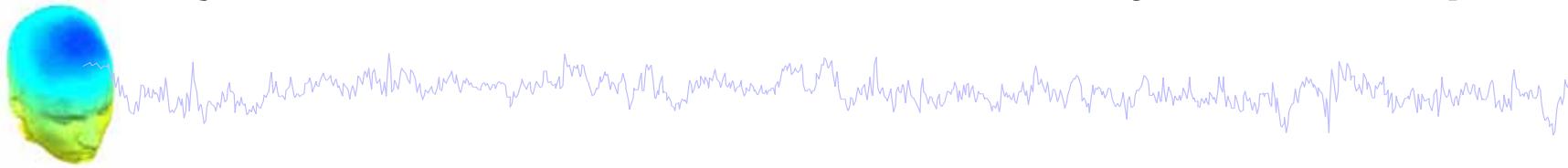
### Edit/select data



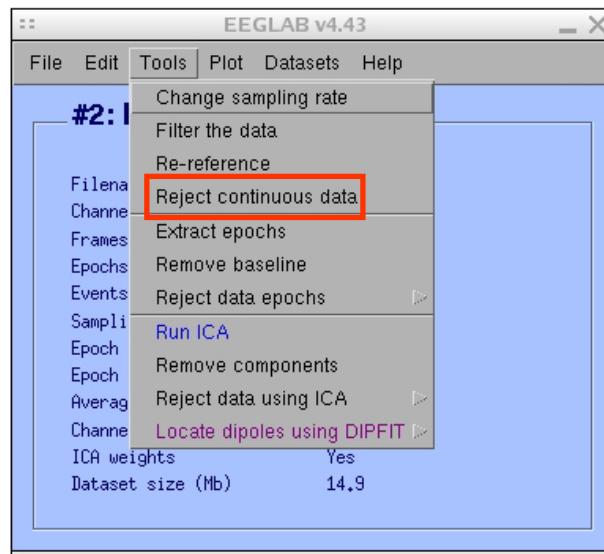
### Preprocessing data



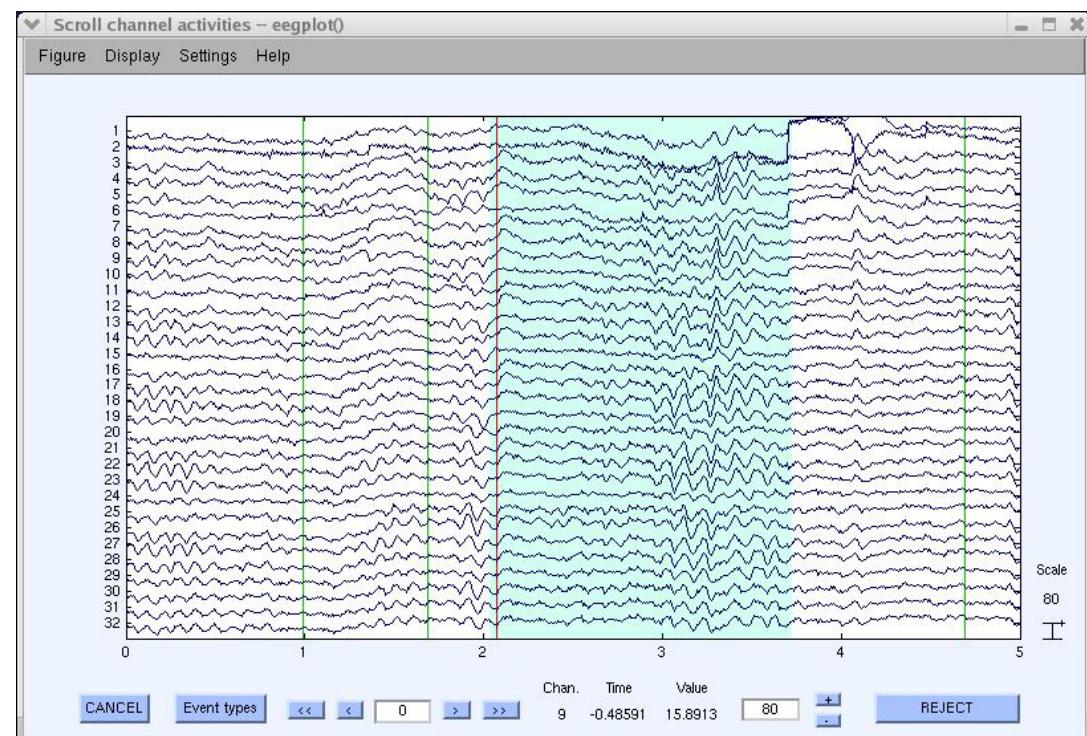
### 3. Reject artifacts in continuous data by visual inspection



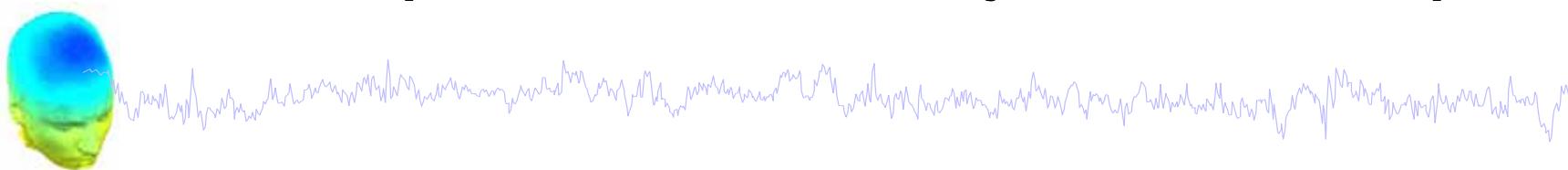
#### Data info



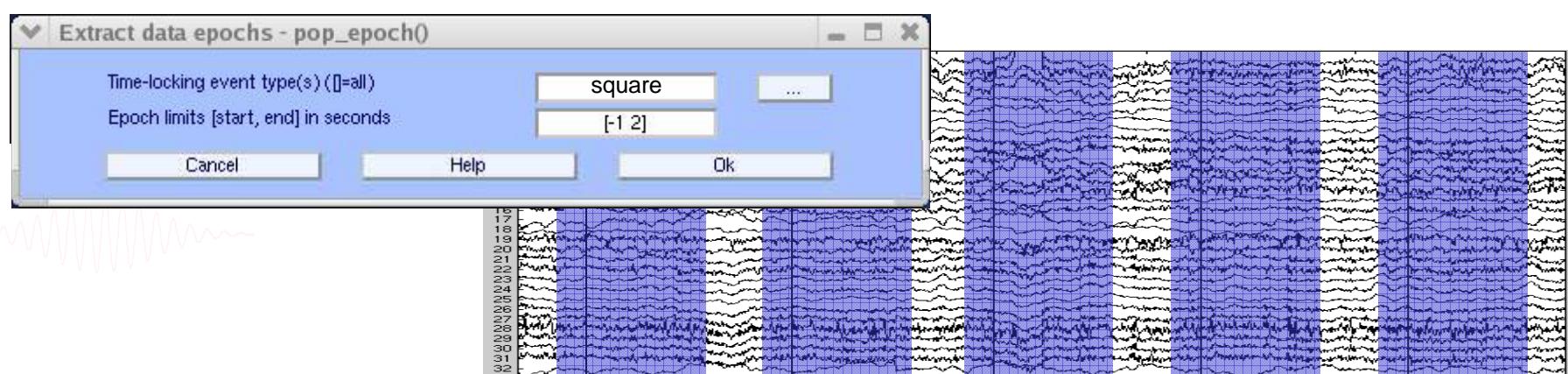
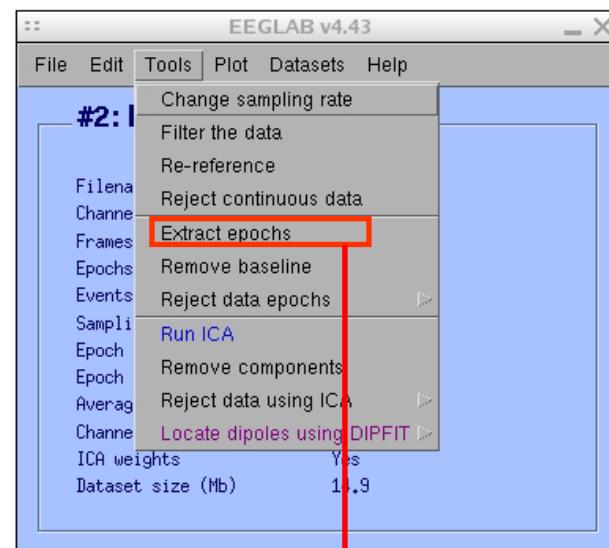
#### Reject portions of continuous data



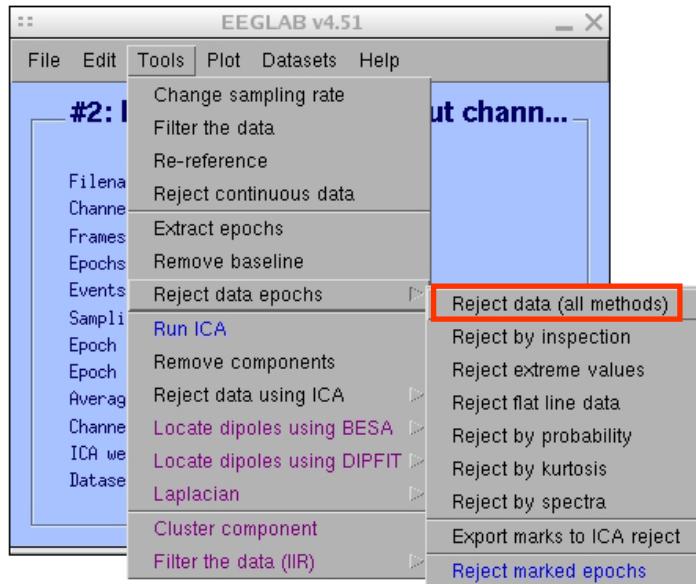
## 4. Extract epochs from data & reject artifactual epochs



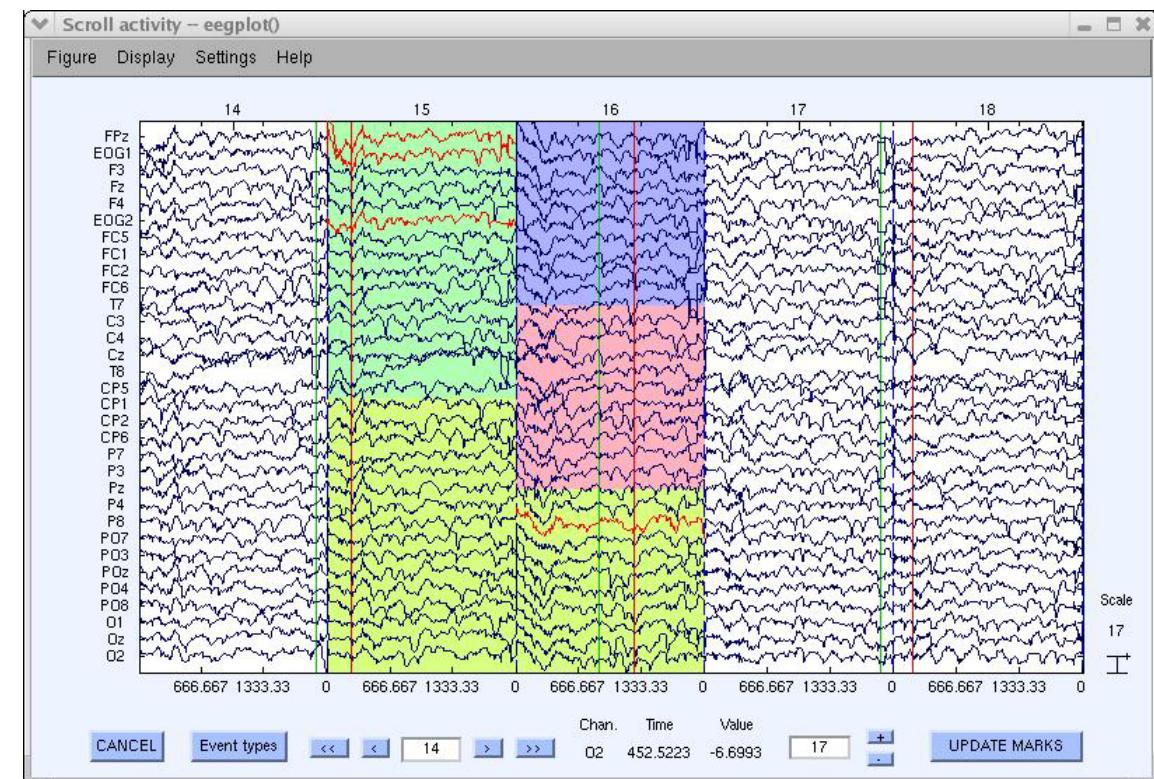
### Preprocessing data



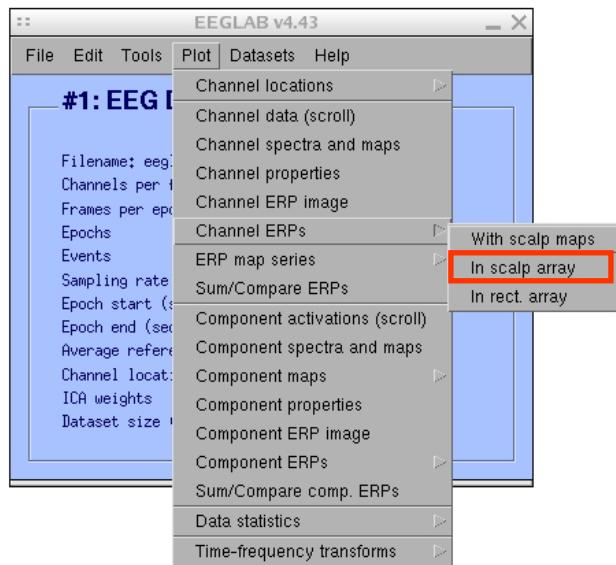
## 4. Extract epochs from data & reject artifactual epochs



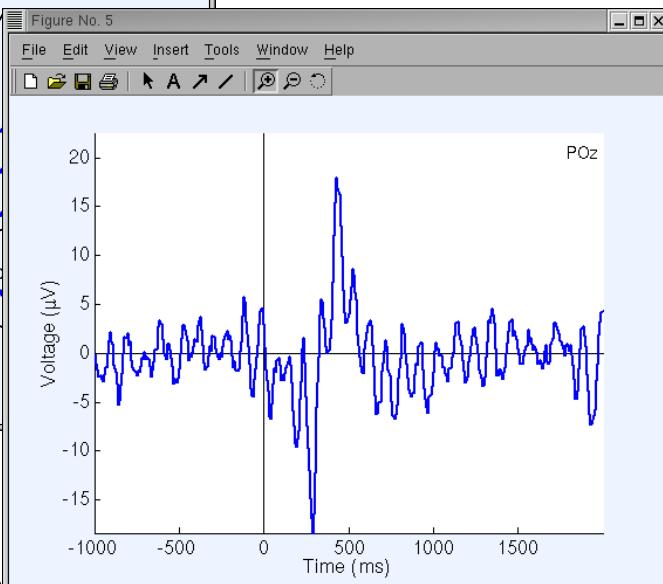
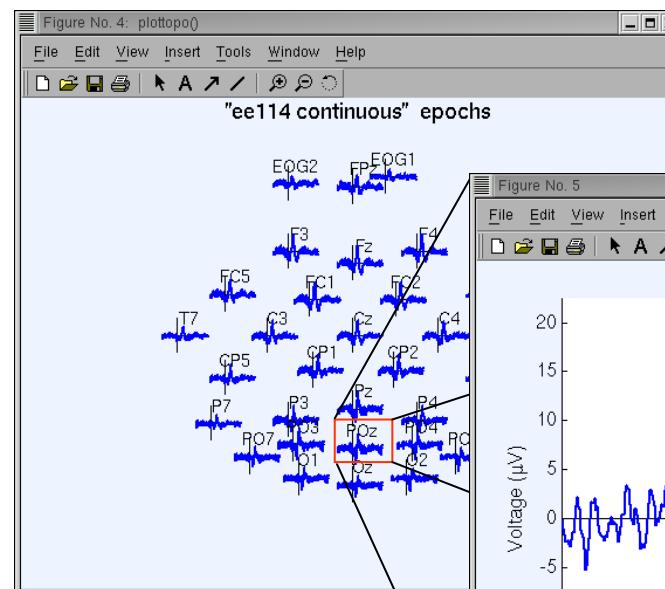
Different color = different rejection methods



# 5. Visualize data measures

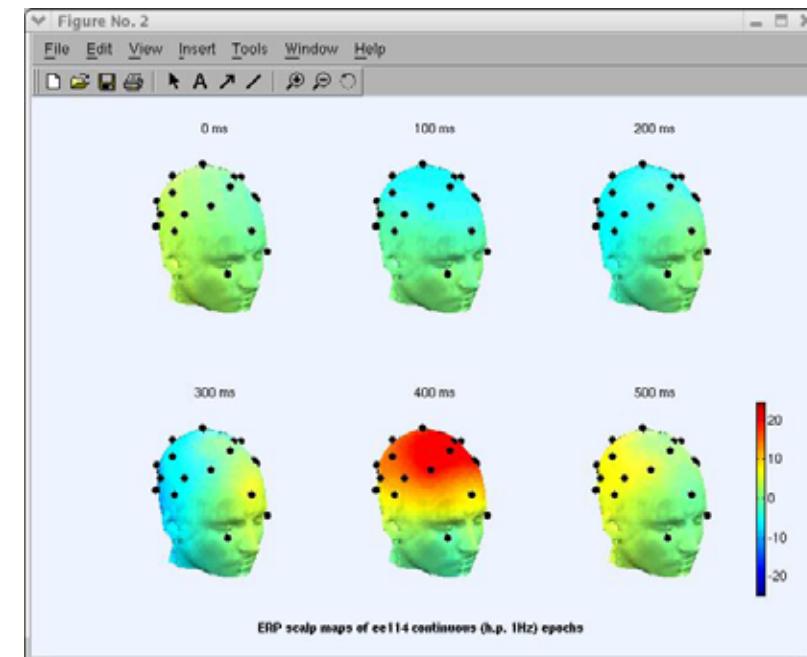
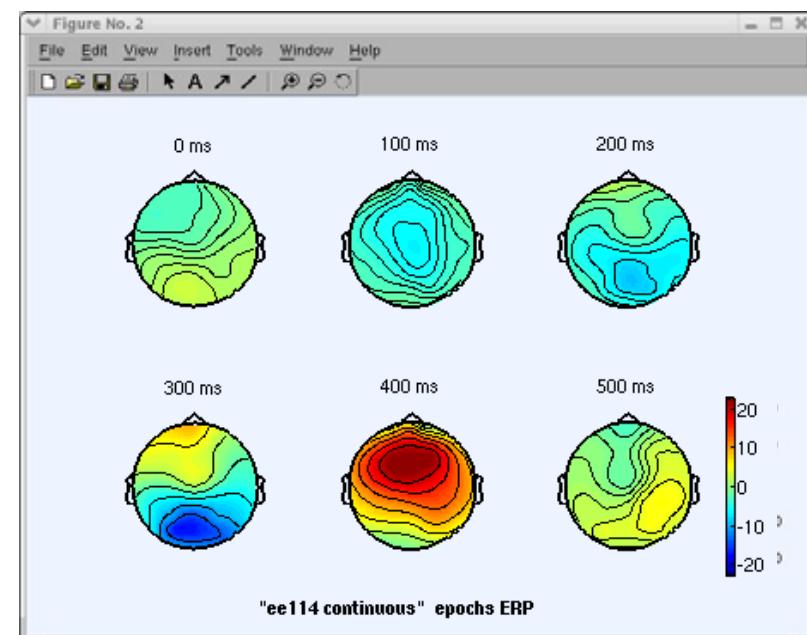
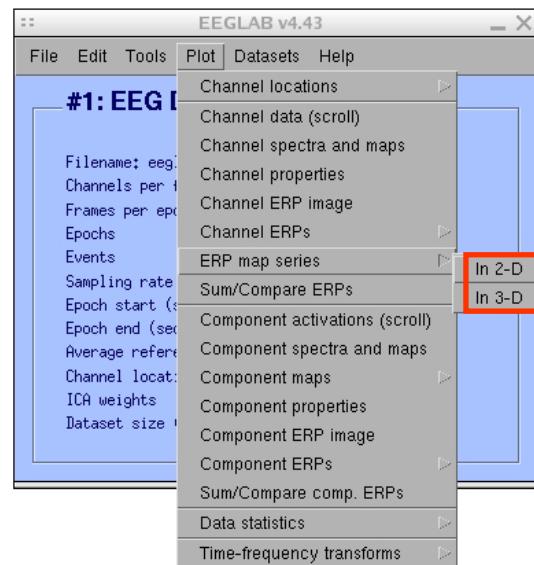


## Plot ERP



# 5. Visualize data measures

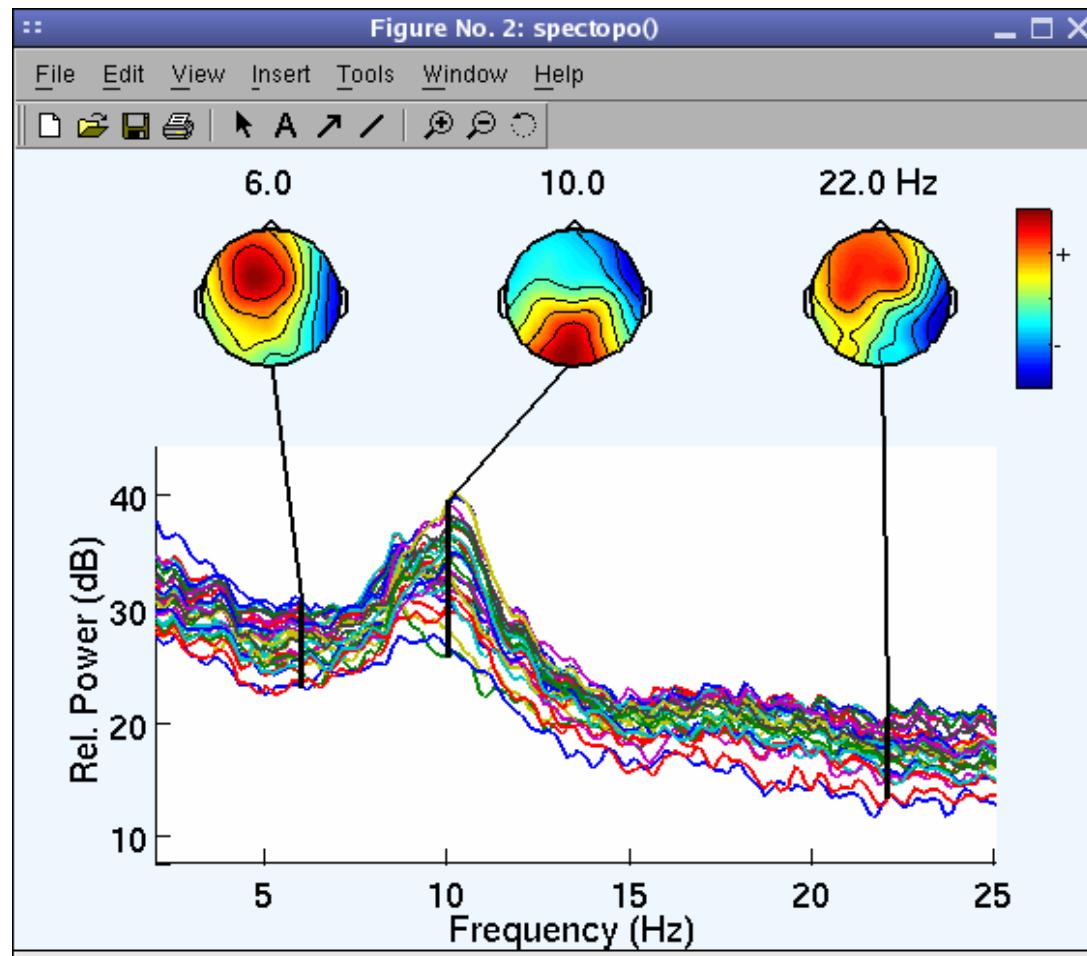
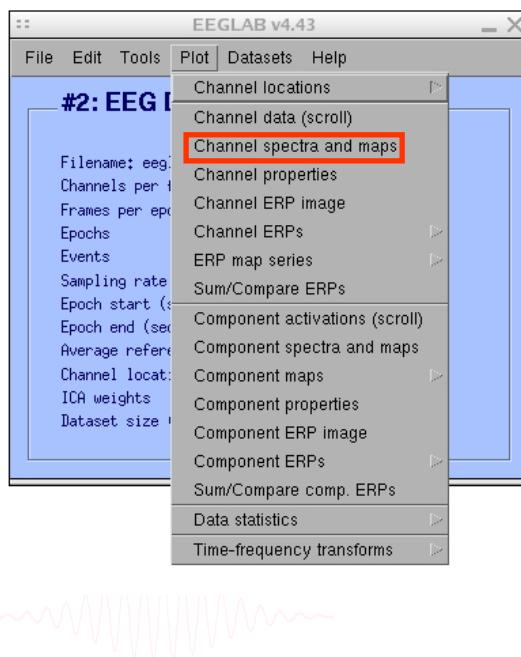
## Plot ERP map series



# 5. Visualize data measures



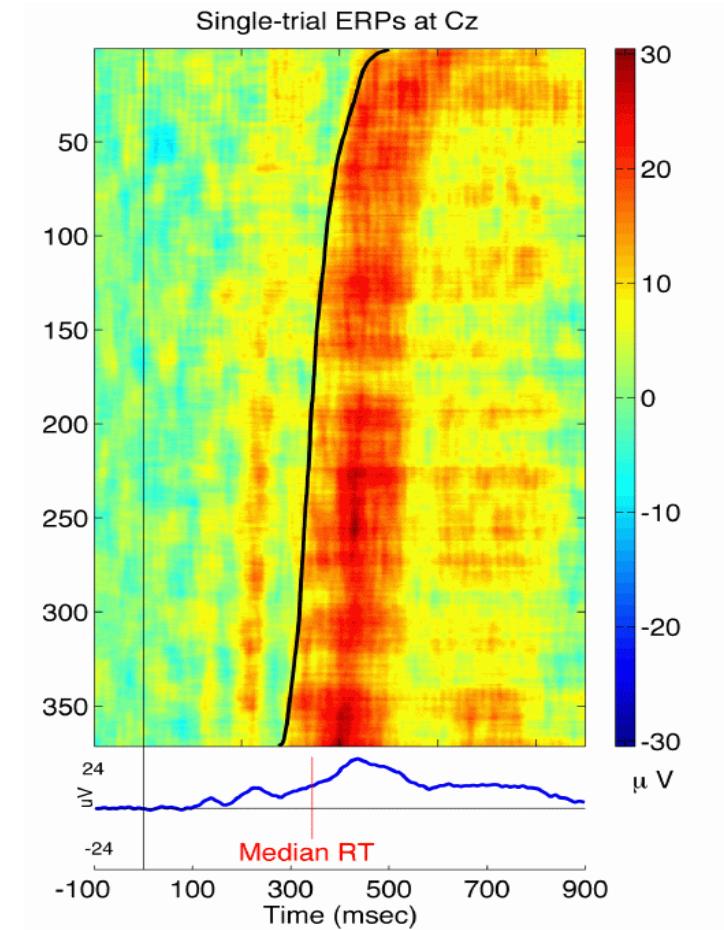
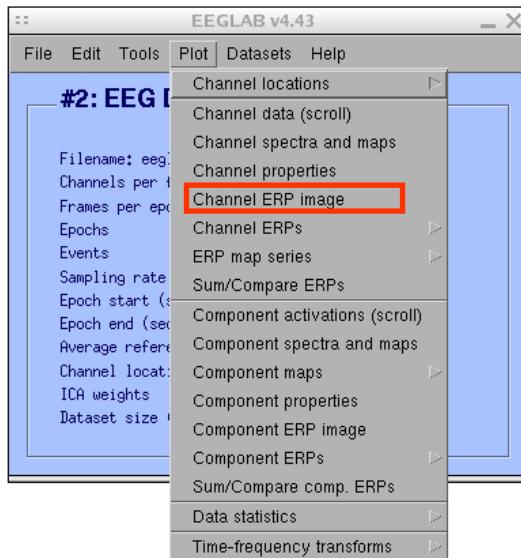
Plot data  
spectrum and  
maps



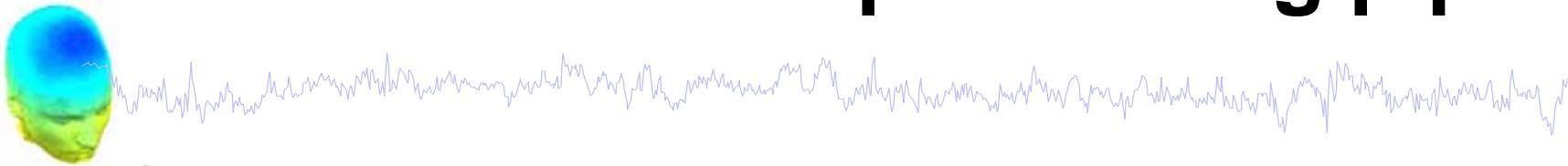
# 5. Visualize data measures



Plot channel ERPimage



# EEGLAB standard processing pipeline



## Single subject

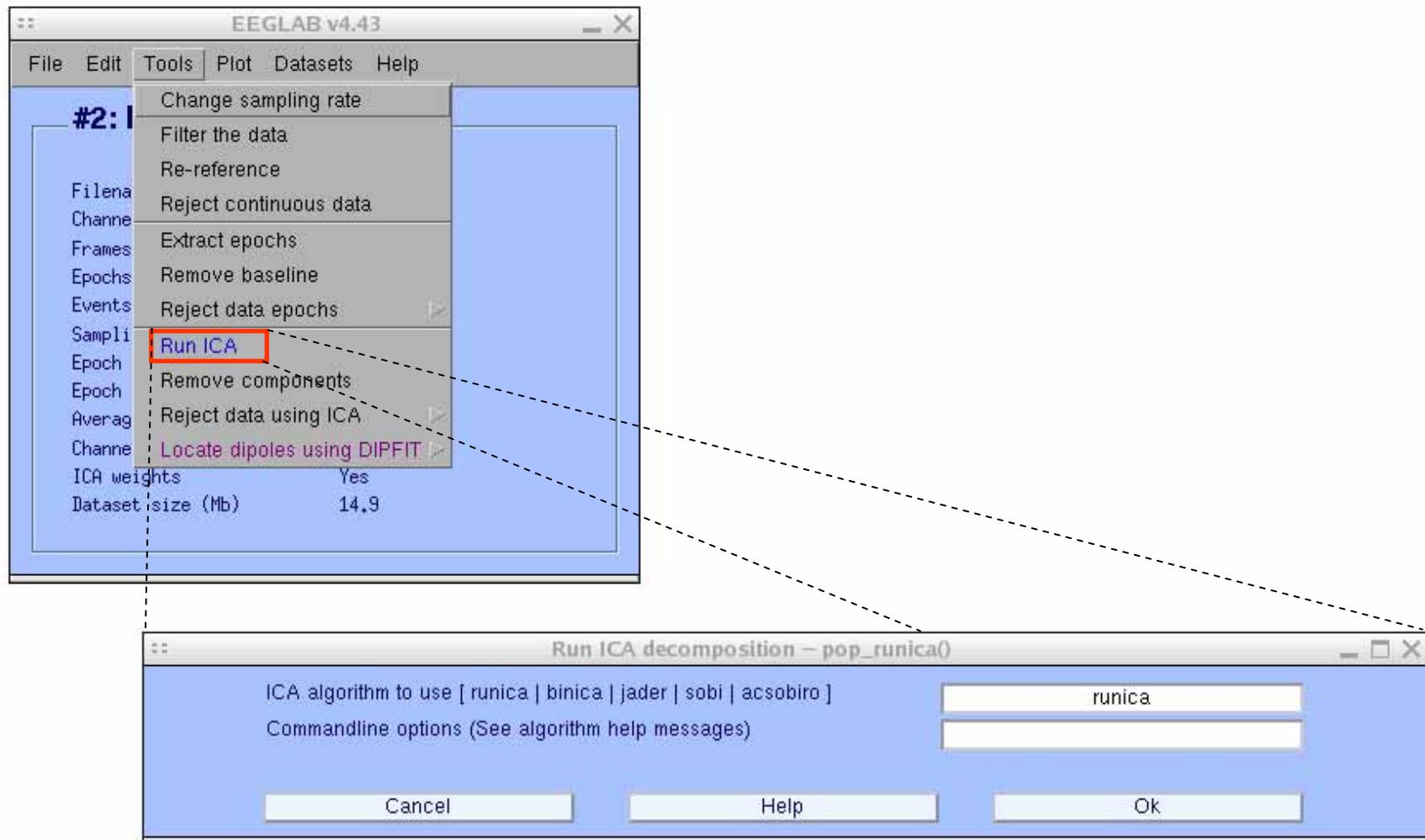
1. Import binary data, events and channel location
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4. Extract epochs from data & reject artifactual epochs
5. Visualize data measures
6. Perform ICA decomposition
  - Perform source localization of components
  - Analyze components contribution to ERP
  - Analyze components contribution to spectrum

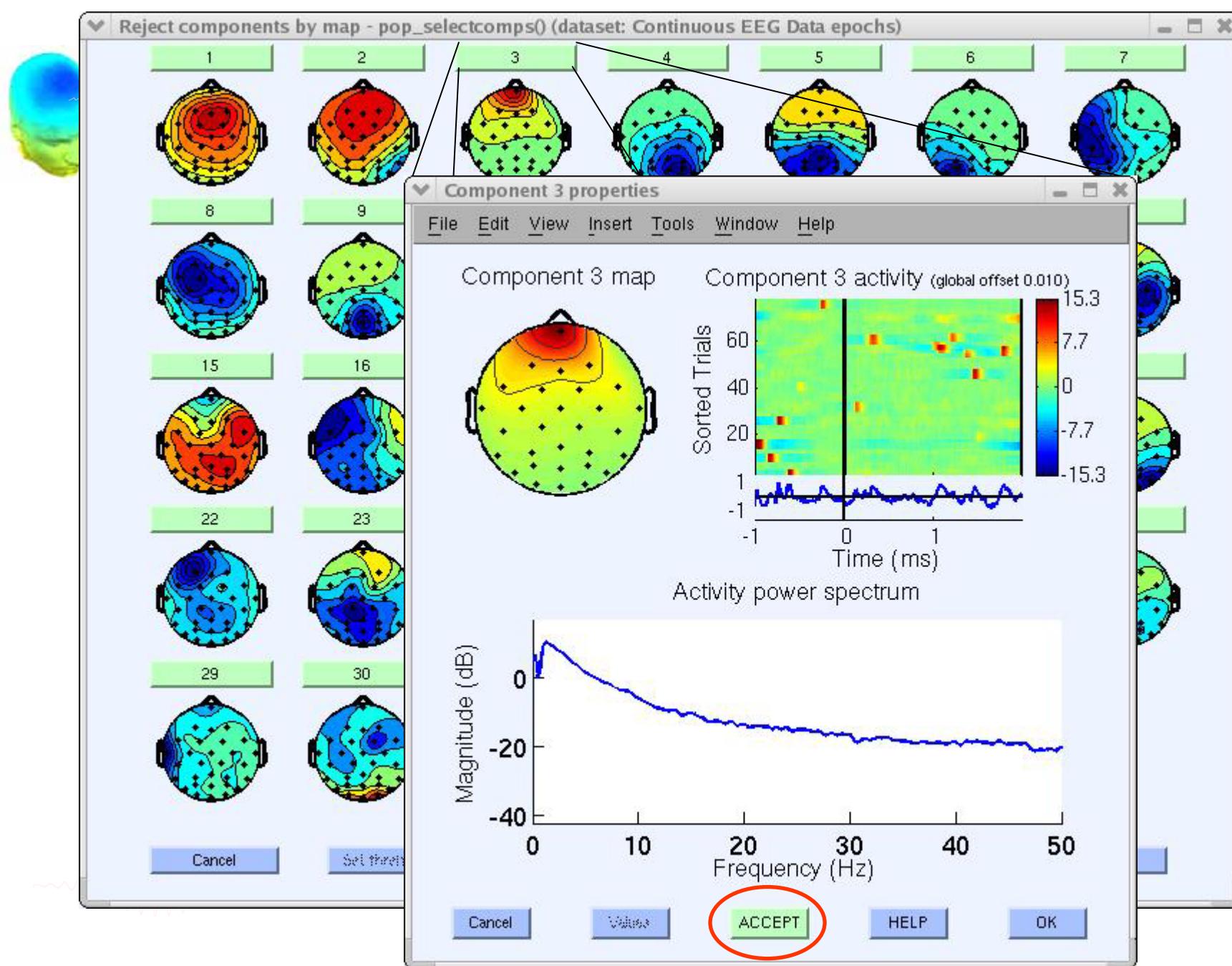
## Multi-subjects

1. Build study
2. Pre-compute measures
3. Cluster components
4. Analyze clusters

Advanced analysis using scripting and EEGLAB command line functions

# 6. Perform ICA decomposition





# Localizing components

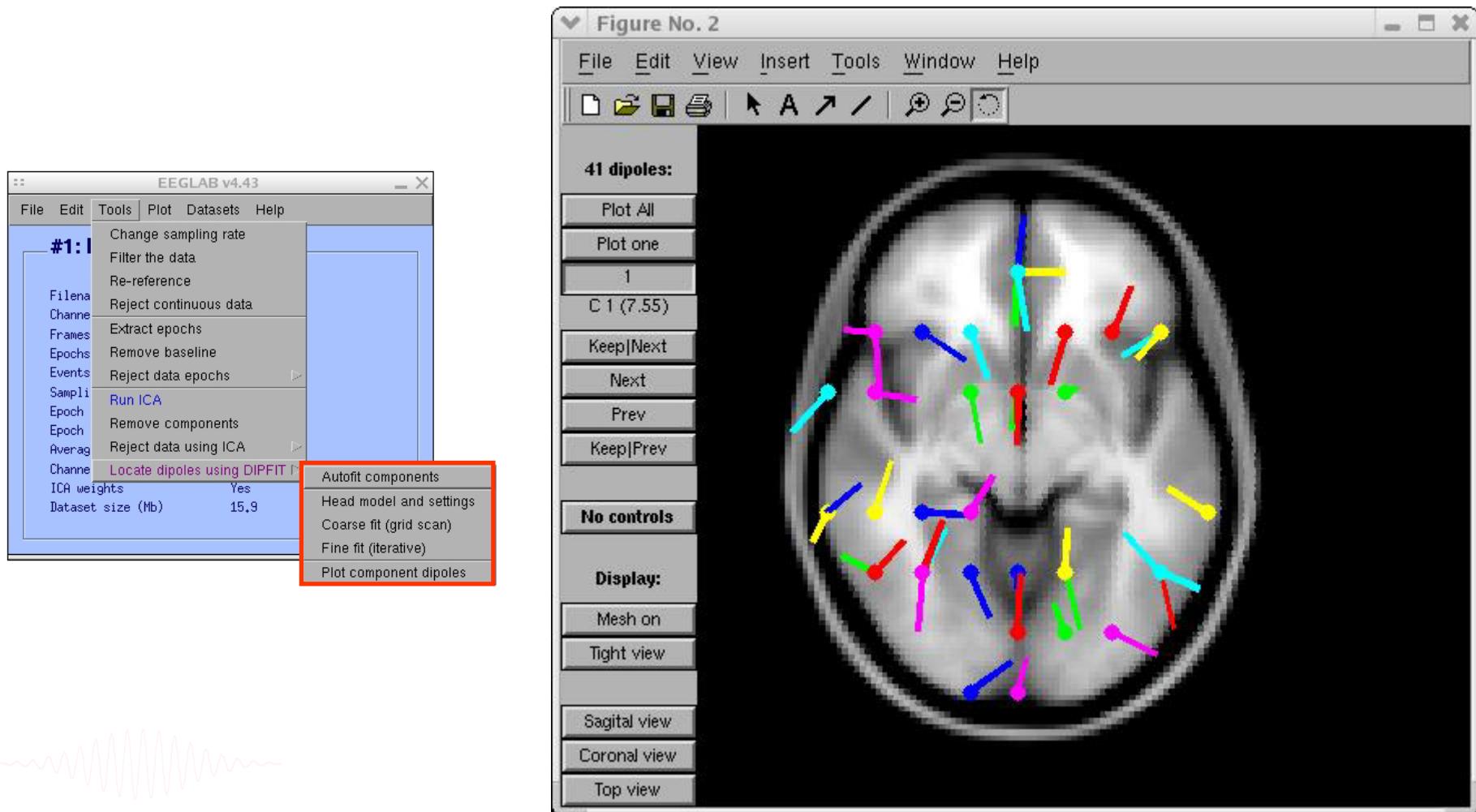
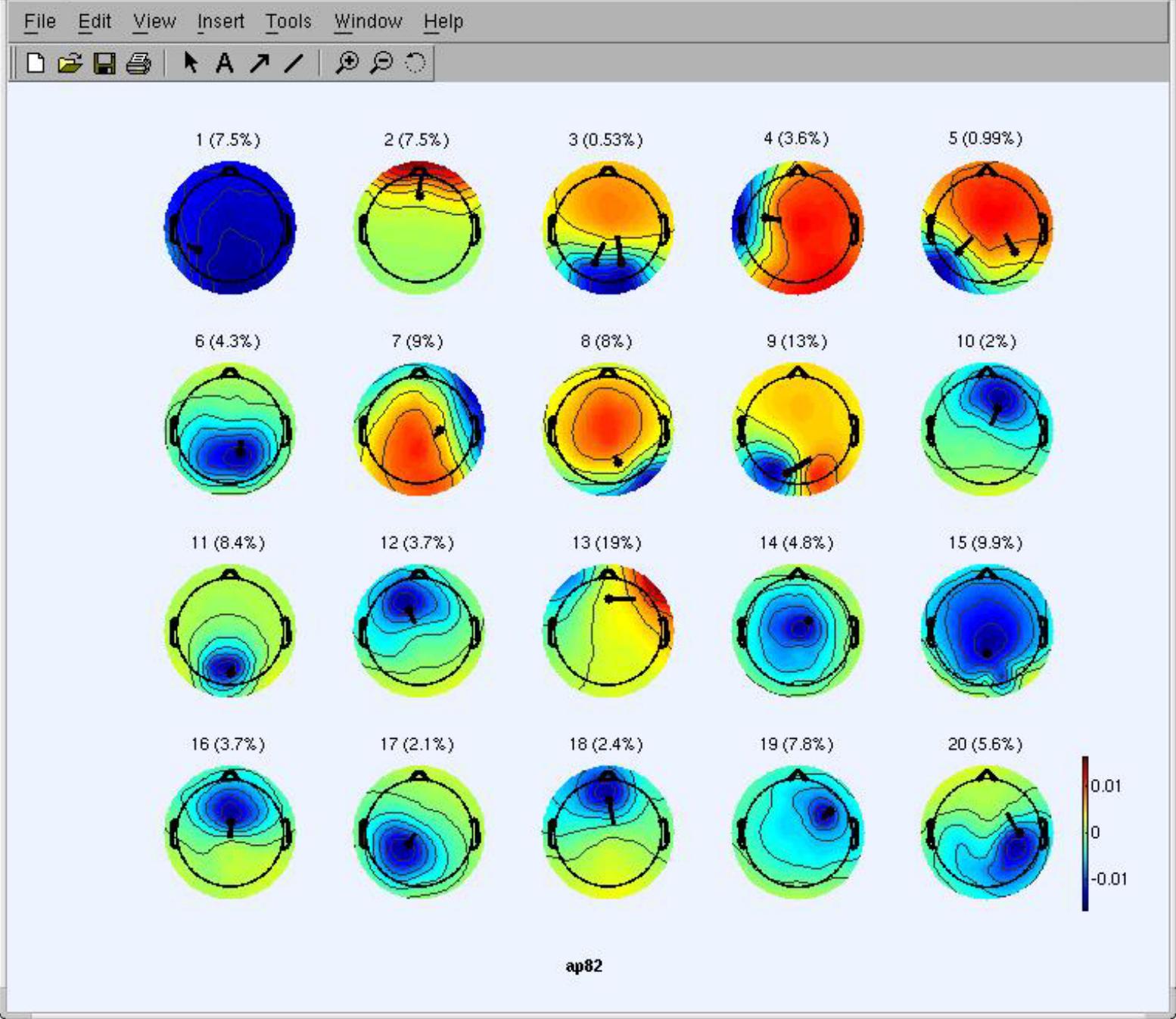
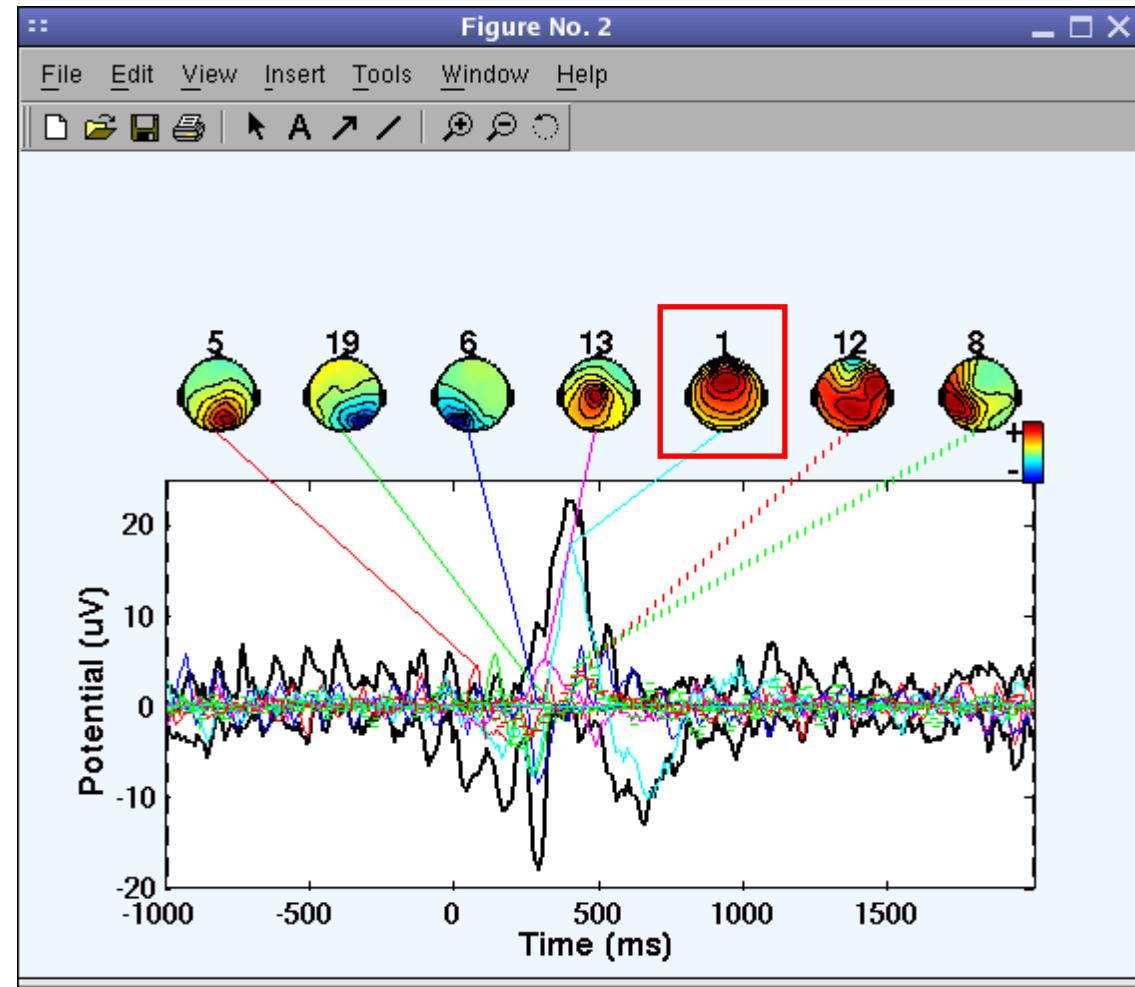
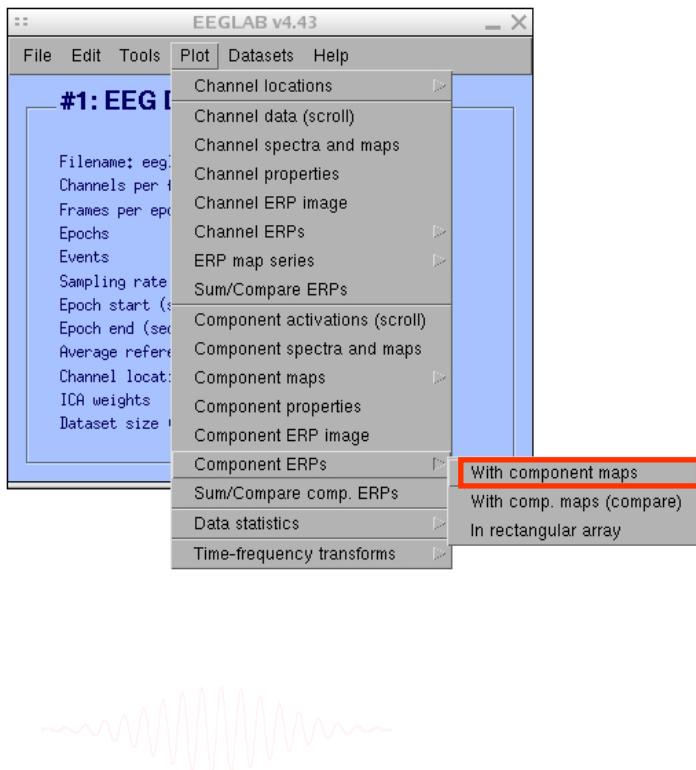
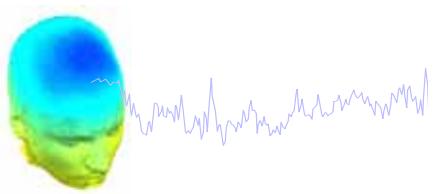


Figure No. 4

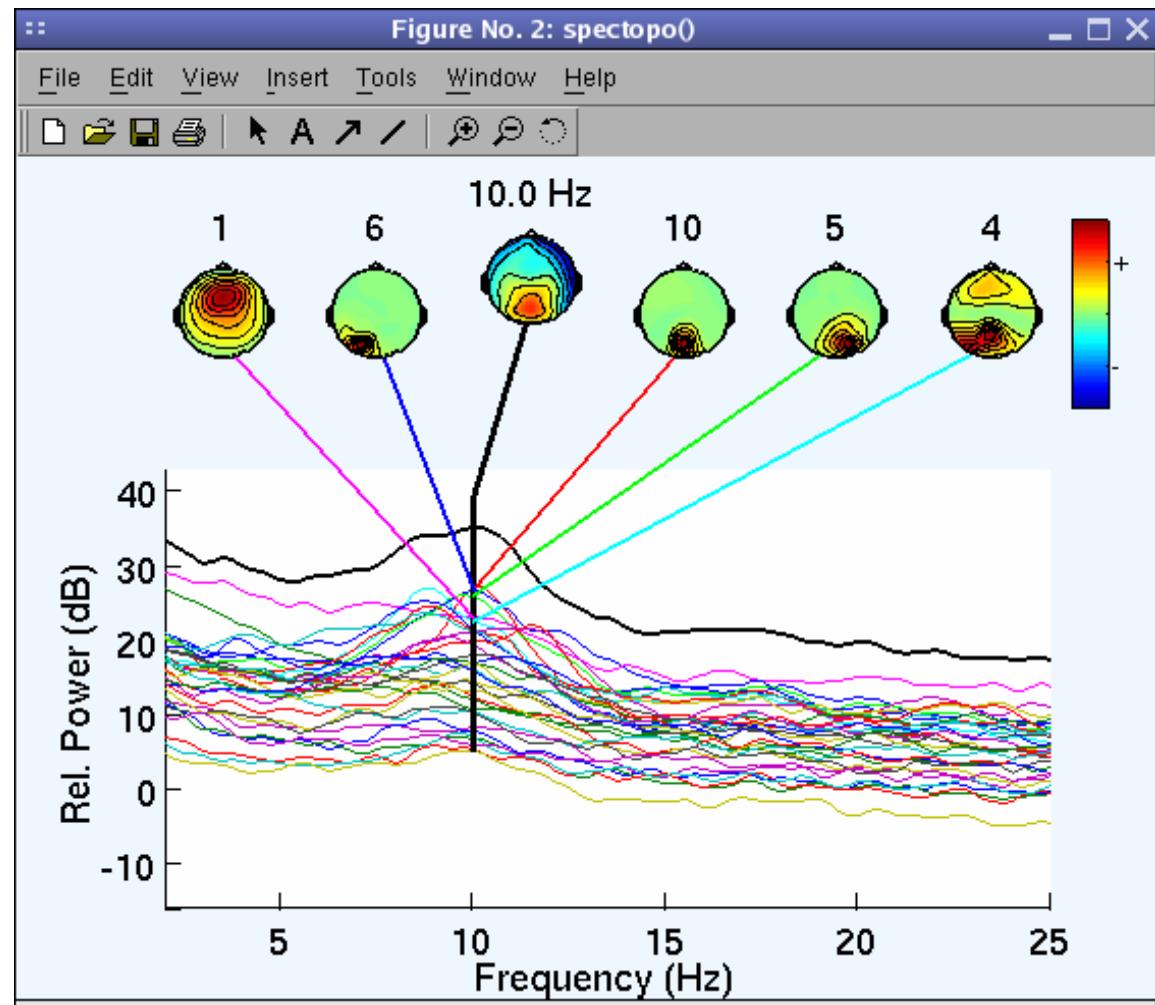
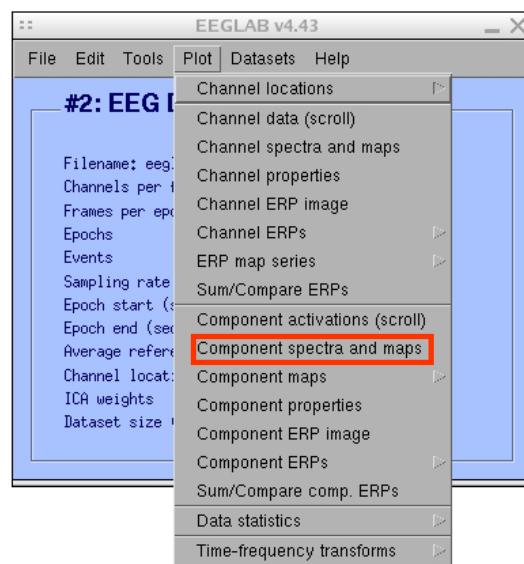


# Component contribution to the ERP

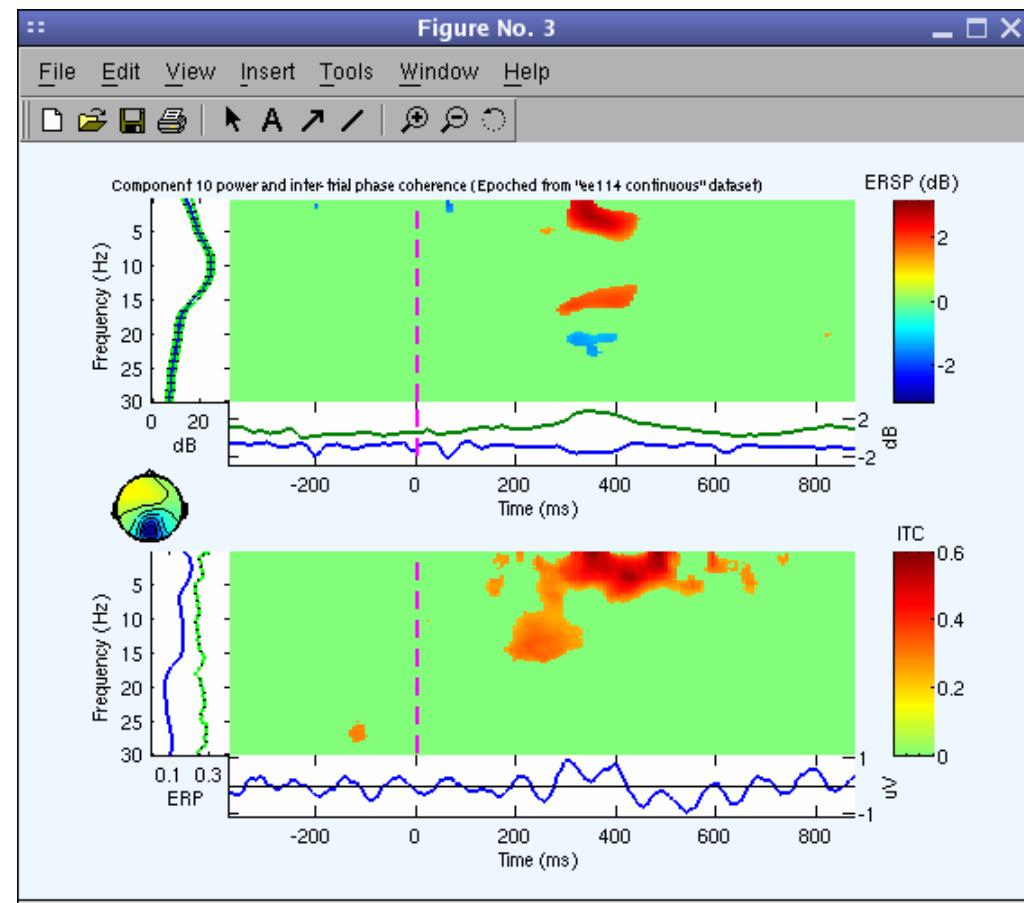
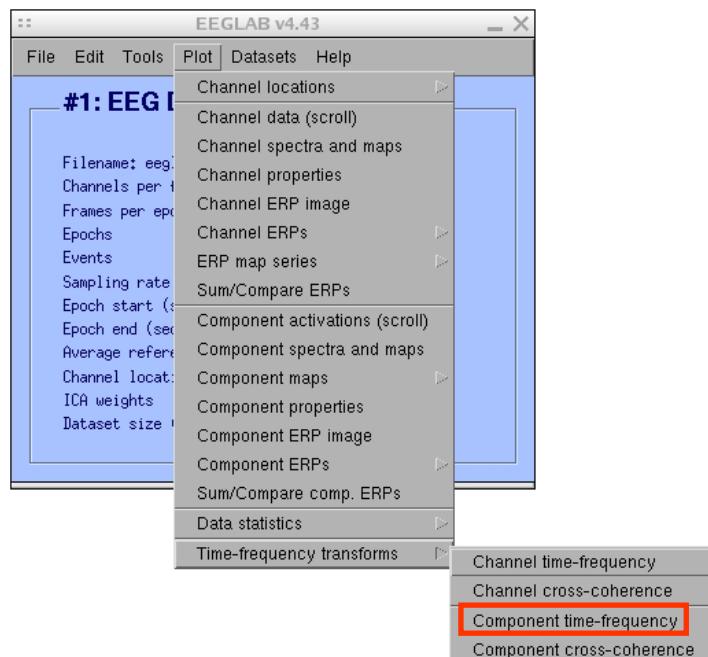




# Component contribution to the EEG spectrum



# Component time-frequency



# EEGLAB standard processing pipeline



## Single subject

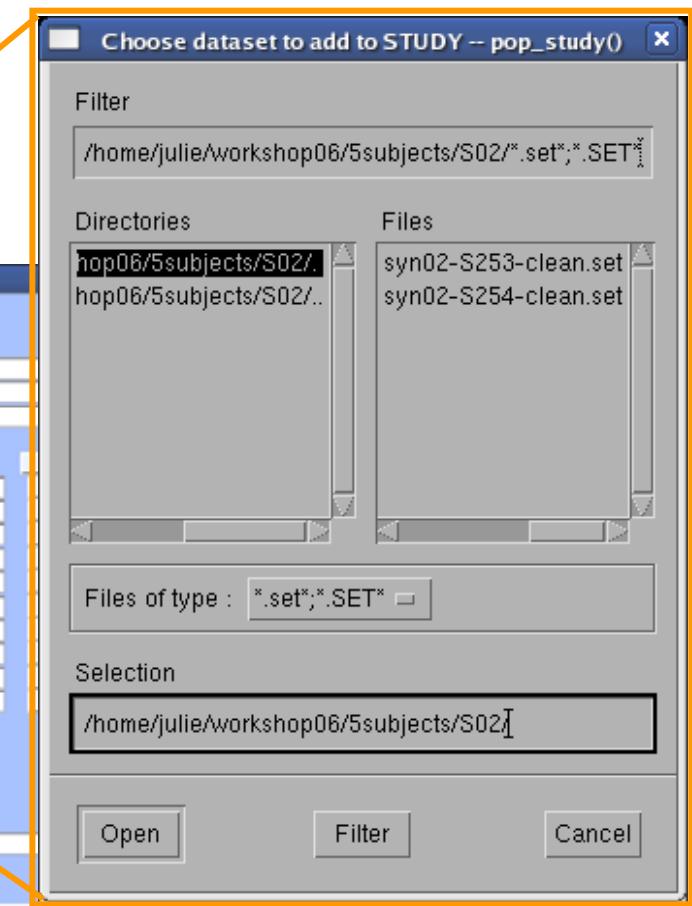
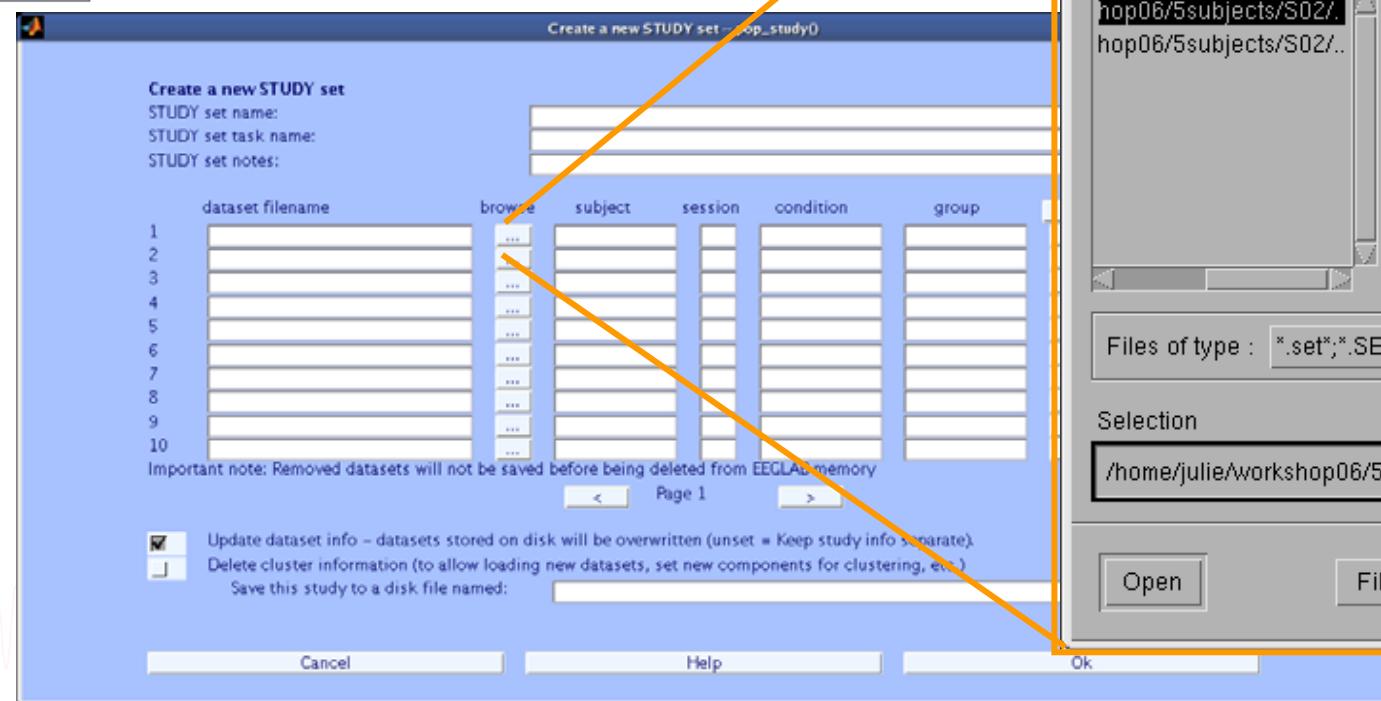
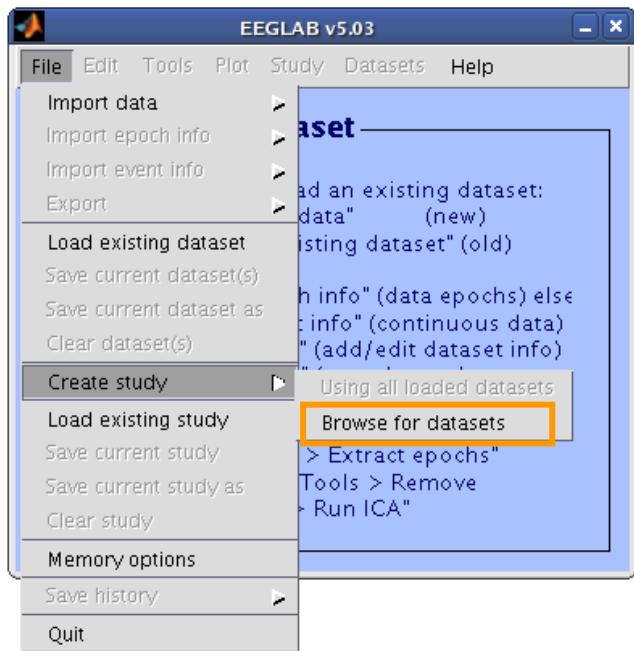
1. Import binary data, events and channel location
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## Multi-subjects

1. Build study
2. Pre-compute measures
3. Cluster components
4. Analyze clusters

Advanced analysis using scripting and EEGLAB command line functions

# 1. Build a STUDY

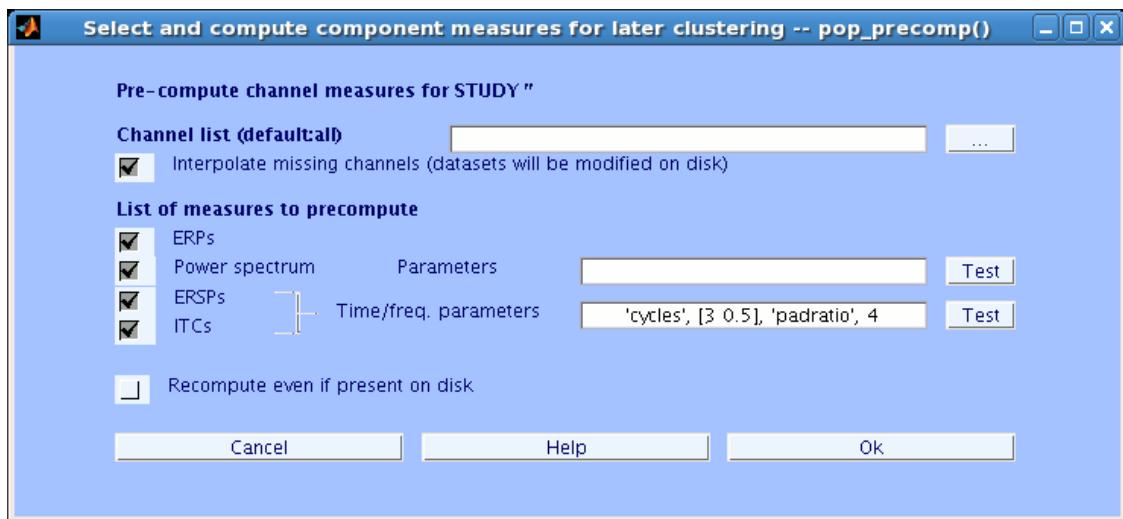
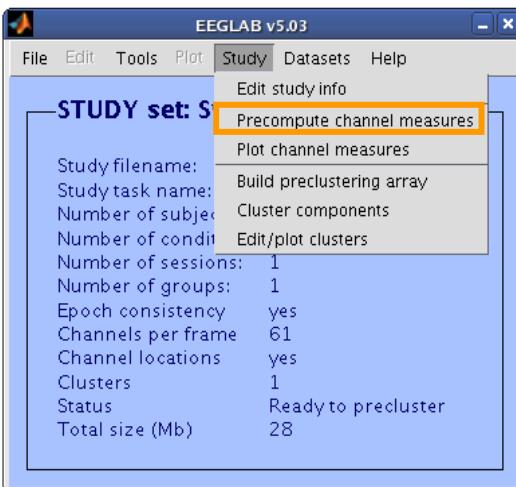
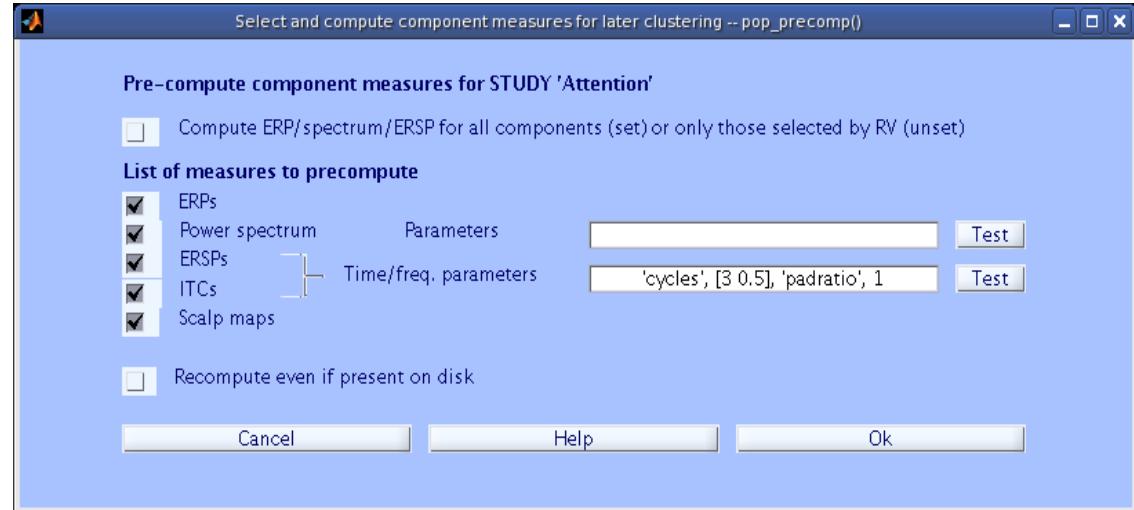
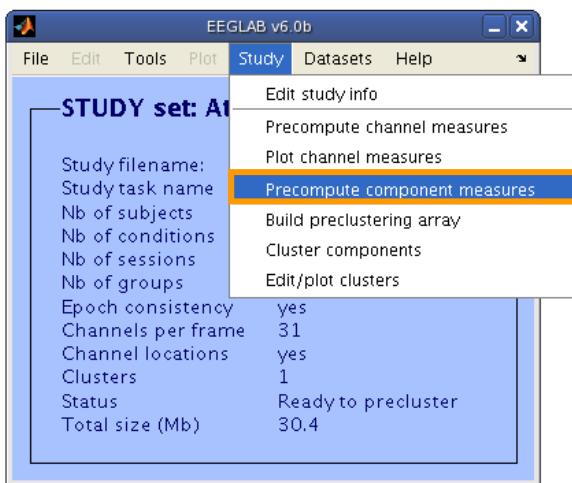


# Channels

# Components



# 2. Pre-compute measures



# 3. Cluster components



EEGLAB v6.0b

File Edit Tools Plot Study Datasets Help

**STUDY set: Attention**

Study filename: Attention  
Study task name: Attention  
Nb of subjects: 181  
Nb of conditions: 1  
Nb of sessions: 1  
Nb of groups: 1  
Epoch consistency: yes  
Channels per frame: 31  
Channel locations: yes  
Clusters: 1  
Status: Pre-clustered  
Total size (Mb): 32.4

Edit study info  
Precompute channel measures  
Plot channel measures  
Precompute component measures  
**Build preclustering array**  
Cluster components  
Edit/plot clusters

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

**Build pre-clustering matrix for STUDY 'Attention'**  
Select the cluster to refine during sub-clustering (any existing sub-hierarchy will be overwritten)  
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
<input checked="" type="checkbox"/> ERPs	10	<input checked="" type="checkbox"/>	1
<input checked="" type="checkbox"/> dipoles	3	<input checked="" type="checkbox"/>	10
<input type="checkbox"/> scalp maps	10	<input checked="" type="checkbox"/>	1
<input checked="" type="checkbox"/> ERSPs	20	<input checked="" type="checkbox"/>	1
<input checked="" type="checkbox"/> ITCs	10	<input checked="" type="checkbox"/>	1
<input type="checkbox"/> Final dimensions	10	<input type="checkbox"/>	Help

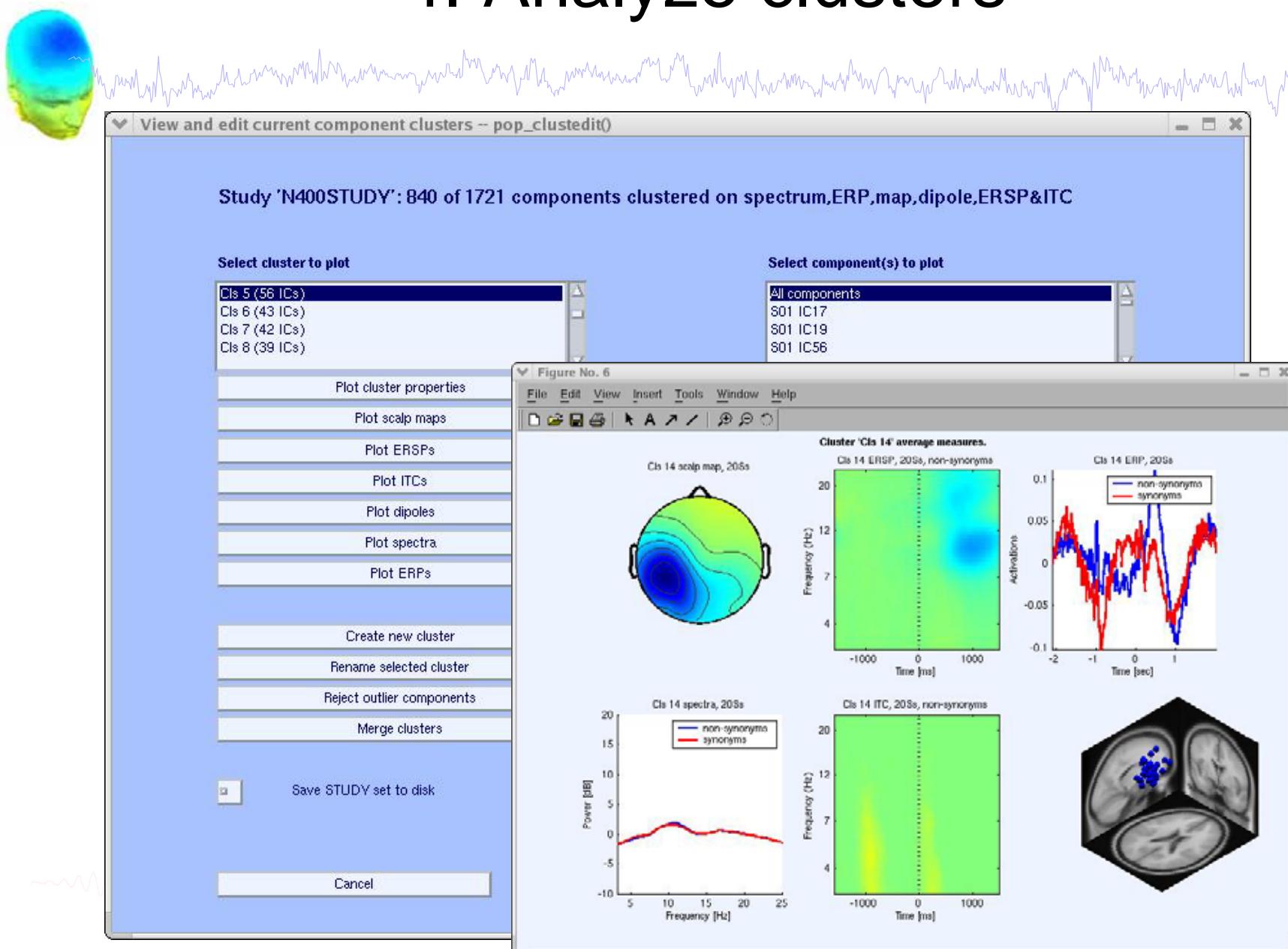
Freq. range [Hz] 3.25  
Time range [ms] 0.600  
Use channel values  
Time range [ms] 0.1500  
Freq. range [Hz] 3.45  
Time range [ms] 0.600  
Freq. range [Hz] 2.30  
 Absolute values

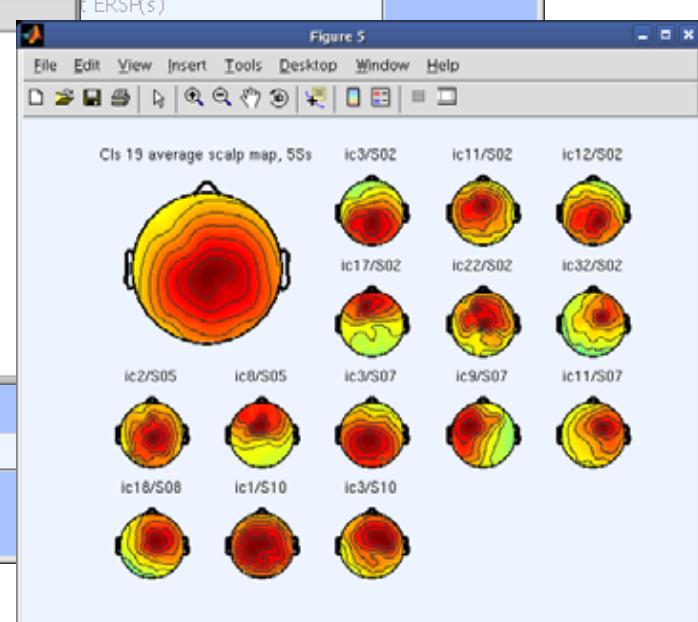
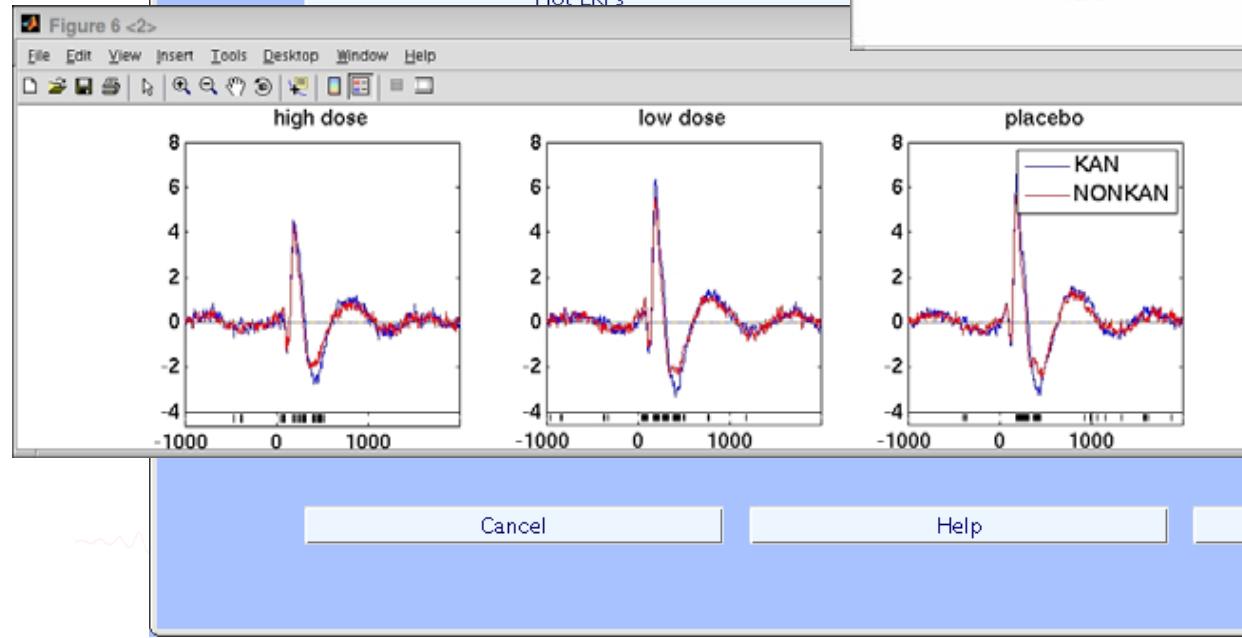
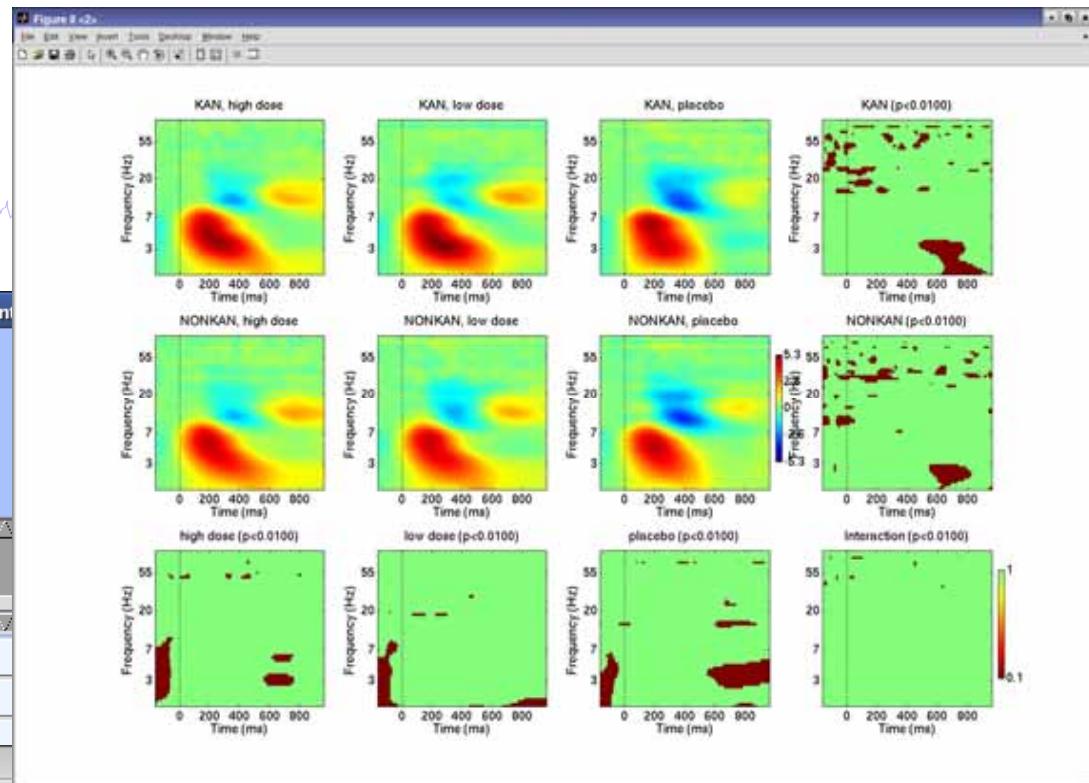
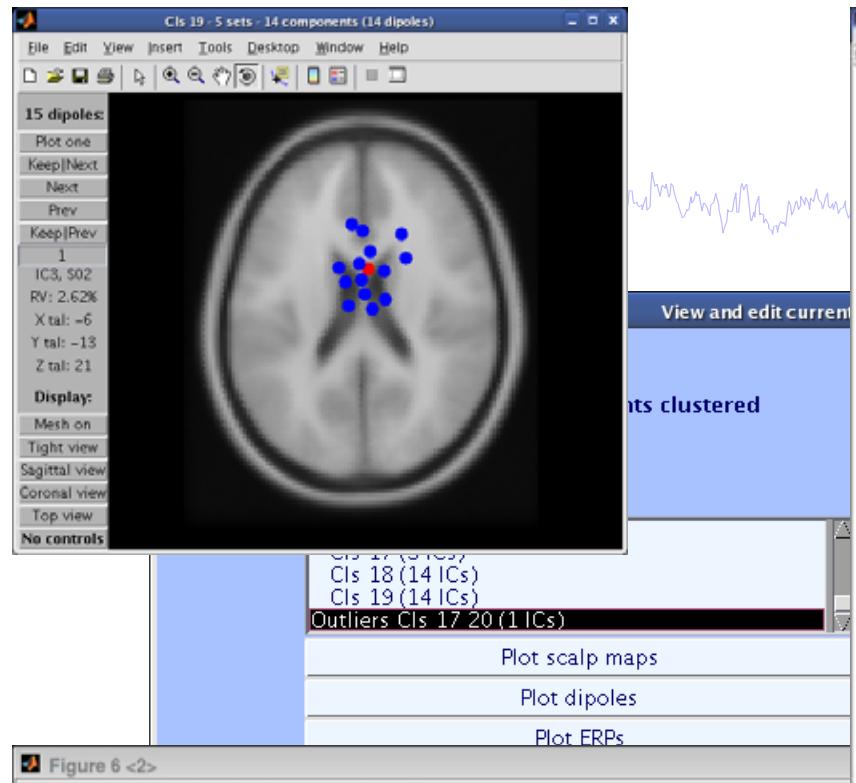
Save STUDY to file /home/julie/WorkshopSD2007/STUDY/attention.study ...

Cancel Help Ok

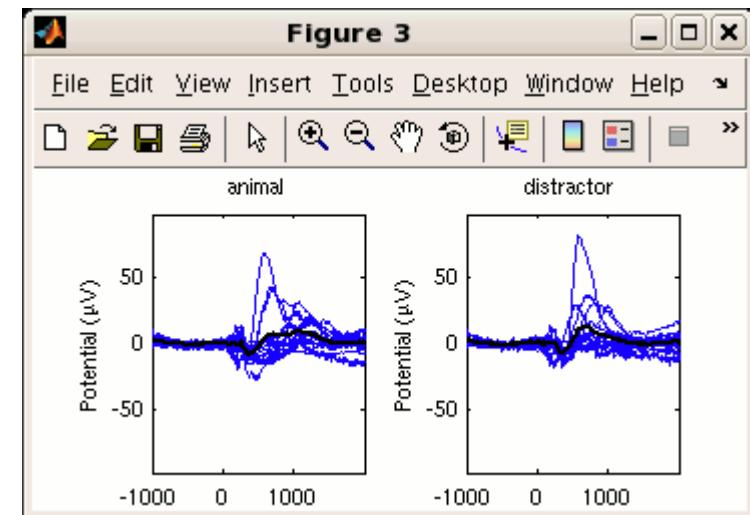
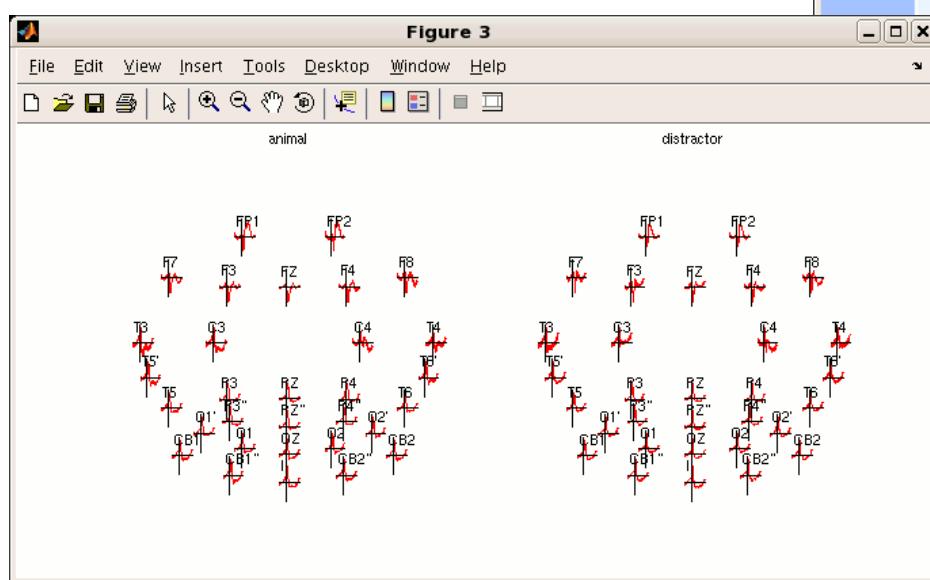
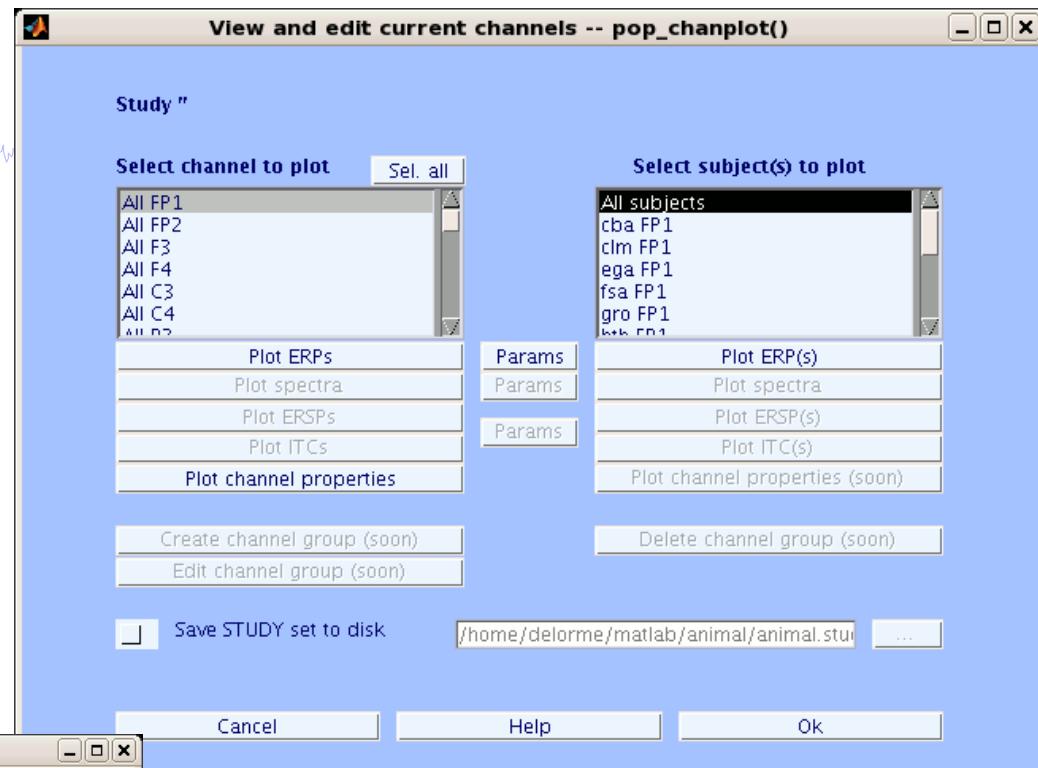
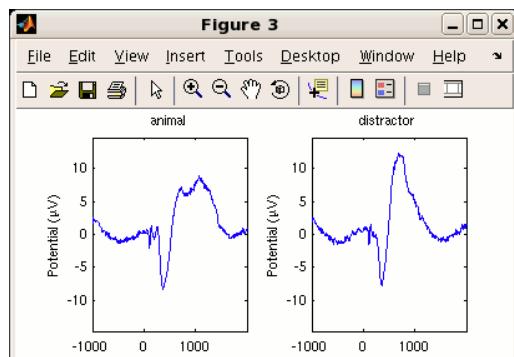


# 4. Analyze clusters





# Channel plotting



# EEGLAB standard processing pipeline



## Single subject

1. Import binary data, events and channel location
2. Edit, Re-reference, Resample, High pass filter data
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  - Analyze components contribution to spectrum

## Multi-subjects

1. Build study
2. Pre-compute measures
3. Cluster components
4. Analyze clusters

**Advanced analysis using scripting and EEGLAB command line functions**

# EEGLAB Data Structures



1. **EEG** - root 'dataset' structure
  - .data - the dataset data (2-D, 3-D matrix)
  - .chanlocs - channel locations substructure
  - .event - data events substructure
  - .epoch - data epochs substructure
2. **ALLEEG** - vector of loaded EEG datasets
3. **CURRENTSET** - index in ALLEEG of current EEG dataset
4. **STUDY** - root 'studyset' structure
  - .cluster - component clustering substructure



# EEG structure

EEG =

```
setname:'Epoched from "ee114 continuous"'  
filename:'ee114squaresepochs.set'  
filepath:'/home/arno/ee114/'  
pnts:384  
nbchan:32  
trials:80  
srate:128  
xmin:-1  
xmax:1.9922  
data:[32x384x80 double]  
icawinv:[32x32 double]  
icasphere:[32x32 double]  
icaweights:[32x32 double]  
icaact:[32x384x80 double]  
event:[1x157 struct]  
epoch:[1x80 struct]  
chanlocs:[1x32 struct]  
comments:[8x150 char]  
averef:'no'  
rt[]  
eventdescription:{1x5 cell}  
epochdescription:{}  
specdata:{}  
specicaact:{}  
reject:[1x1 struct]  
stats:[1x1 struct]  
splinefile:{}  
ref:'common'  
history:[7x138 char]  
urevent:[1x154 struct]  
times:[1x384 double]
```

Number of data points per trial  
Number of channels  
Number of trials  
Sampling rate  
Time limits  
Data  
ICA scalp maps  
ICA activity  
Epoch/event information  
Channel location

# 3 levels of functions



Administrative functions: handle EEG and ALLEEG structures

`eeglab()`, `eeg_checkset()`, `pop_delset()`, ...

Pop functions: interactive functions using EEG structure

`pop_erpimage()`, `pop_topoplot()`, `pop_envtopo()`, ...

Signal processing functions: perform signal processing

`erpimage()`, `topoplot()`, `envtopo()`, ...



# 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...



# Future directions



- Signal processing and source localization
- Analysis of large studies and parallel processing
- Multi-modality imaging
- Study design
- Improved memory mapping features
  
- Improved plug-in facility and script library
- Shared data resource (BIRN)
- Better binary format handling
- Wiki documentation <http://sccn.ucsd.edu/wiki/EEGLAB>
- Open source community development (CVS server)

