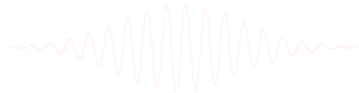


# EEGLAB Processing

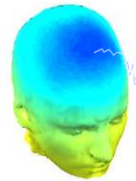
Data import/Preprocessing

Basic ERP visualization

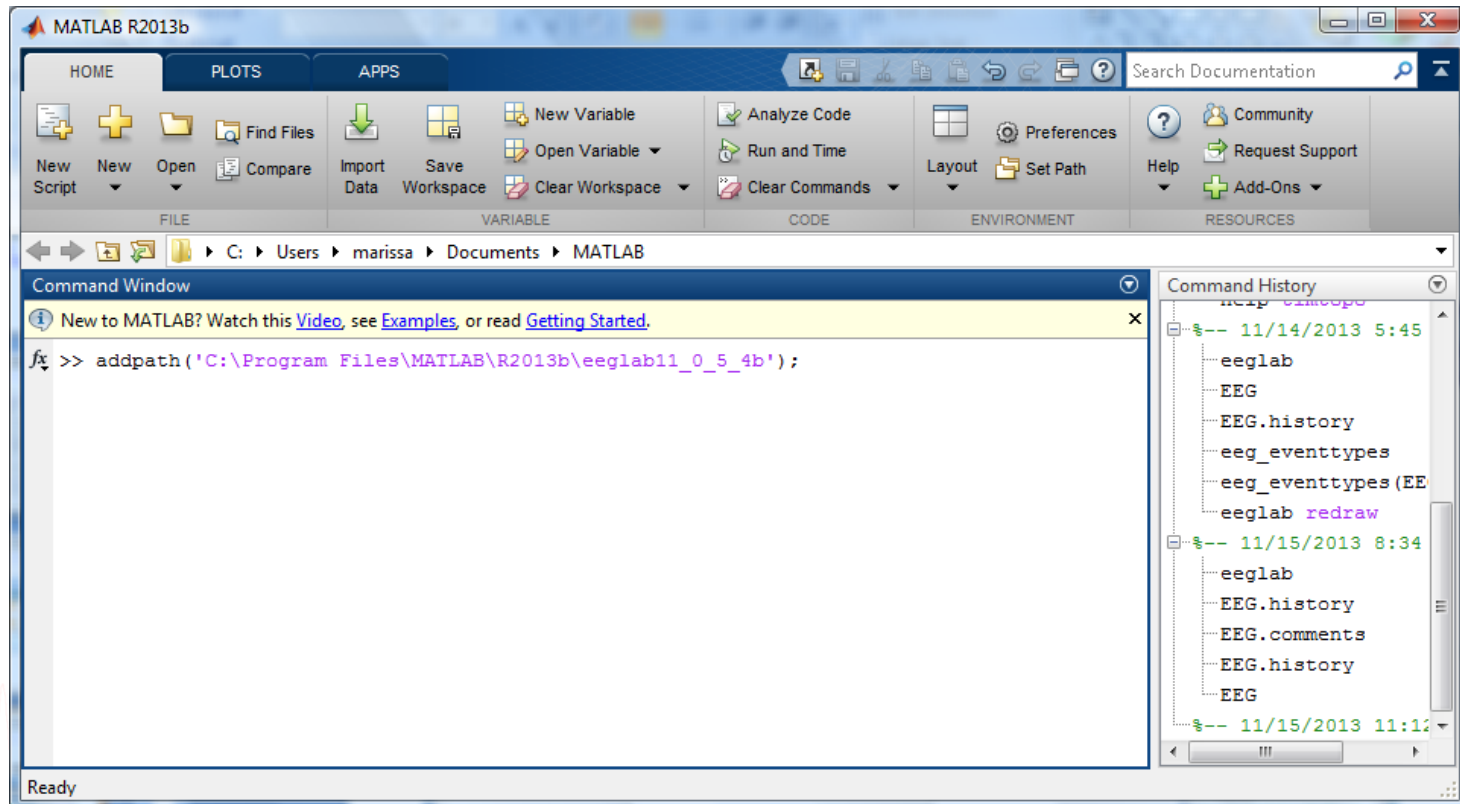




# Installing EEGLAB and data folder

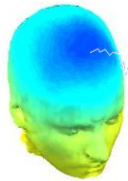


- Start Matlab
- Add the EEGLAB folder to your Matlab path:

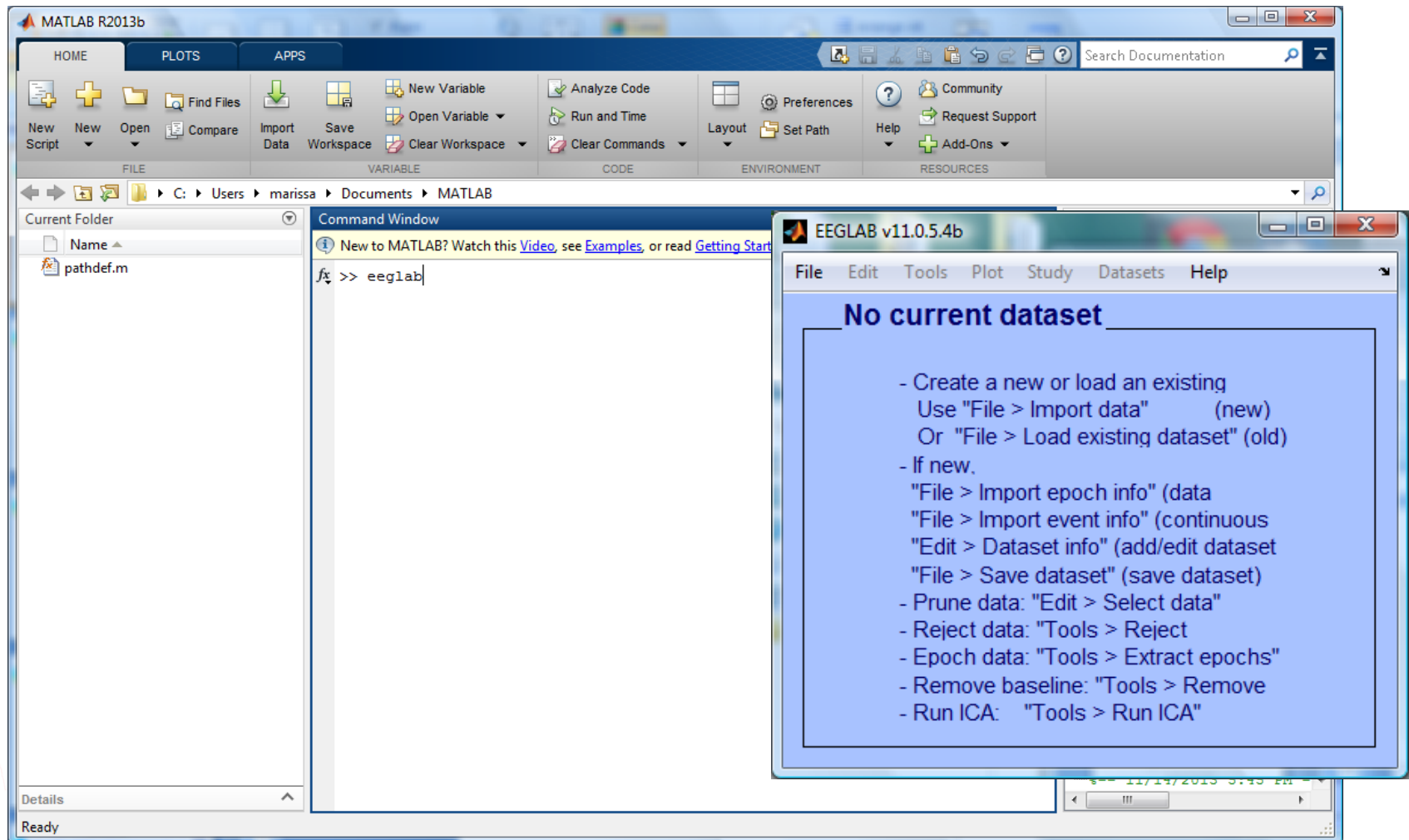




# The EEGLAB Matlab software



## main graphic interface





# Sample data: basic P300 paradigm



## File

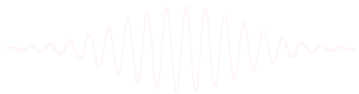
SimpleOddball.set

## Data

68 channel EEG, 256 Hz sampling rate, Biosemi system, re-referenced during import to averaged left and right mastoid electrodes

## Task

speeded button press response to star shape (no response to circle shape), 100 ms presentation duration, 200 trials





# Pre-processing pipeline



**Collect high-density  
EEG data (>30 chan)**

**Import into EEGLAB**

**Import event markers  
and channel locations**

**Re-reference/  
down-sample  
(if necessary)**

**High pass filter  
(~.5 – 1 Hz)**

**Examine raw data**

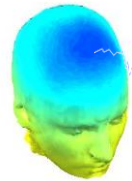
**Identify/reject  
bad channels**

**Reject large artifact  
time points**

**Run ICA**



# Importing a dataset



EEGLAB v11.0.5.4b

File Edit Tools Plot Study Datasets Help

Import data

- Using EEGLAB functions and plugins
- Using the FILE-IO interface
- Using the BIOSIG interface
- Troubleshooting data formats...

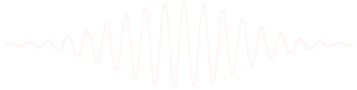
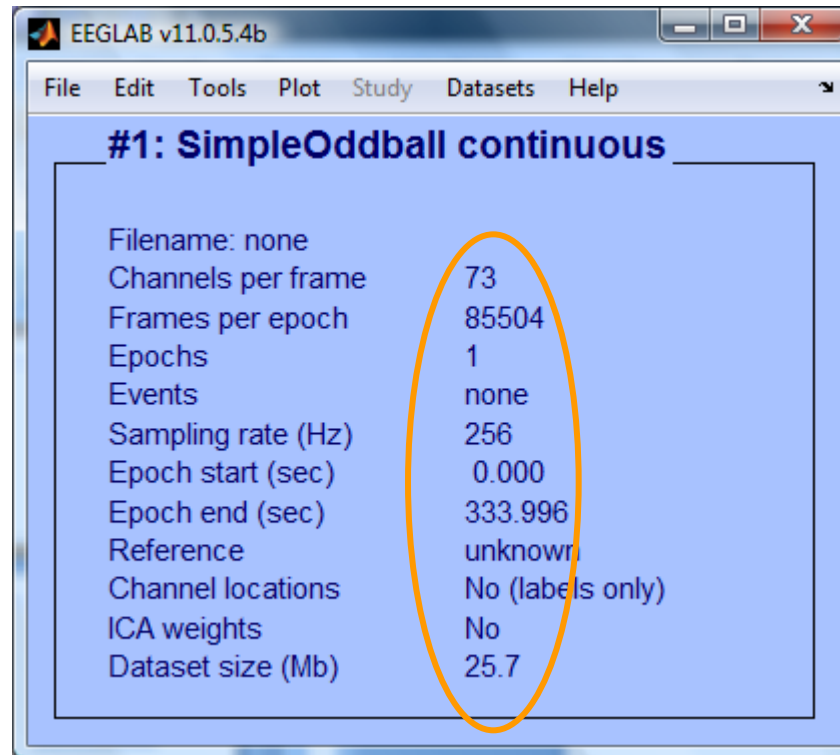
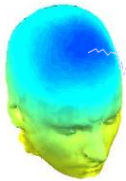
Load existing dataset

- From ASCII/float file or Matlab array
- From Netstation .mff (FILE-IO toolbox)
- From Netstation binary simple file
- From Multiple seg. Netstation files
- From Netstation Matlab files
- From BCI2000 ASCII file
- From Snapmaster .SMA file
- From Neuroscan .CNT file
- From Neuroscan .EEG file
- From Biosemi BDF file (BIOSIG toolbox)
- From Biosemi BDF and EDF files (BDF plugin)**
- From EDF/EDF+/GDF files (BIOSIG toolbox)
- From ANT EEProbe .CNT file
- From ANT EEProbe .AVR file
- From BCI2000 .DAT file
- From BIOPAC MATLAB files
- From Brain Vis. Rec. .vhdr file
- From Brain Vis. Anal. Matlab file
- From CTF folder (MEG)
- From ERPSS .RAW or .RDF file
- From INStep .ASC file
- From 4D .m4d pdf file
- From Procom Infinity Text File

Tip for Biosemi users:  
Use the 'BDF plugin' version  
of the Biosemi BDF/EDF importer



# Imported EEG data





# Pre-processing pipeline



**Collect high-density  
EEG data (>30 chan)**

**Import into EEGLAB**

**Import event markers  
and channel locations**

**Re-reference/  
down-sample  
(if necessary)**

**High pass filter  
(~.5 – 1 Hz)**

**Remove line noise  
(if necessary)**

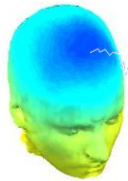
**Identify/reject  
bad channels**

**Reject large artifact  
time points**

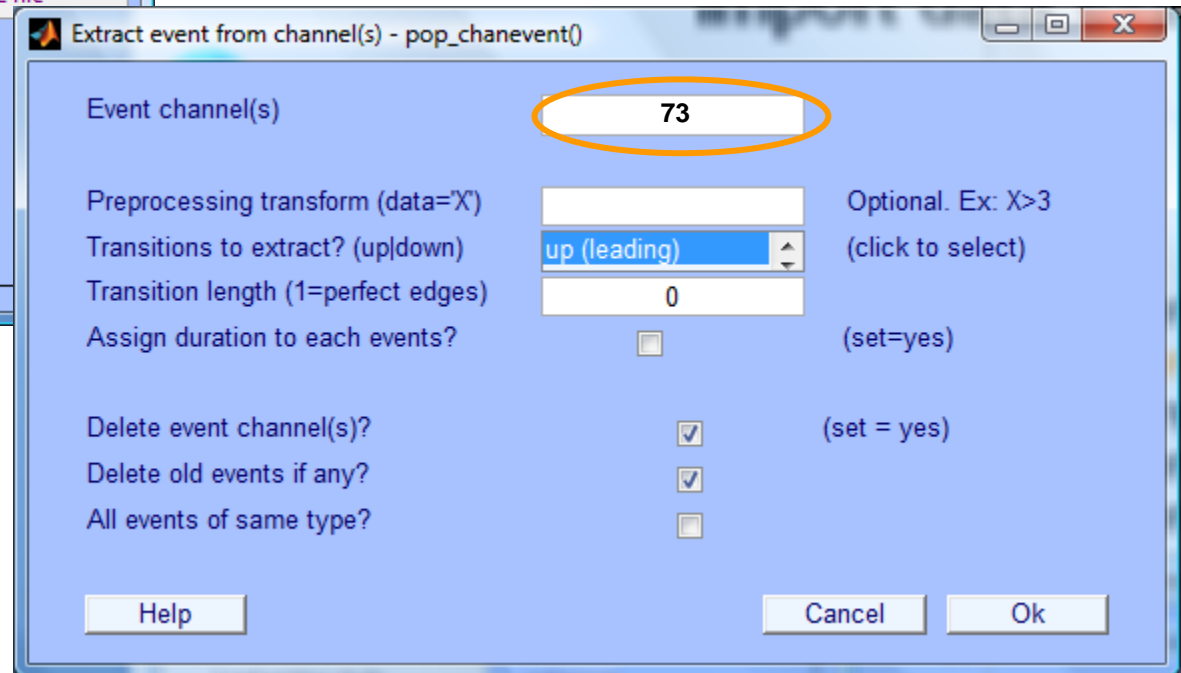
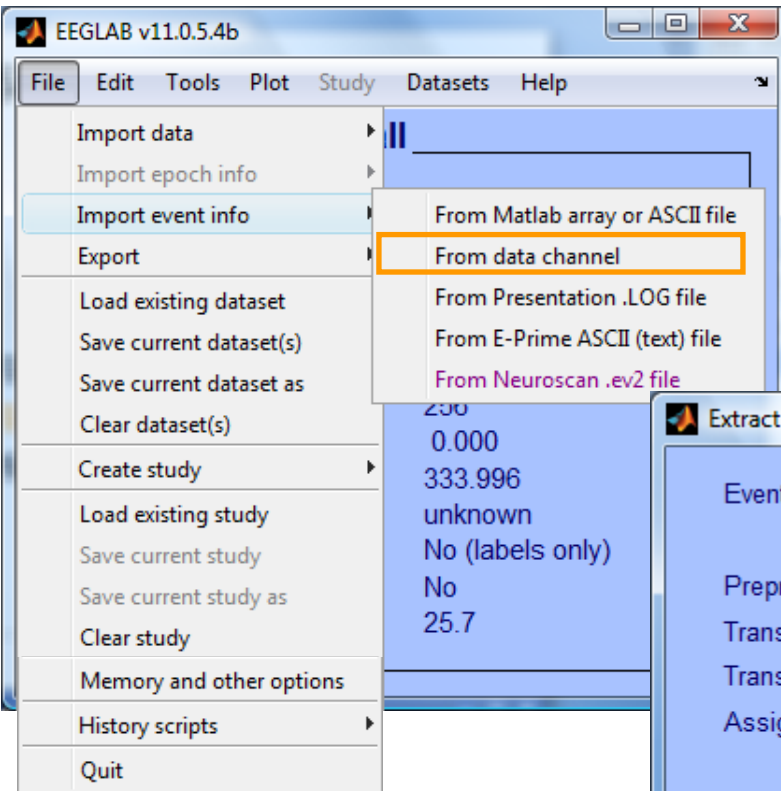
**Run ICA**



# Import data events

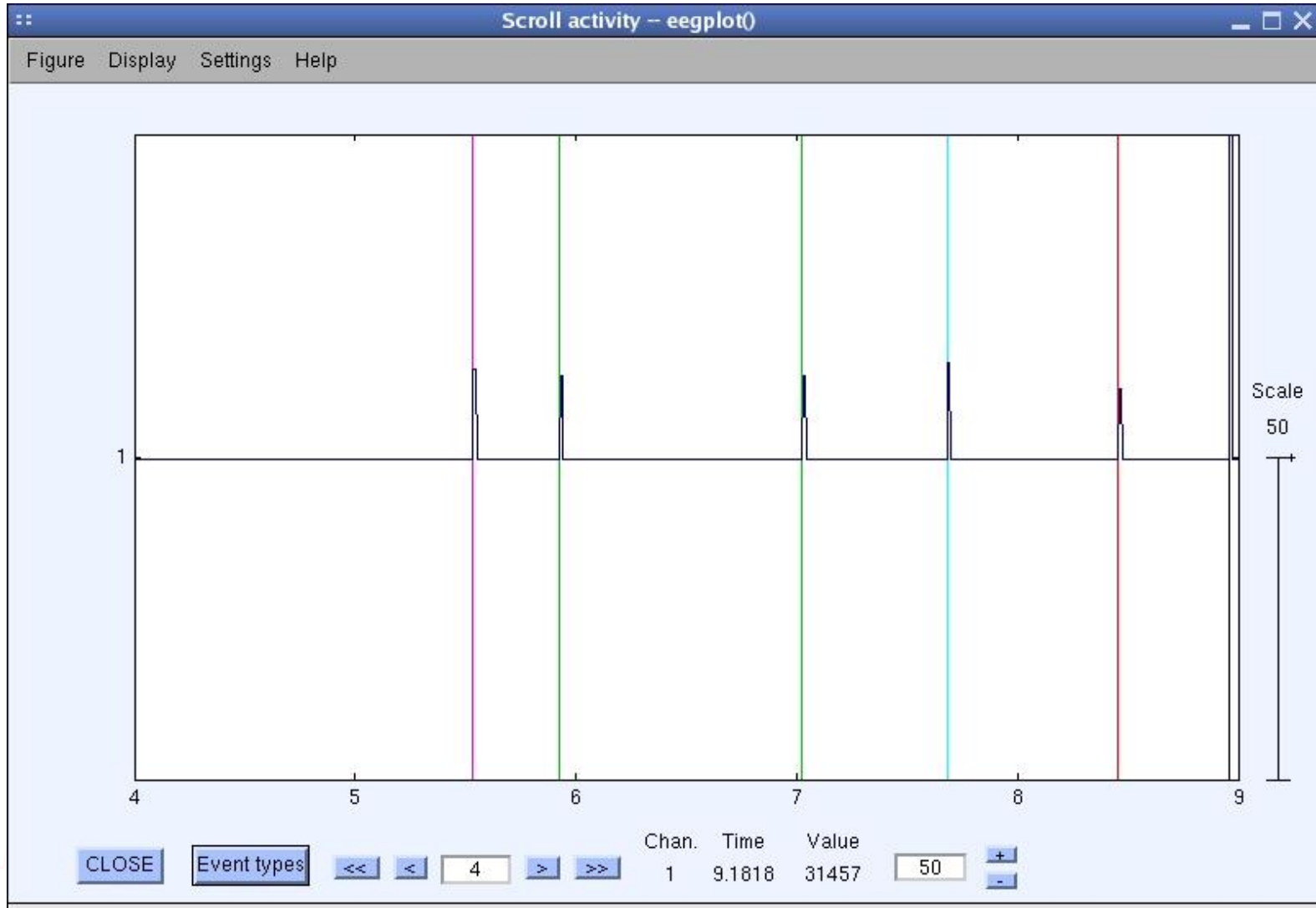
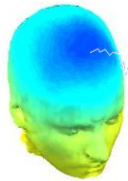


(Often imported automatically during data import)



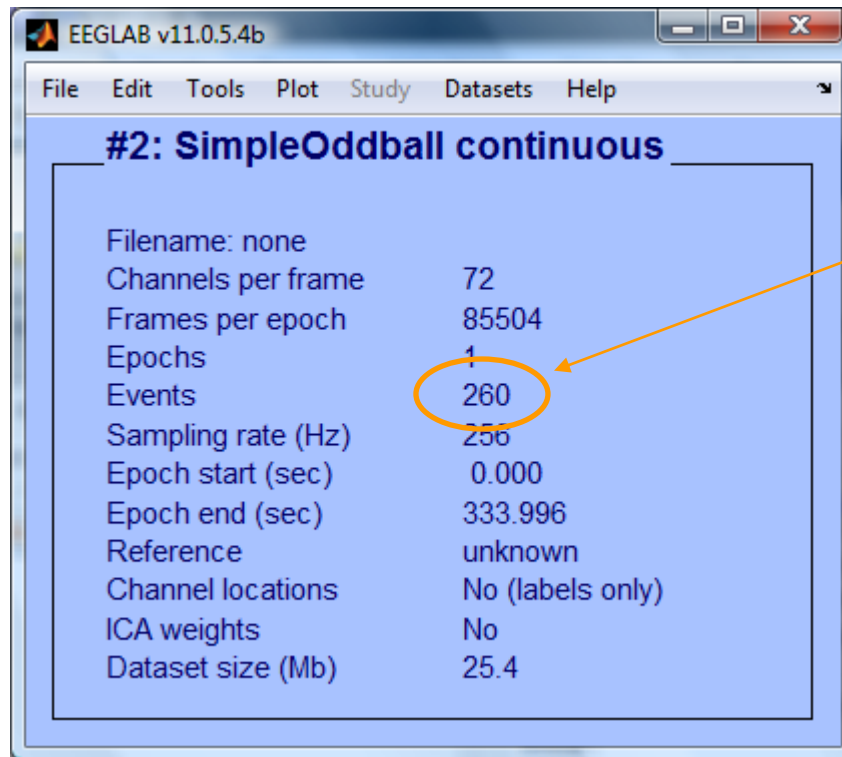
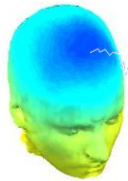


# Appearance of an event channel in raw data

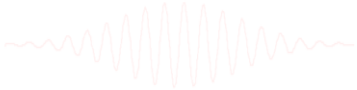




# Imported data events

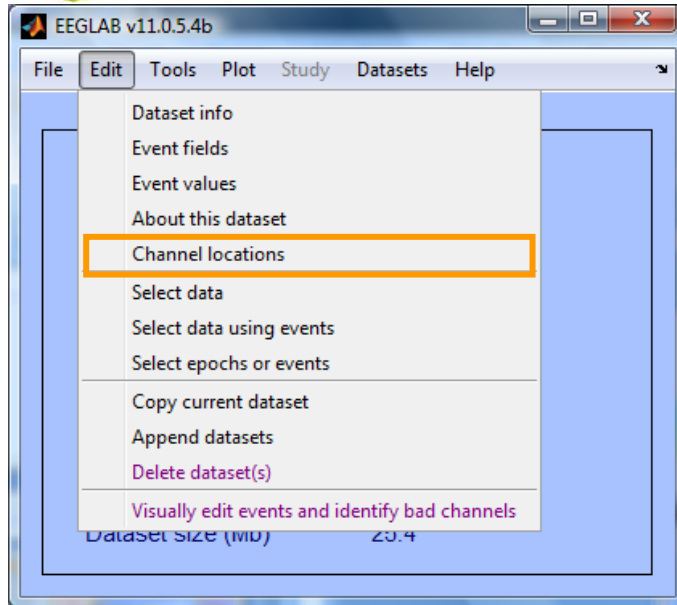
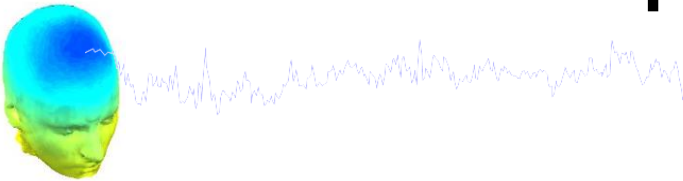


If event import was successful, you will see an appropriate number here



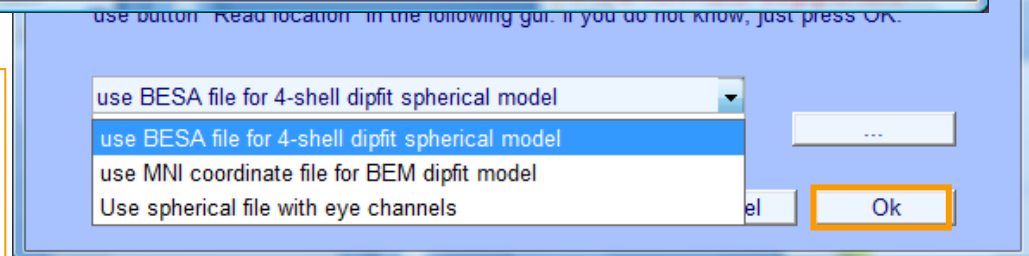
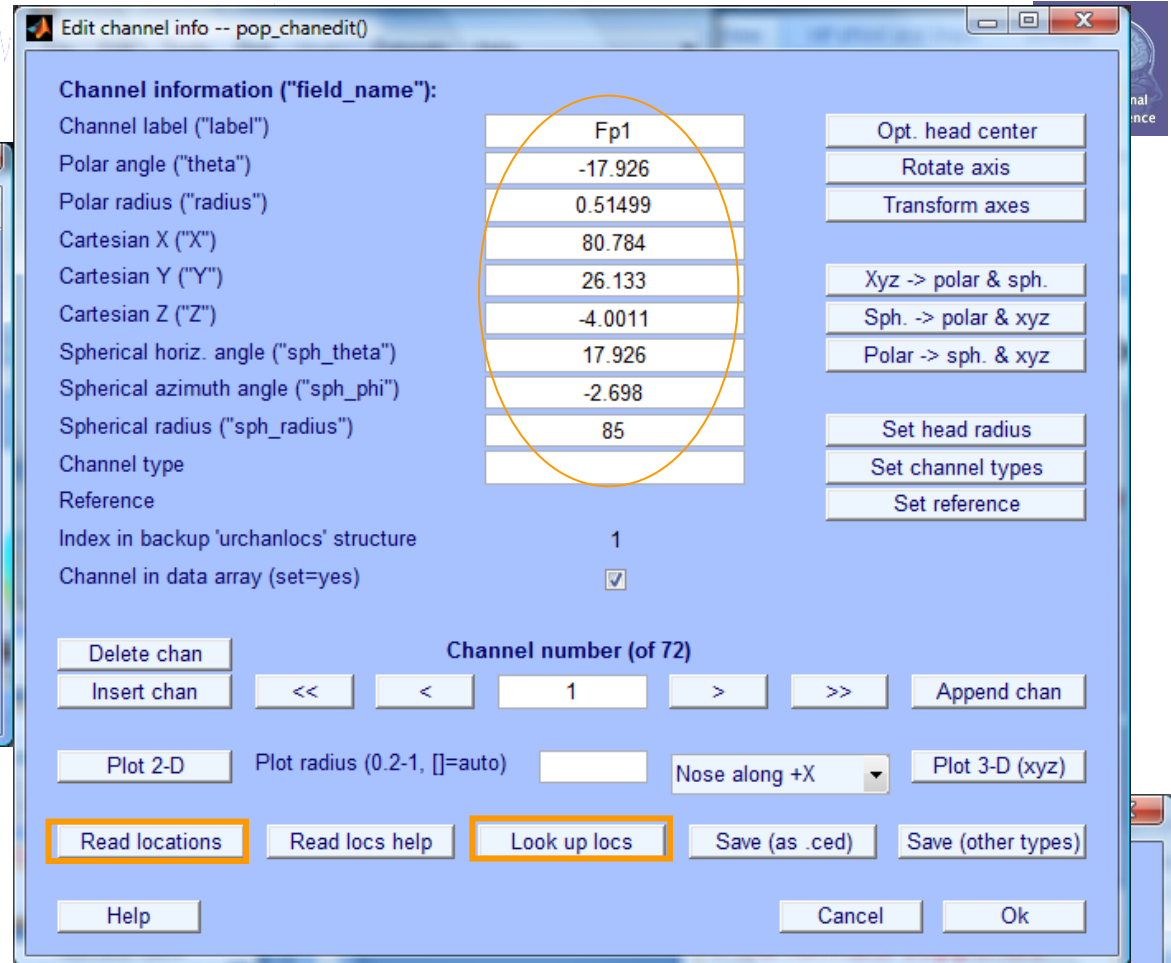


# Import channel locations



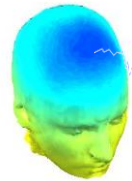
7 file formats supported  
(Polhemus, BESA, ...)

Standard locations offered  
if you do not have locations  
and labels are standard





# Import channel locations



**Edit channel info -- pop\_chanedit()**

**Channel information ("field\_name"):**

Channel label ("label")	LEYE
Polar angle ("theta")	-45.1543
Polar radius ("radius")	0.54374
Cartesian X ("X")	0.79487
Cartesian Y ("Y")	0.79917
Cartesian Z ("Z")	-0.15585
Spherical horiz. angle ("sph_theta")	45.1543
Spherical azimuth angle ("sph_phi")	-7.8725
Spherical radius ("sph_radius")	1.1379
Channel type	EEG
Reference	
Index in backup 'urchanlocs' structure	
Channel in data array (set=yes)	<input checked="" type="checkbox"/>

**Channel number (of 71)**

1

**Buttons:** Delete chan, Insert chan, <<, <, >, >>, Append chan, Plot 2-D, Plot radius (0.2-1, []=auto), Nose along +X, Plot 3-D (xyz), Read locations, Read locs help, Look up locs, Save (as .ced), Save (other types), Help, Cancel, Ok

**Opt. head center**  
**Rotate axis**  
**Transform axes**

XYZ -> polar & sph.  
Sph. -> polar & xyz  
Polar -> sph. & xyz

Set head radius  
Set channel types  
Set reference

**Convert channel locations -- pop\_chancenter()**

**Optimize center location** ☒ or specify center 0 0 0

Channel indices to ignore for best-sphere matching

Help Cancel Ok

**Force electrode location -- forclocs()**

X/Y value Coordinate Electrode list

0 X (rotate X-Z plane) Cz Pick

Help Cancel Ok

**Set channel ...**

Channel indices 1:71

Type (e.g. EEG) EEG

Help Cancel Ok



Edit channel info -- pop\_chanedit()

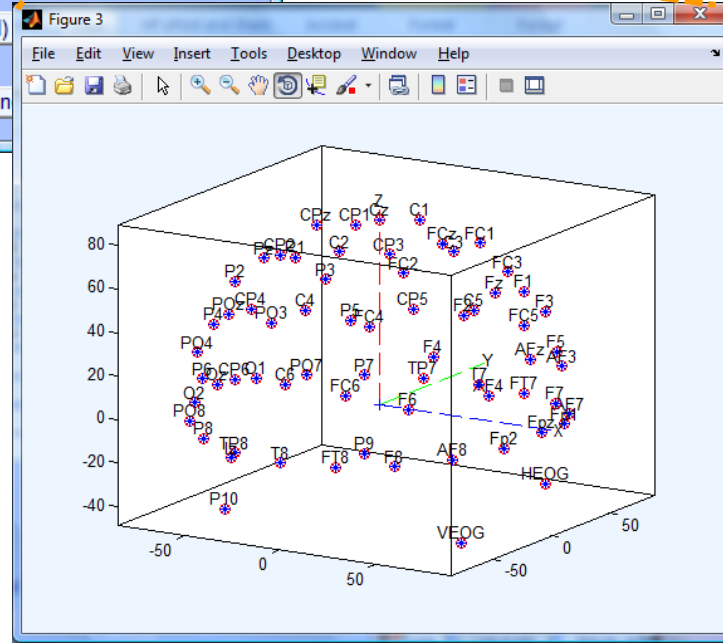
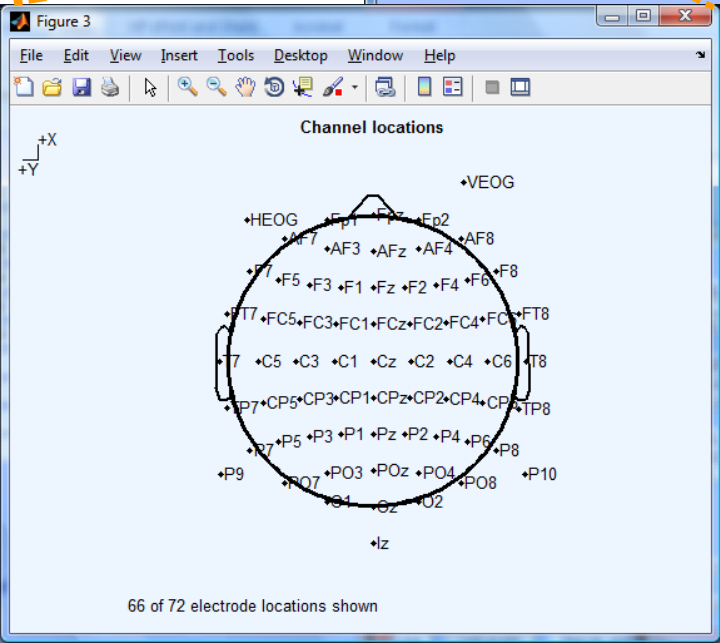
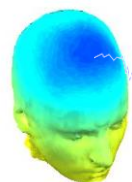
Channel information ("field\_name"):

Channel label ("label")	HEOG	Opt. head center
Polar angle ("theta")	-42	Rotate axis
Polar radius ("radius")	0.65556	Transform axes
Cartesian X ("X")	55.7734	XYZ -> polar & sph.
Cartesian Y ("Y")	50.2186	Sph. -> polar & xyz
Cartesian Z ("Z")	-39.9051	Polar -> sph. & xyz
Spherical horiz. angle ("sph_theta")	42	Set head radius
Spherical azimuth angle ("sph_phi")	-28	Set channel types
Spherical radius ("sph_radius")	85	Set reference
Channel type		
Reference		
Index in backup 'urchanlocs' structure	68	
Channel in data array (set=yes)	<input checked="" type="checkbox"/>	

Channel number (of 72)

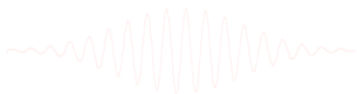
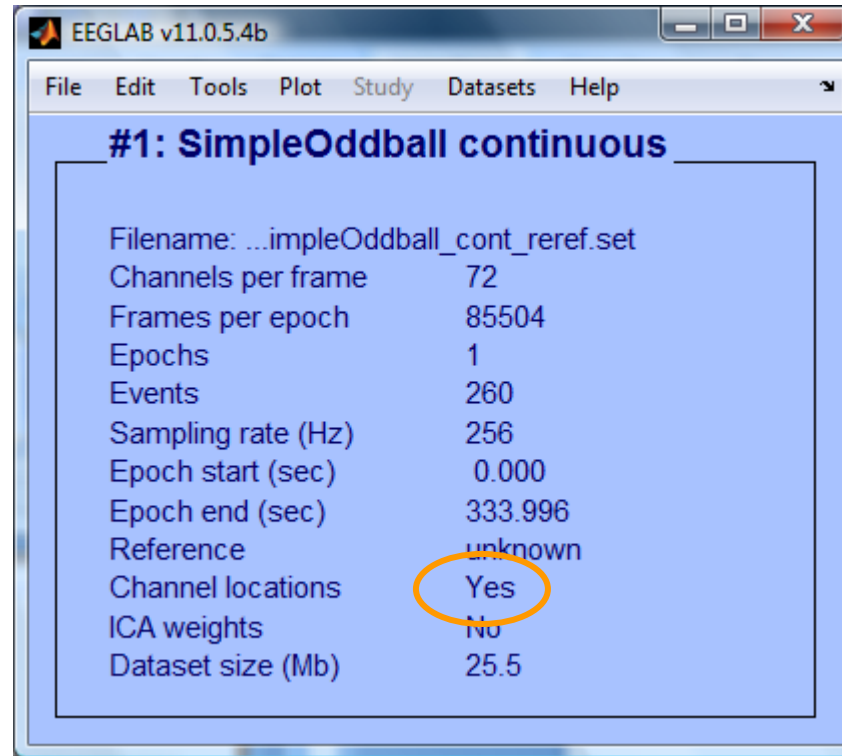
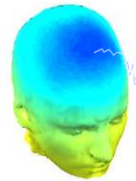
68

Plot 2-D Plot radius (0.2-1, []=auto) Nose along +X Plot 3-D (xyz)





# Imported channel locations





# Pre-processing pipeline



**Collect high-density  
EEG data (>30 chan)**

**Import into EEGLAB**

**Import event markers  
and channel locations**

**Re-reference/  
down-sample  
(if necessary)**

**High pass filter  
(~.5 – 1 Hz)**

**Remove line noise  
(if necessary)**

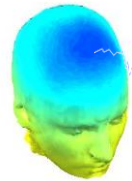
**Identify/reject  
bad channels**

**Reject large artifact  
time points**

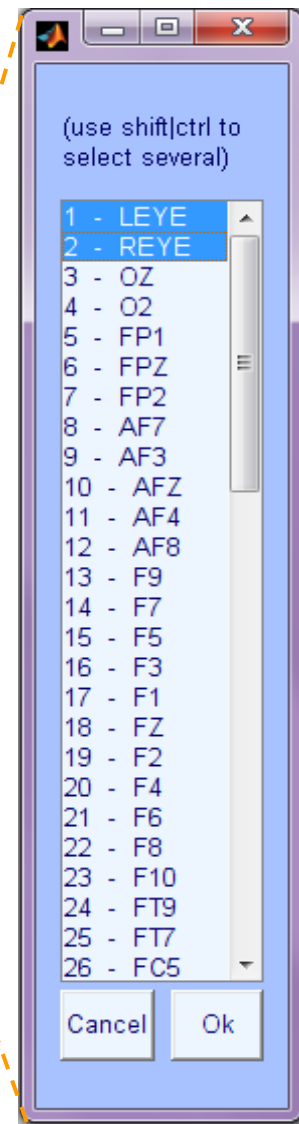
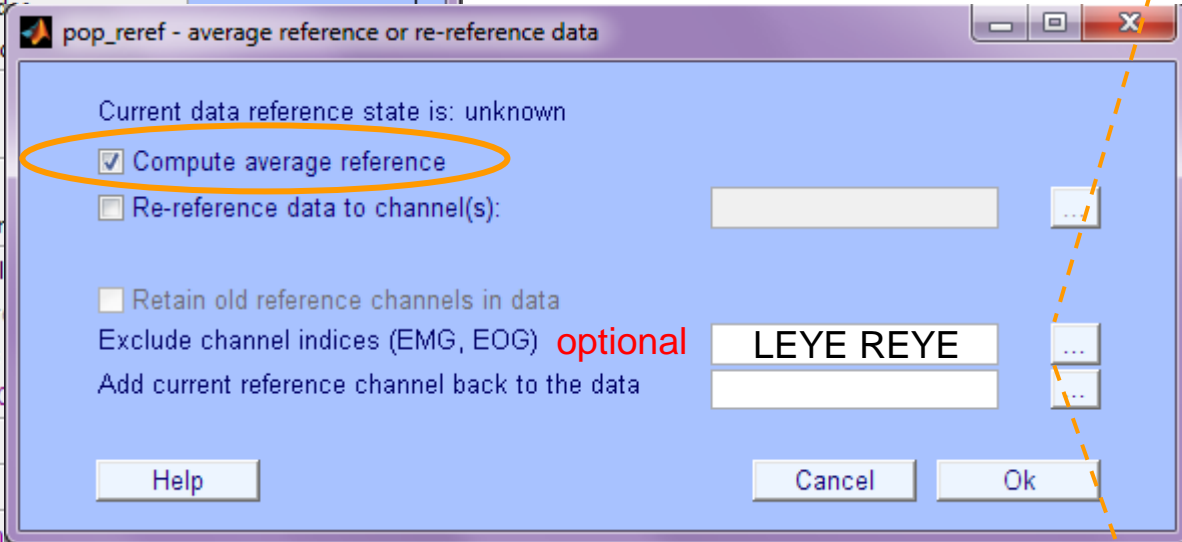
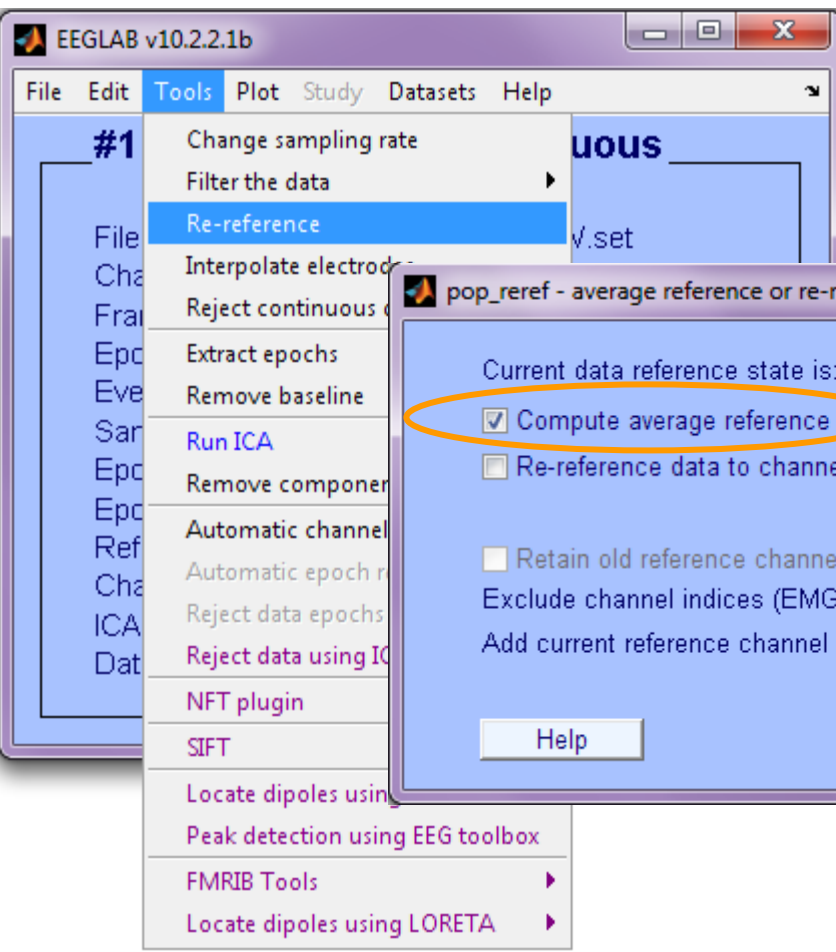
**Run ICA**



# Re-reference data (if necessary/desired)

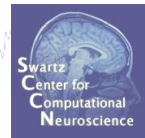
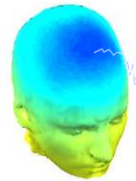


For example,  
average reference

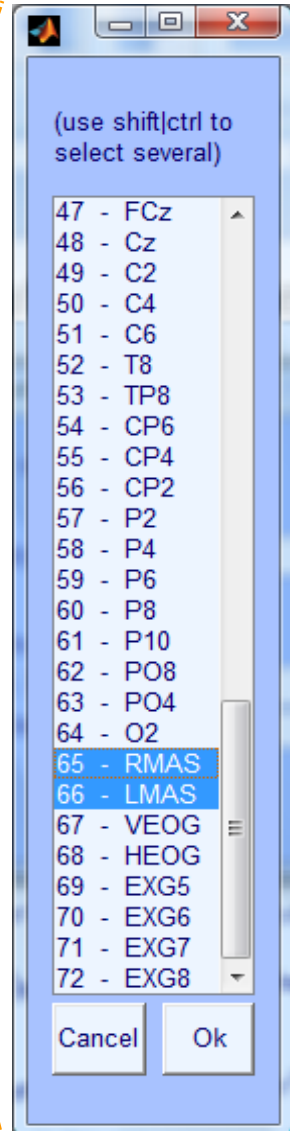
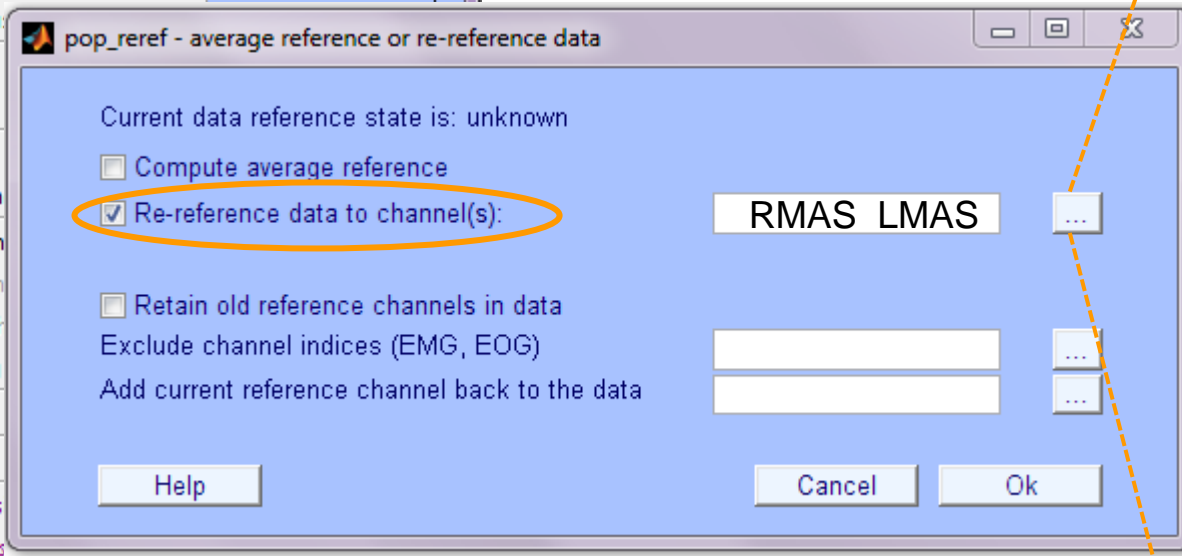
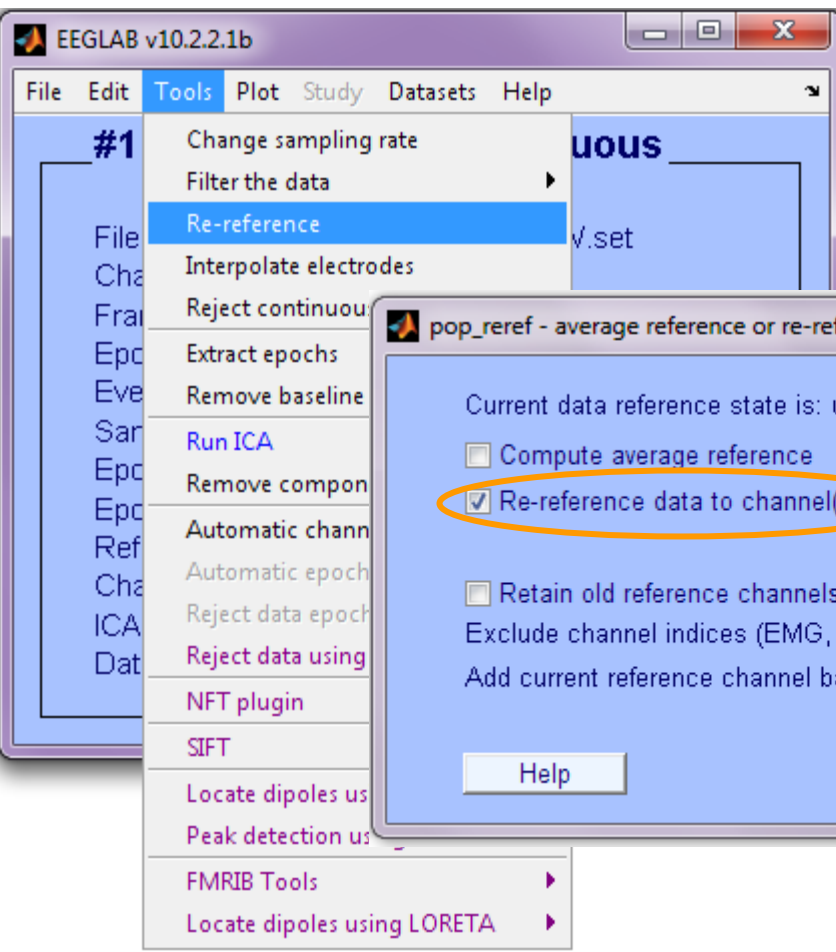




# Re-reference data (if necessary/desired)



OR, re-reference to  
(i.e.) 'linked mastoids'



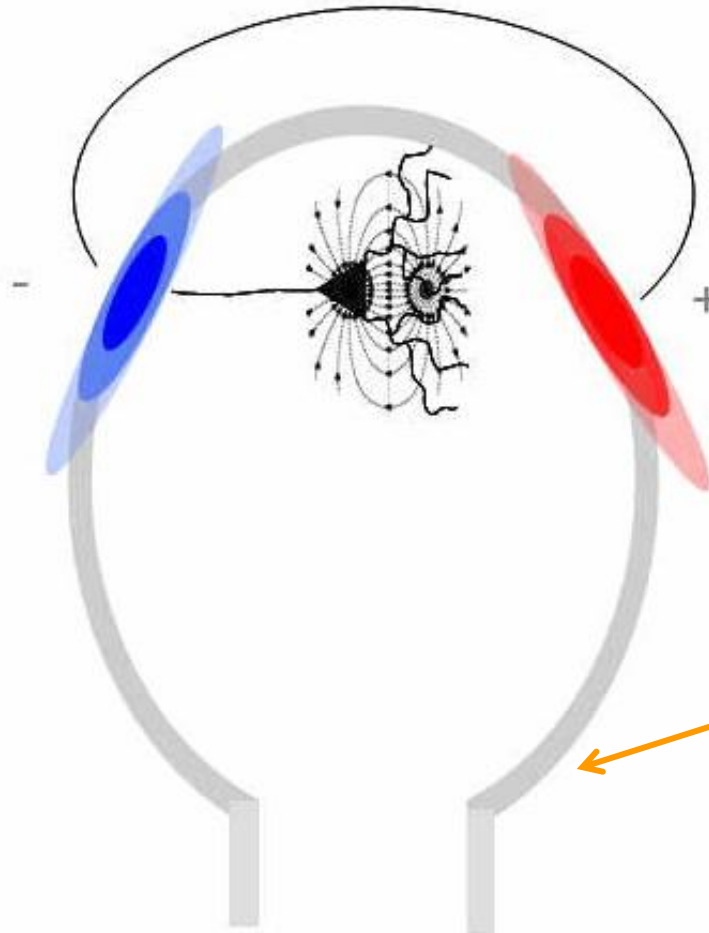


# Choice of reference

The “perfect” reference is probably impossible to find



True average reference requires **even electrode distribution over the head**

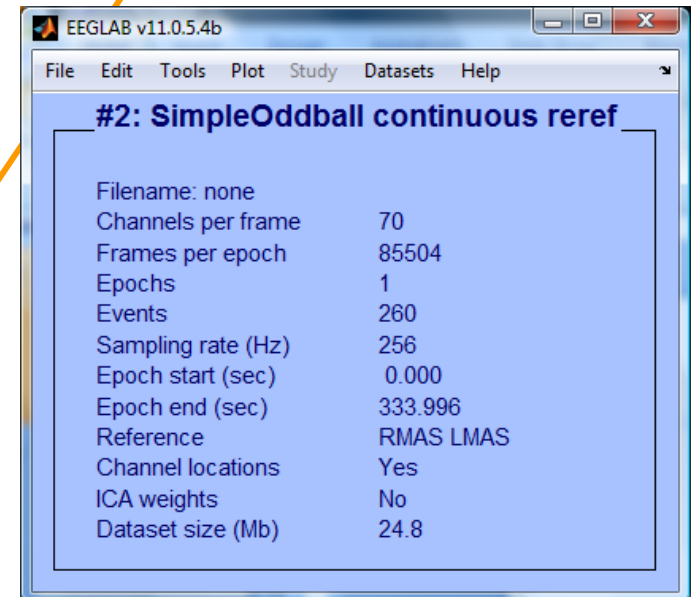
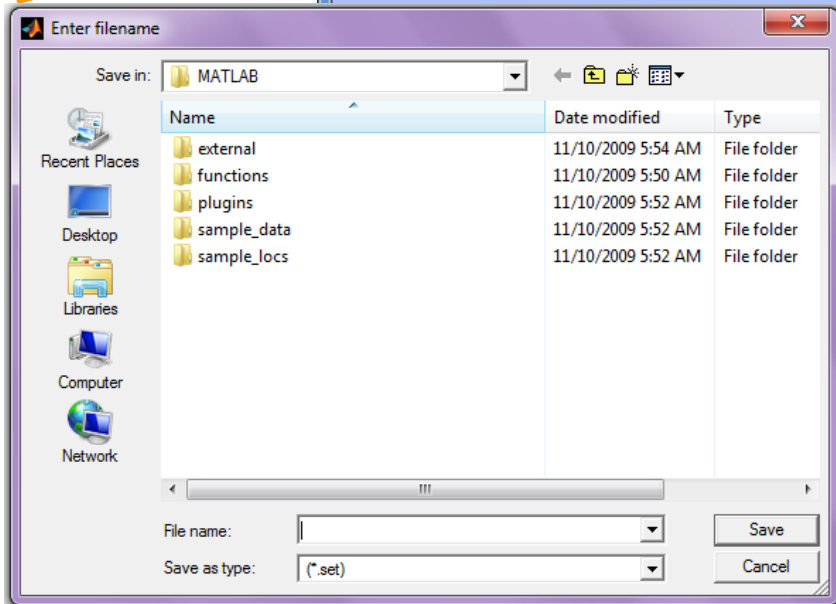
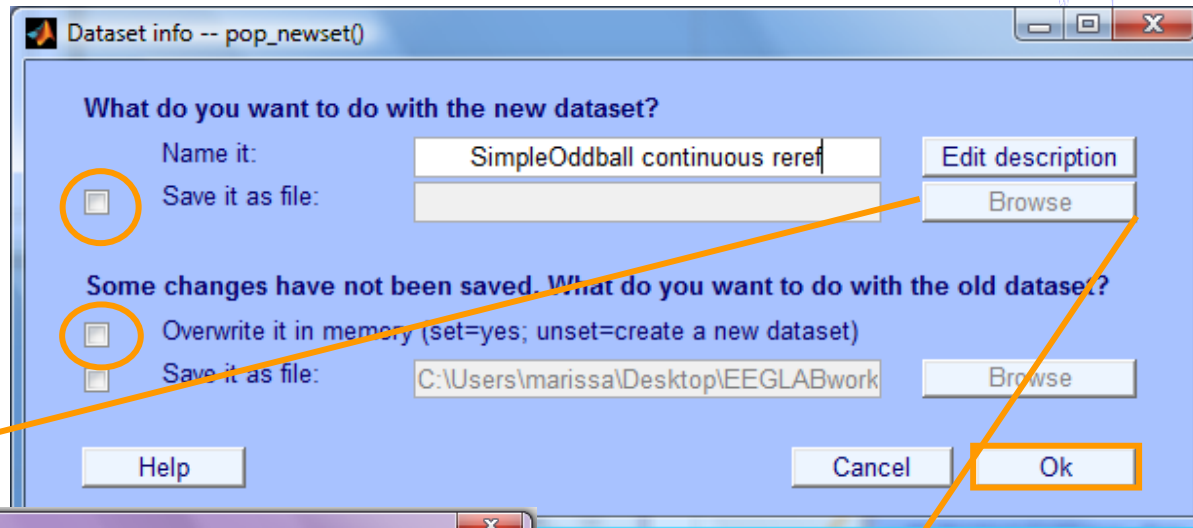


Typical montages have **fewer (if any) on the lower half of the head surface**

**Rationale for average reference:** outward positive and negative currents, summed across an entire (electrically isolated) sphere, will sum to 0 (by Ohm's law)

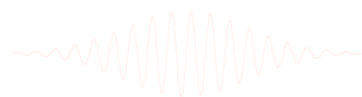
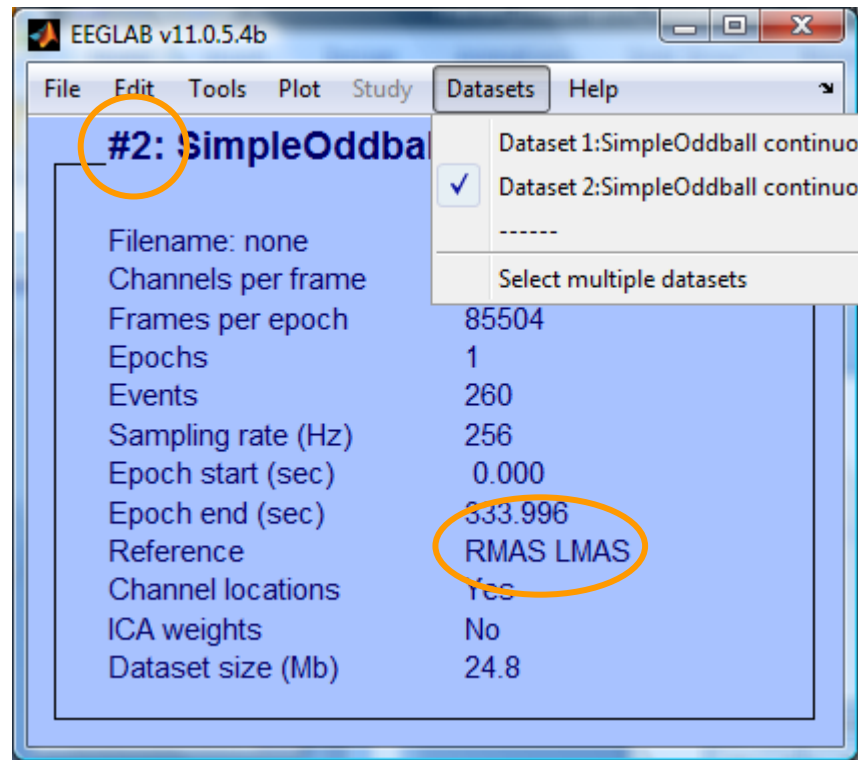
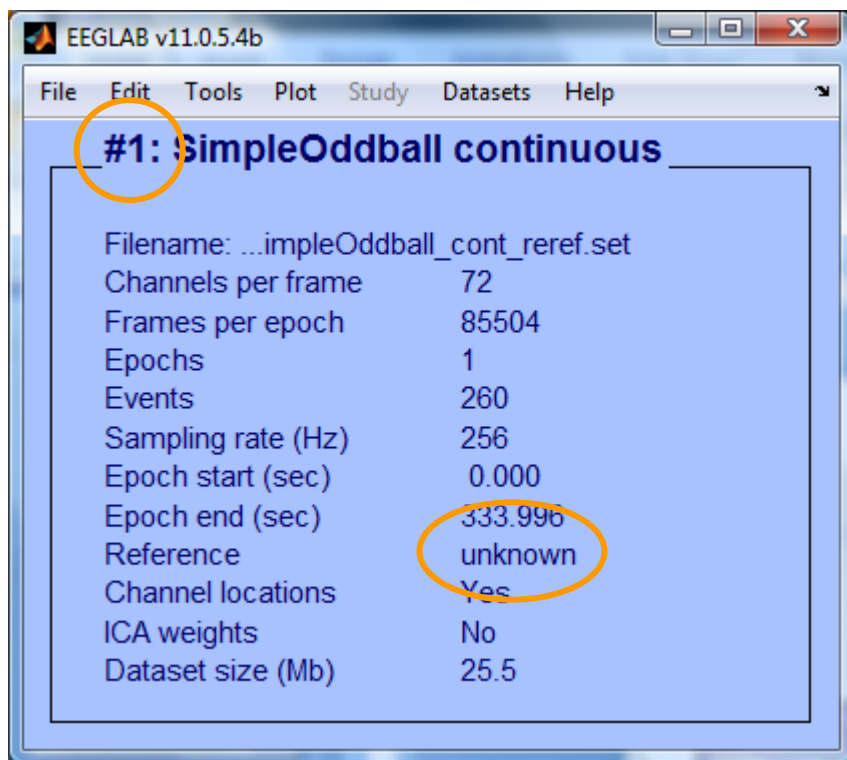


# Save new dataset, keep old one



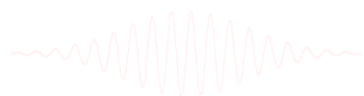
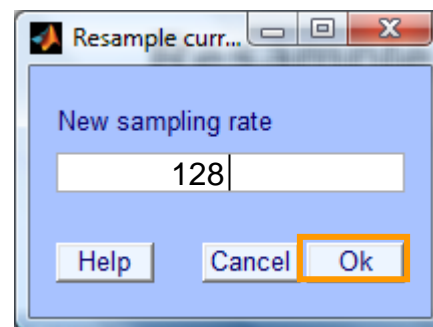
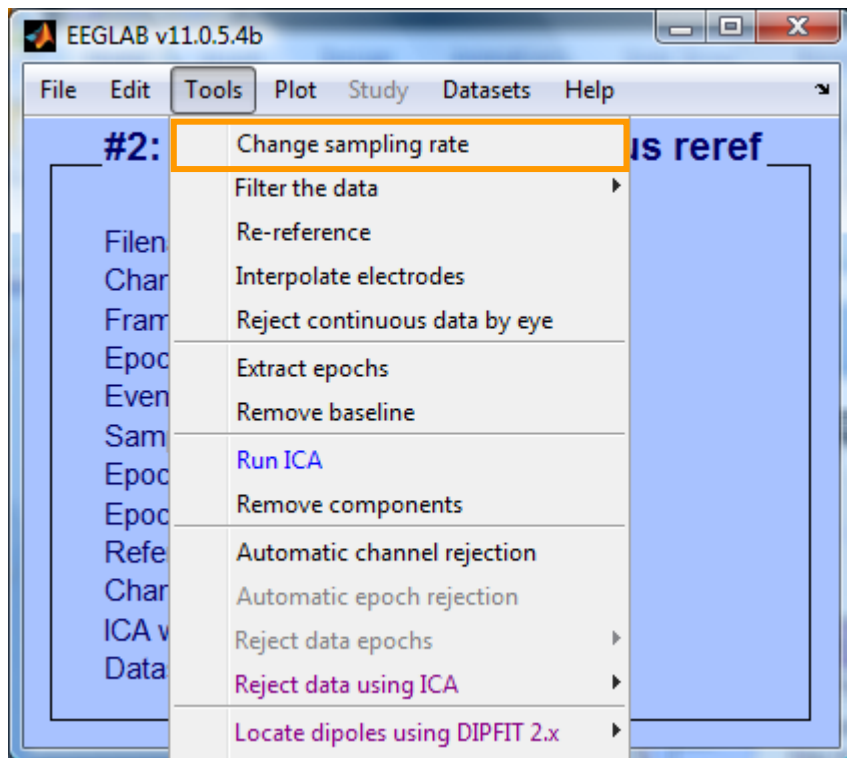


# Multiple active datasets (ALLEEG)





# Resample data (if necessary)





# Pre-processing pipeline



**Collect high-density  
EEG data (>30 chan)**

**Import into EEGLAB**

**Import event markers  
and channel locations**

**Re-reference/  
down-sample  
(if necessary)**

**High pass filter  
(~.5 – 1 Hz)**

**Remove line noise  
(if necessary)**

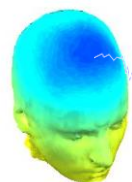
**Identify/reject  
bad channels**

**Reject large artifact  
time points**

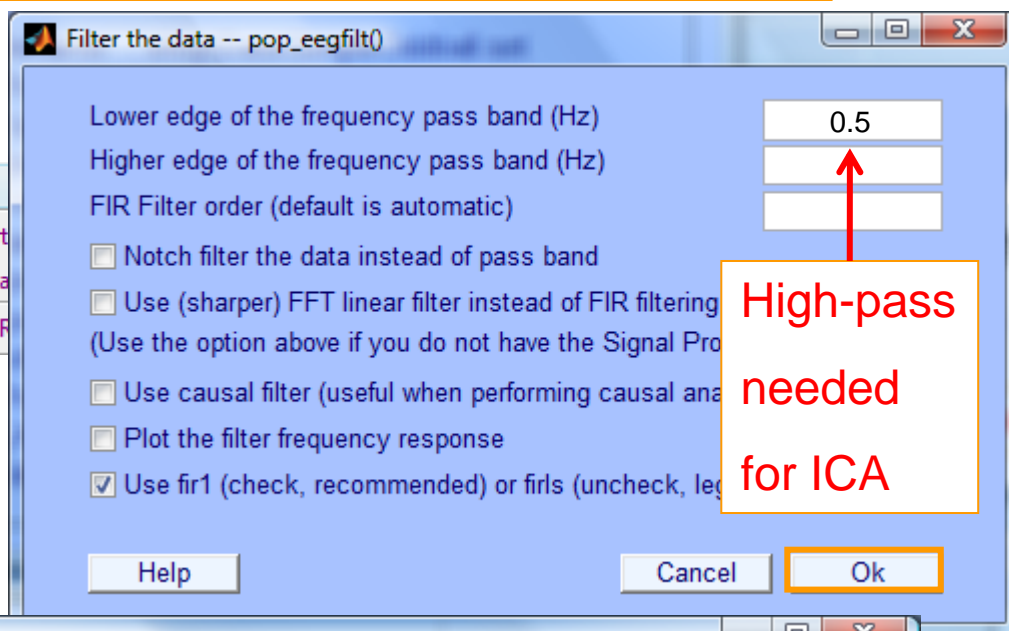
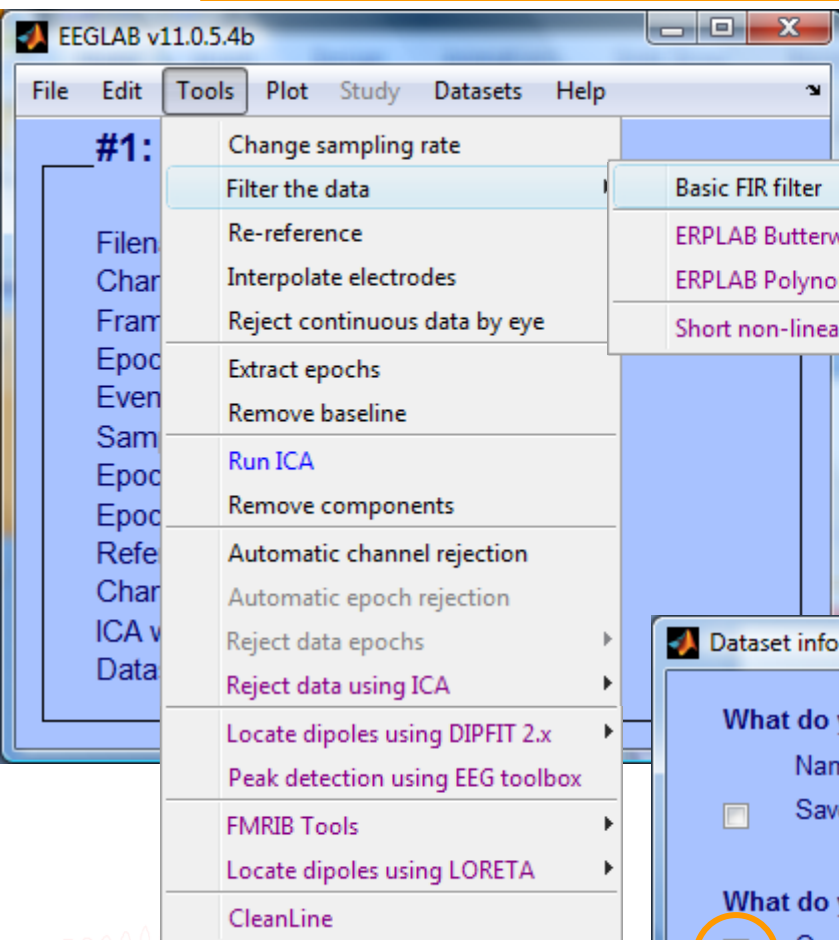
**Run ICA**



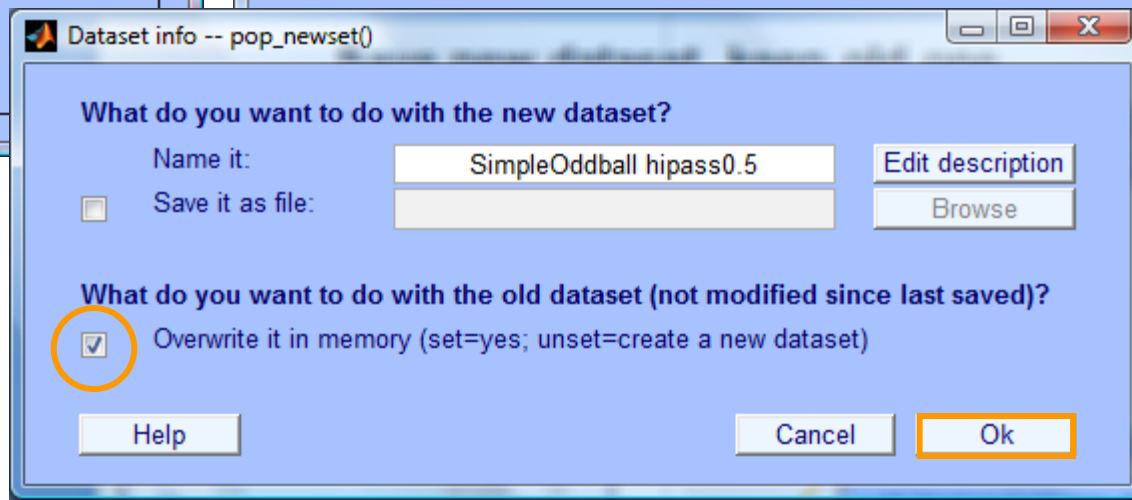
# Filter the data (if necessary/desired)



Lower cut off frequencies require longer stretches of continuous data



High-pass  
needed  
for ICA





# Pre-processing pipeline



**Collect high-density  
EEG data (>30 chan)**

**Import into EEGLAB**

**Import event markers  
and channel locations**

**Re-reference/  
down-sample  
(if necessary)**

**High pass filter  
(~.5 – 1 Hz)**

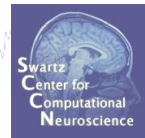
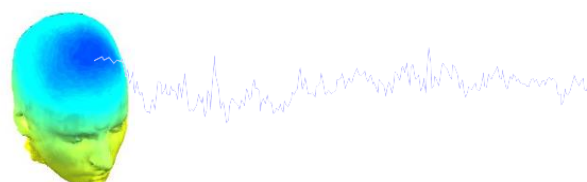
**Remove line noise  
(if necessary)**

**Identify/reject  
bad channels**

**Reject large artifact  
time points**

**Run ICA**





EEGLAB v12.0.2.1b

File Edit Tools Plot Study Datasets Help

- 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
- History scripts
- Manage plugins**
  - Manage data import plugins
  - Manage data processing plugins
  - Manage deactivated plugins
- Quit

Plutings available for install on the internet

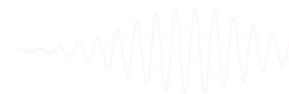
Install	Plugin	Version	Description	
<input type="checkbox"/>	ERPLABfilters	1.00	Interface ERPLAB filters (requires seperate ERPLAB instalati...	Doc
<input type="checkbox"/>	<b>Cleanline</b>	1.21	Automatic artifact rejection	Doc
<input type="checkbox"/>	BERGEN	1.1	Removal of fMRI-related gradient artifacts from simultaneous...	Doc

Update Deactivate

Installed plutings

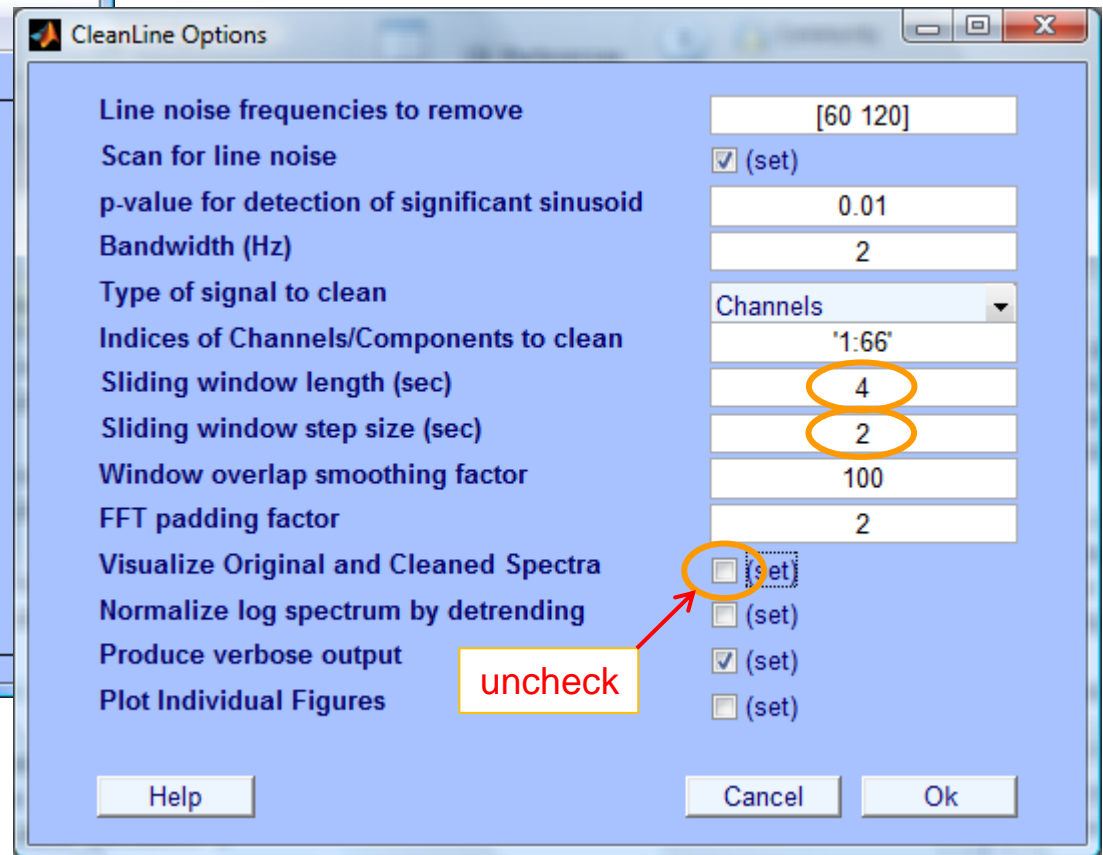
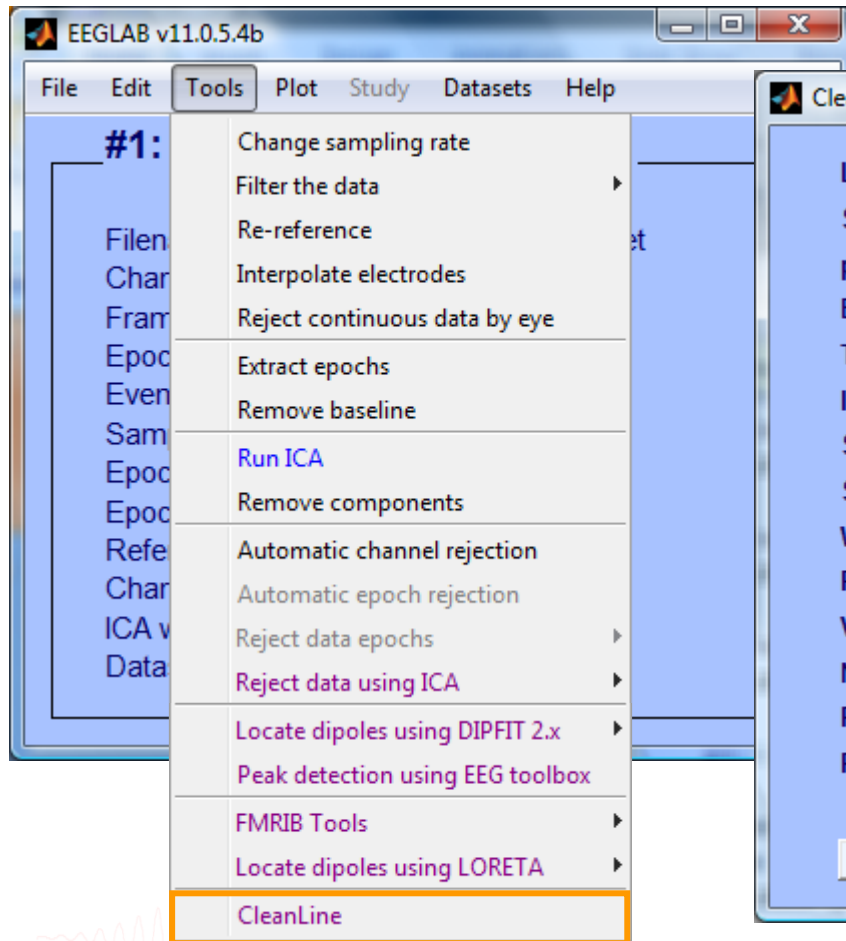
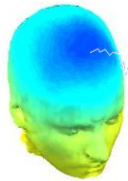
Update	Deactivate	Plugin	Version	Description	
<input type="checkbox"/>	<input type="checkbox"/>	brainmovie	0.1	Brainmovies (command line only)	Doc
<input type="checkbox"/>	<input type="checkbox"/>	corrmap	2.00	New version 1.03 available. Click update to install.	Doc
<input type="checkbox"/>	<input type="checkbox"/>	eeg_toolbox	1.0	Interface EEG toolbox functions for ERP peak detection	Doc
<input type="checkbox"/>	<input type="checkbox"/>	fMRIb	1.21	Remove fMRI artifacts from EEG	Doc
<input type="checkbox"/>	<input type="checkbox"/>	MP_clustering	1.00	Measure projection clustering of ICA components	Doc
<input type="checkbox"/>	<input type="checkbox"/>	MutualInfoClustering	1.00	Mutual information clustering	Doc
<input type="checkbox"/>	<input type="checkbox"/>	StudyEnvtopo	0.9	Add envtopo capabilities to STUDY	Doc
<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	VisEd	1.05	New version 1.04 available. Click update to install.	Doc
<input type="checkbox"/>	<input type="checkbox"/>	iirfilt	1.02	Non linear filtering	Doc
<input type="checkbox"/>	<input type="checkbox"/>	loreta	1.1	New version 1.0 available. Click update to install.	Doc

Cancel Ok



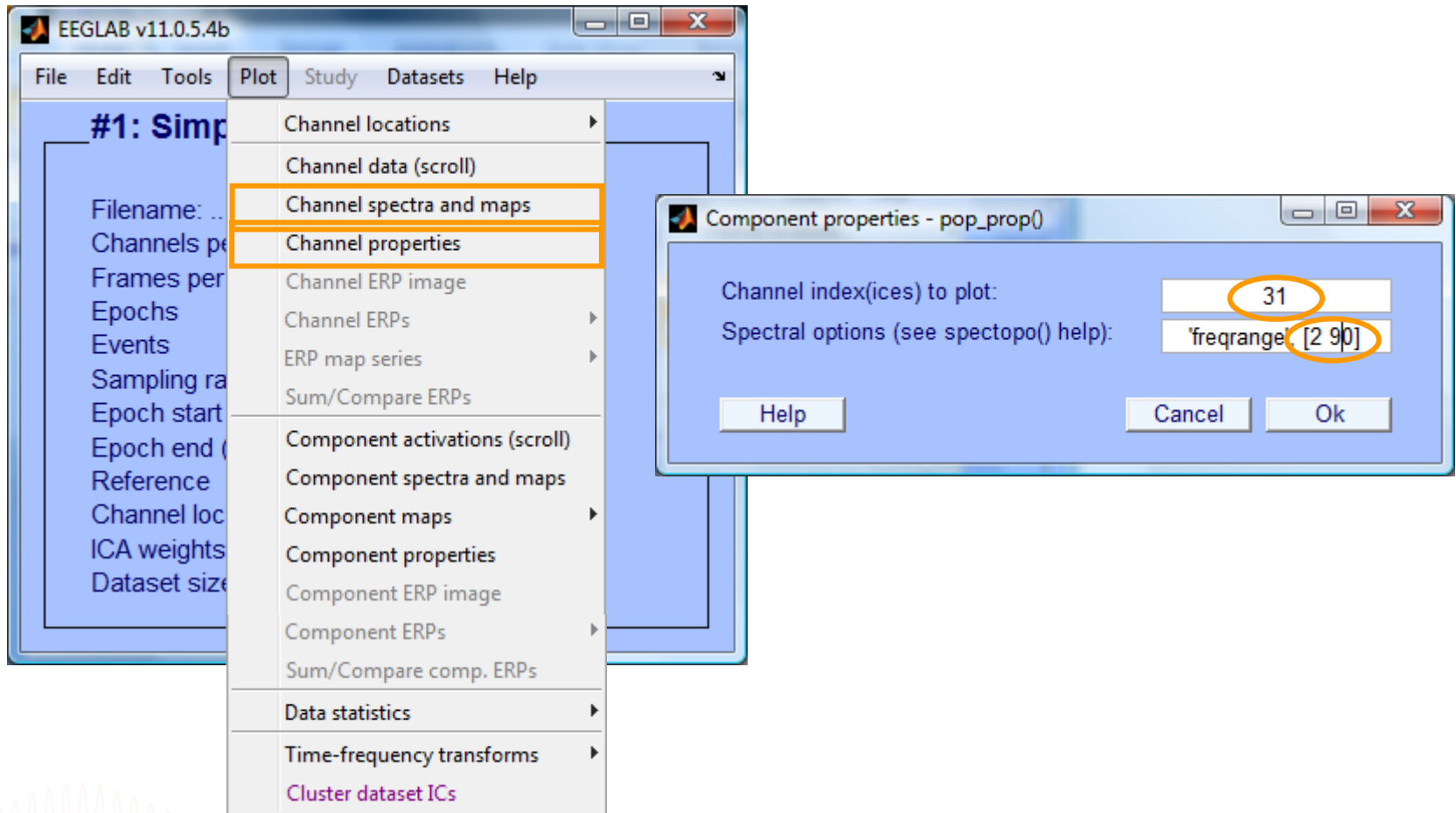
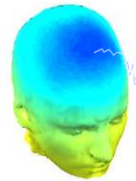


# Remove line noise (Cleanline)





# Plot channel spectra





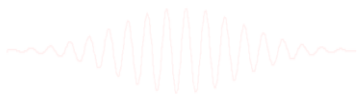
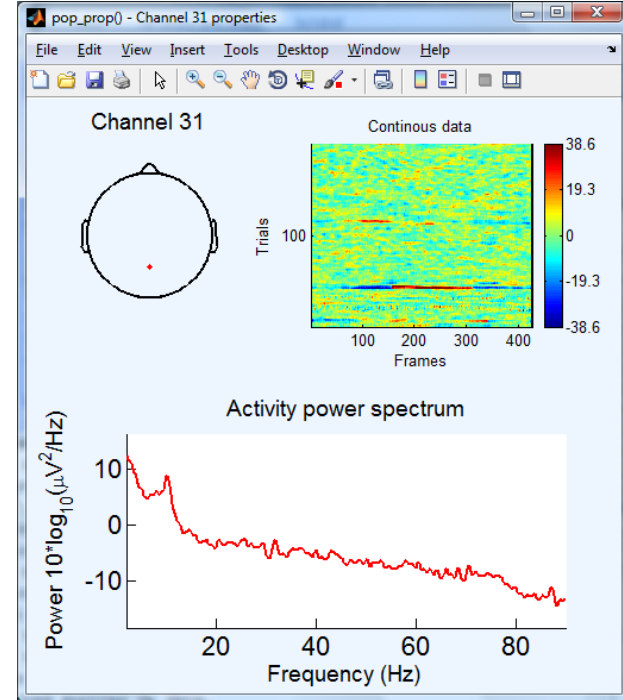
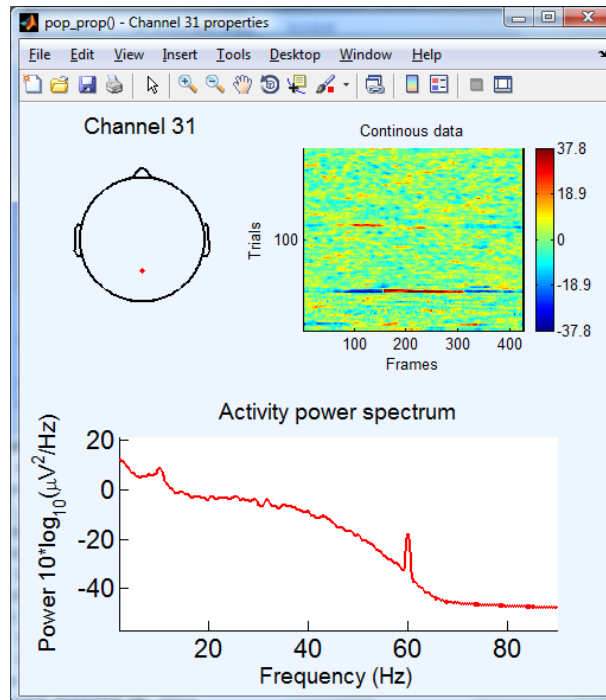
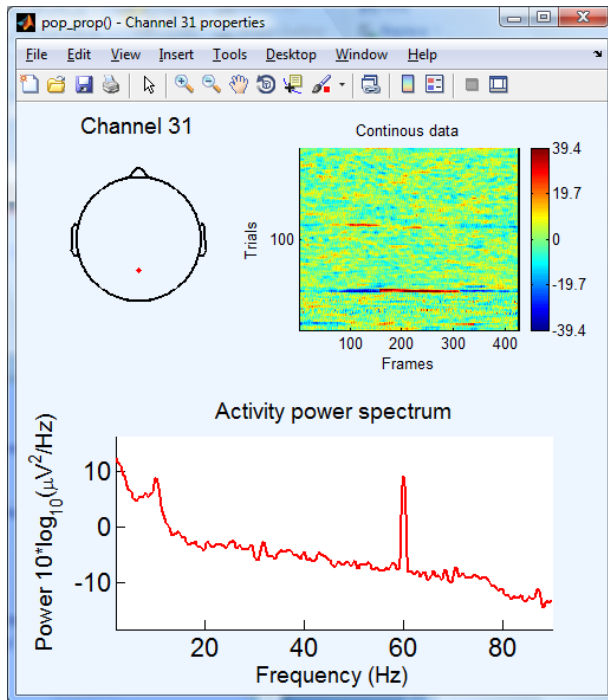
# Filter comparisons



0.5 Hz high-pass filter

0.5 Hz high-pass filter  
50 Hz low-pass filter

0.5 Hz high-pass filter  
Cleanline: 60 Hz





# Pre-processing pipeline



**Collect high-density  
EEG data (>30 chan)**

**Import into EEGLAB**

**Import event markers  
and channel locations**

**Re-reference/  
down-sample  
(if necessary)**

**High pass filter  
(~.5 – 1 Hz)**

**Remove line noise  
(if necessary)**

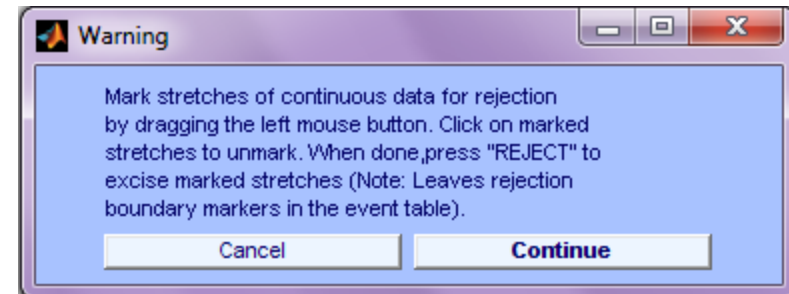
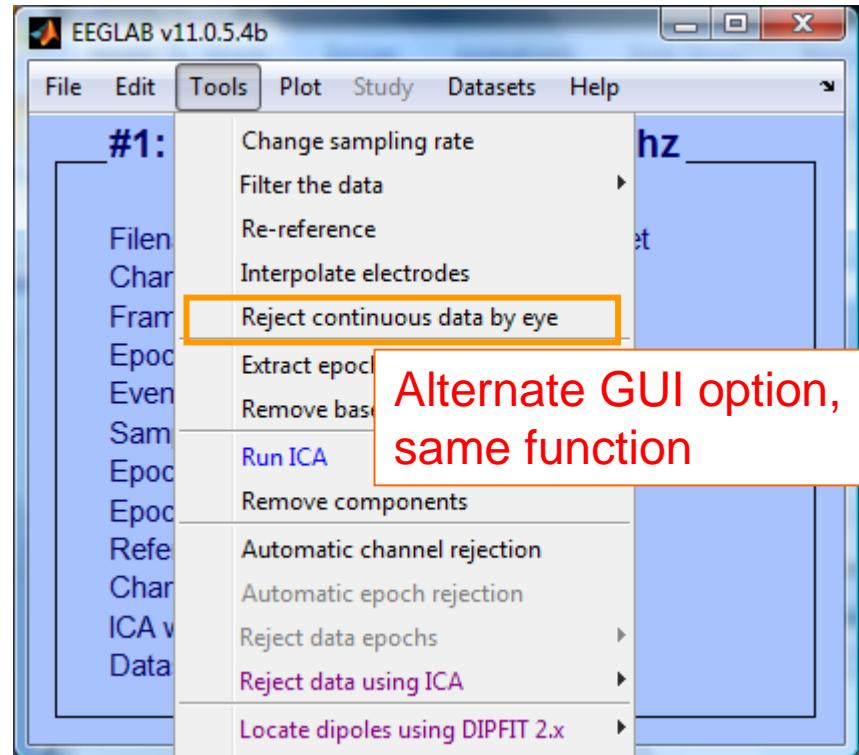
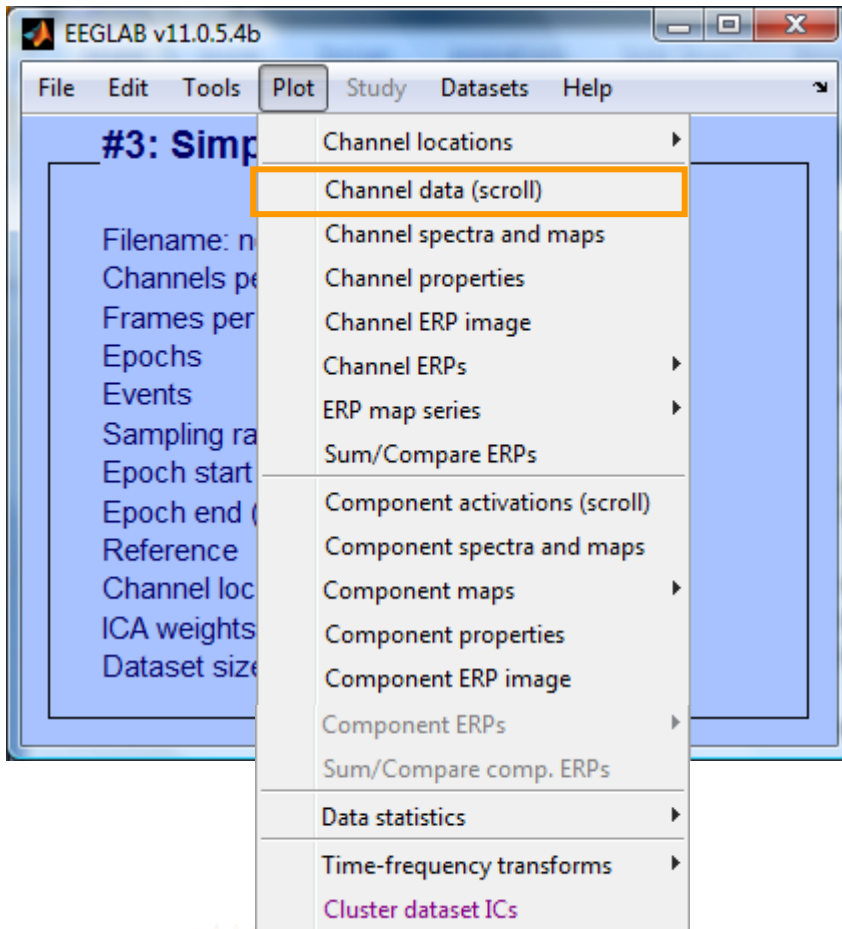
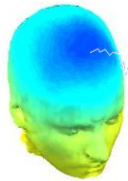
**Identify/reject  
bad channels**

**Reject large artifact  
time points**

**Run ICA**

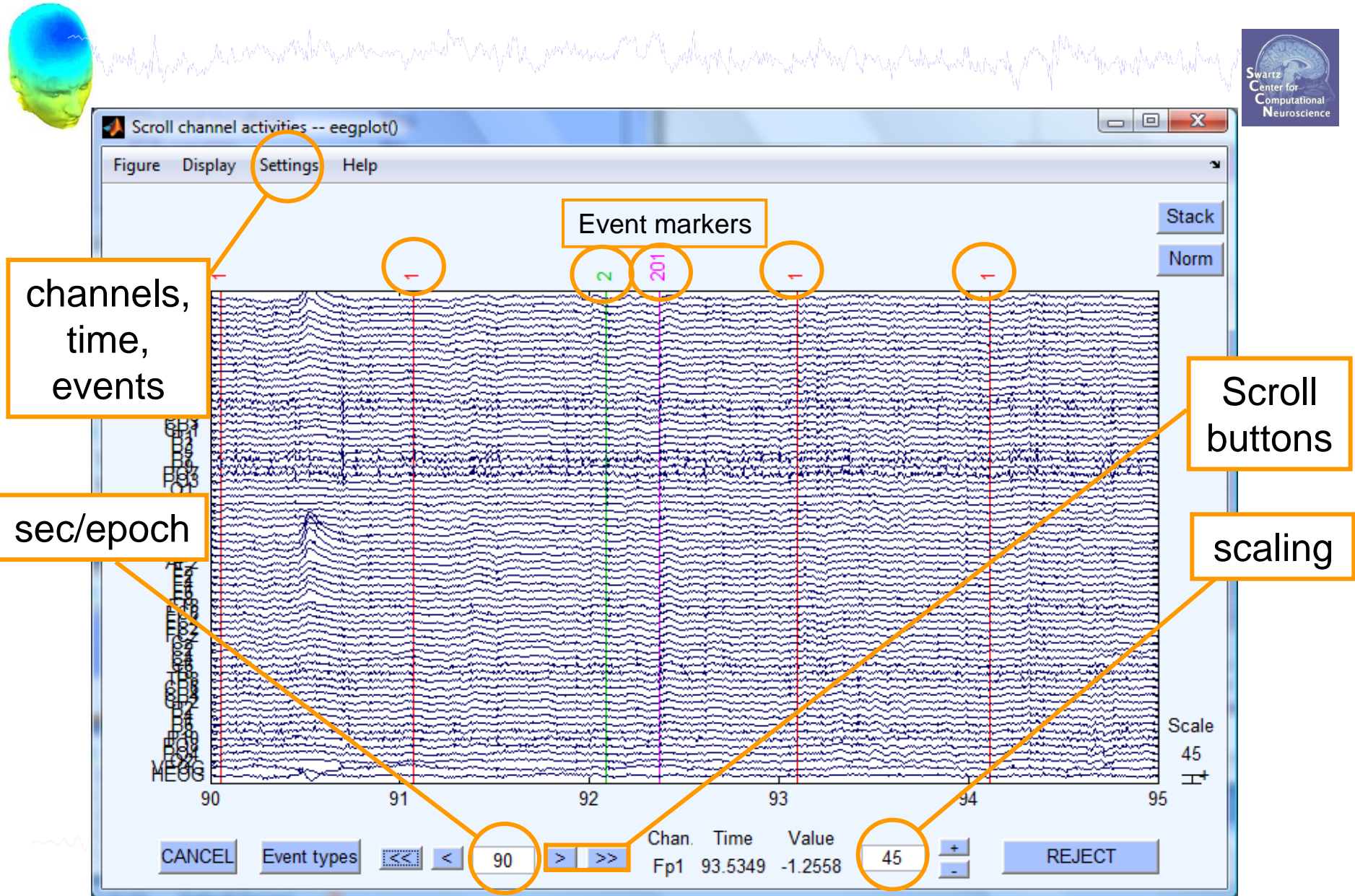


# Scroll channel data



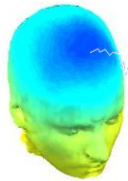


# Scroll channel data





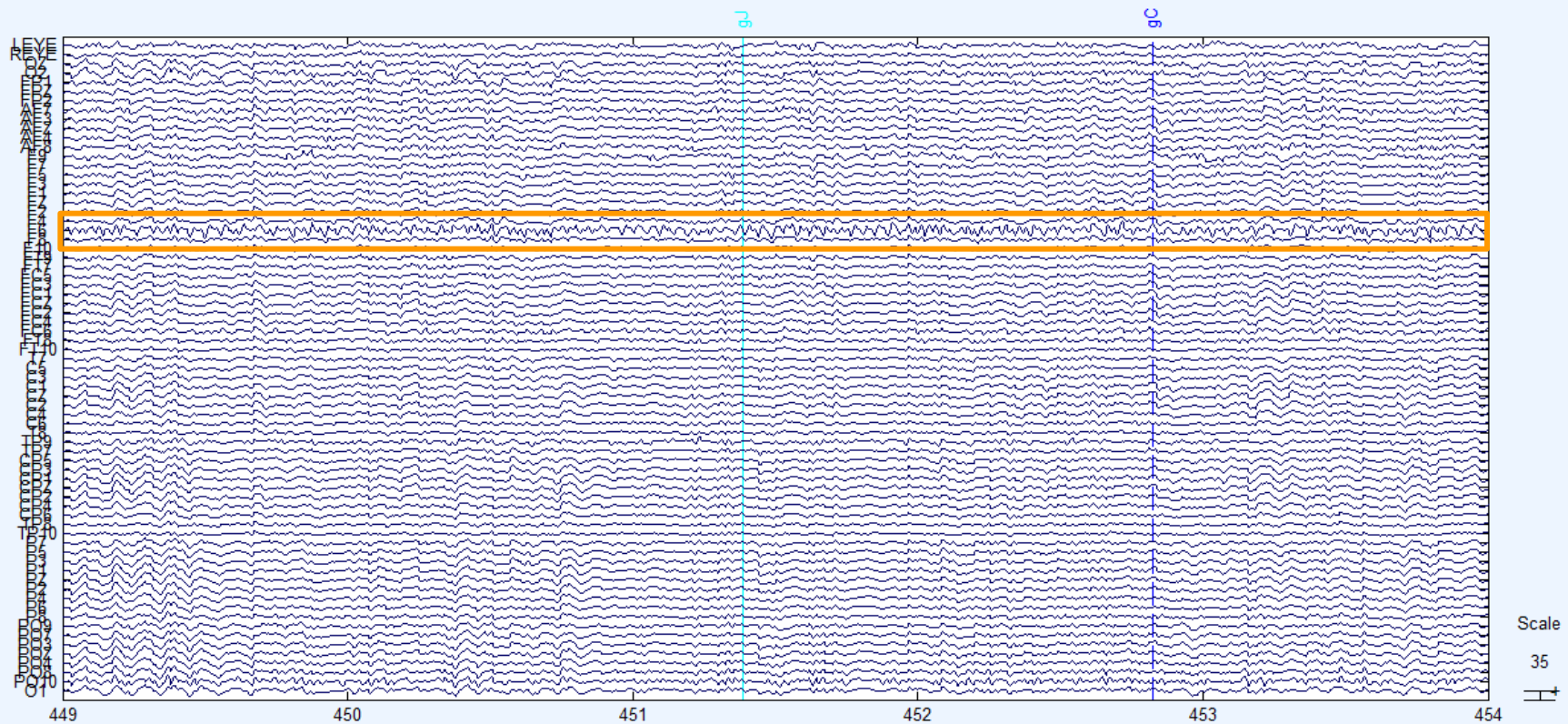
# Manually identifying bad channels



Scroll channel activities -- eegplot()

Figure Display Settings Help

Identify bad channel



Scale  
35

CANCEL

Event types

<<

<

449

>

>>

Chan.

Time

Value

O1

451.0988

3.6619

35

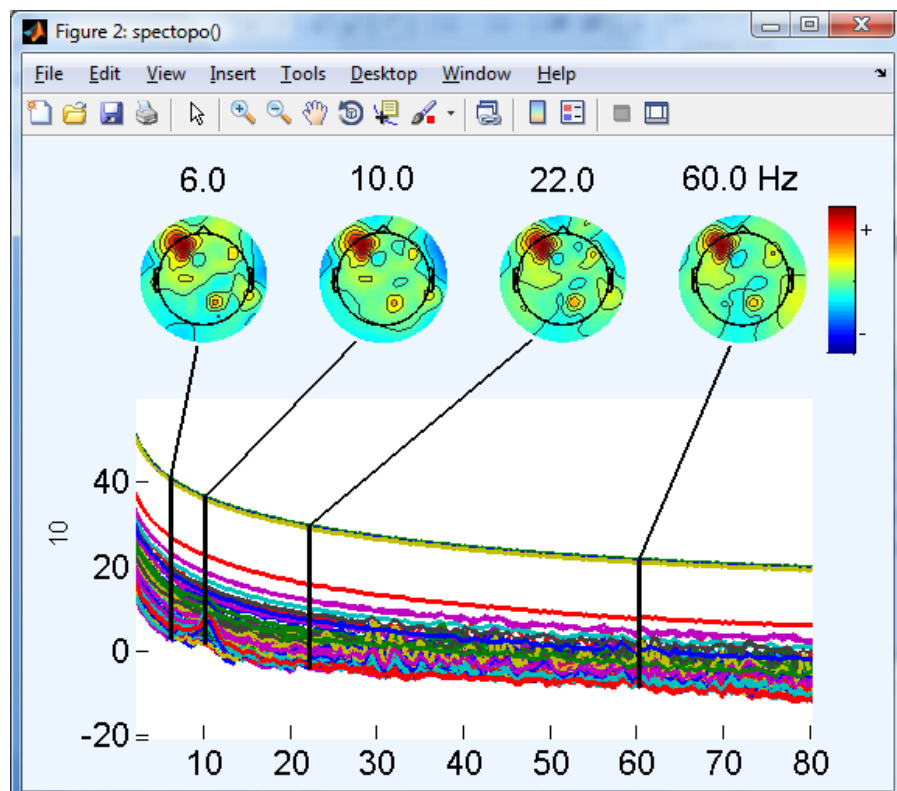
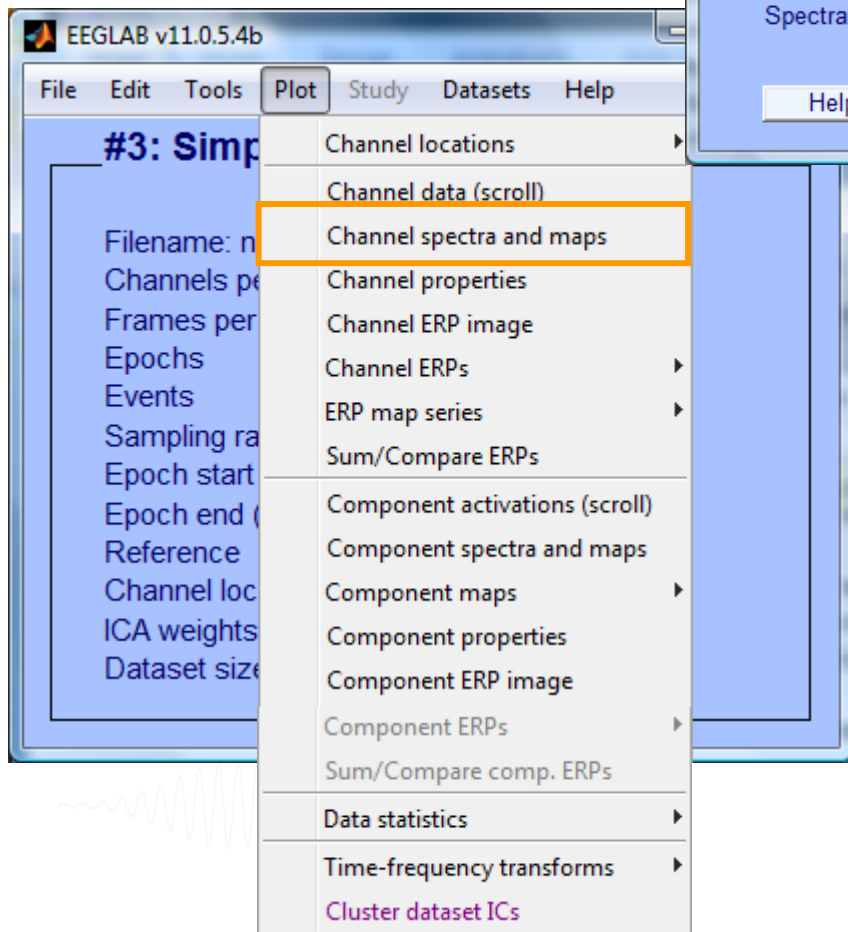
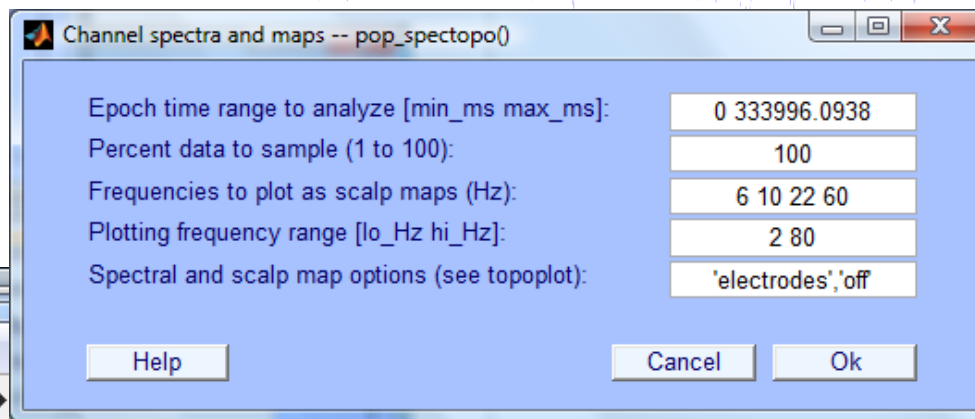
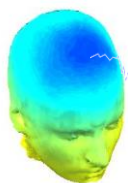
+

-

REJECT

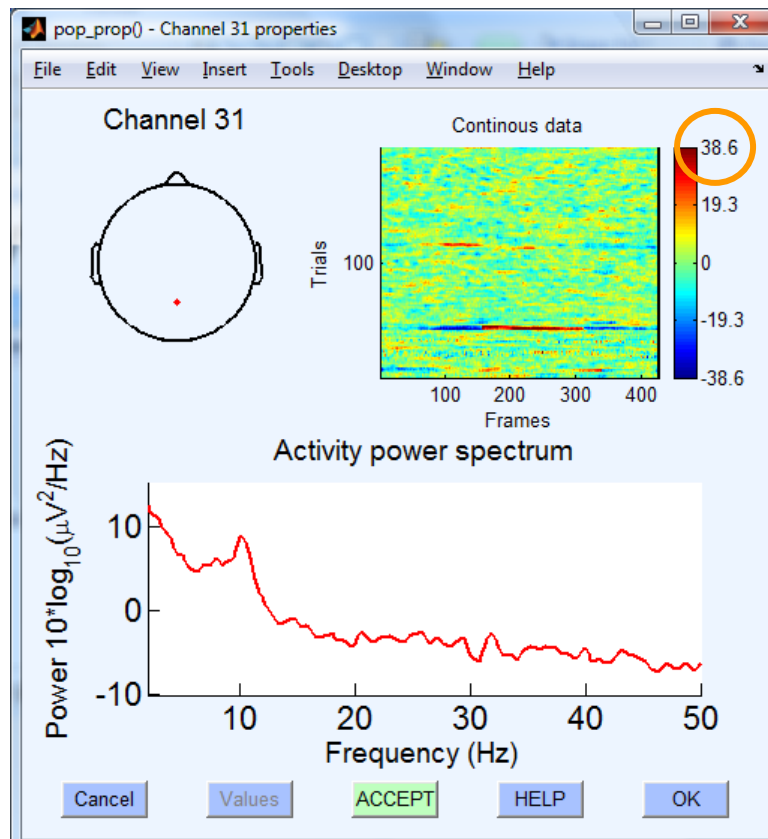
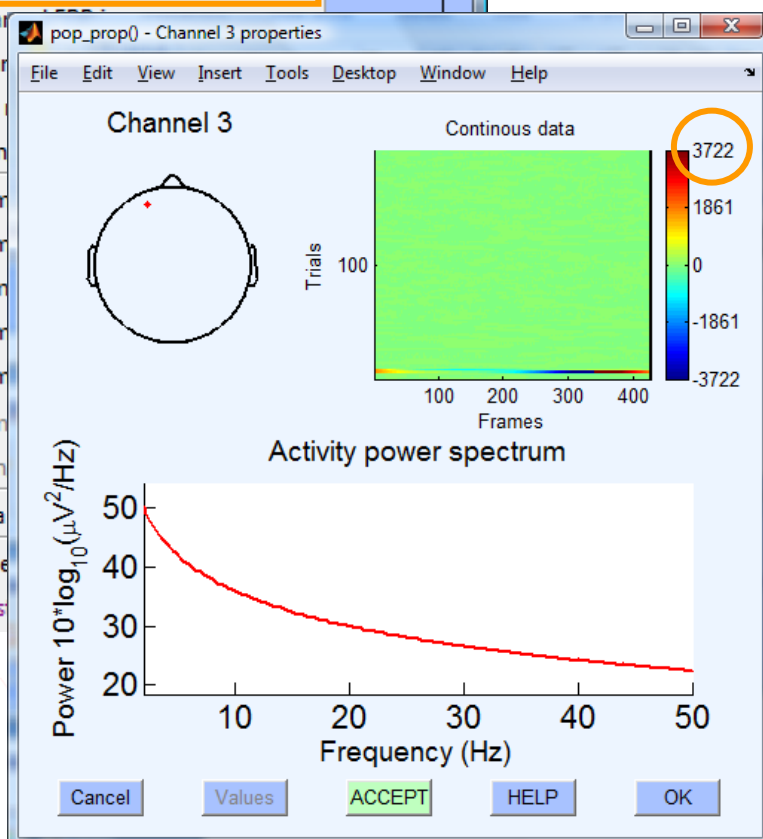
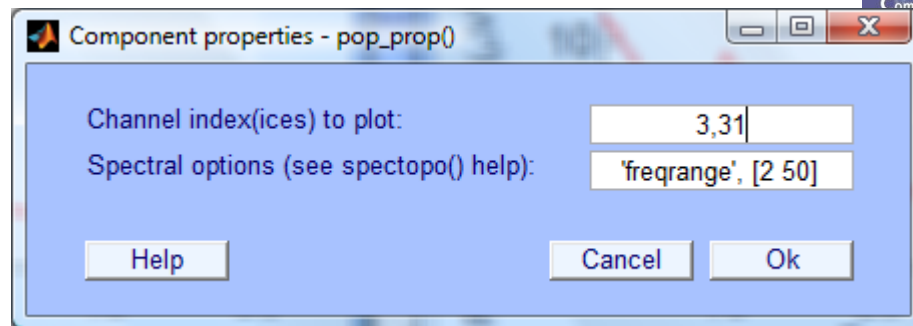
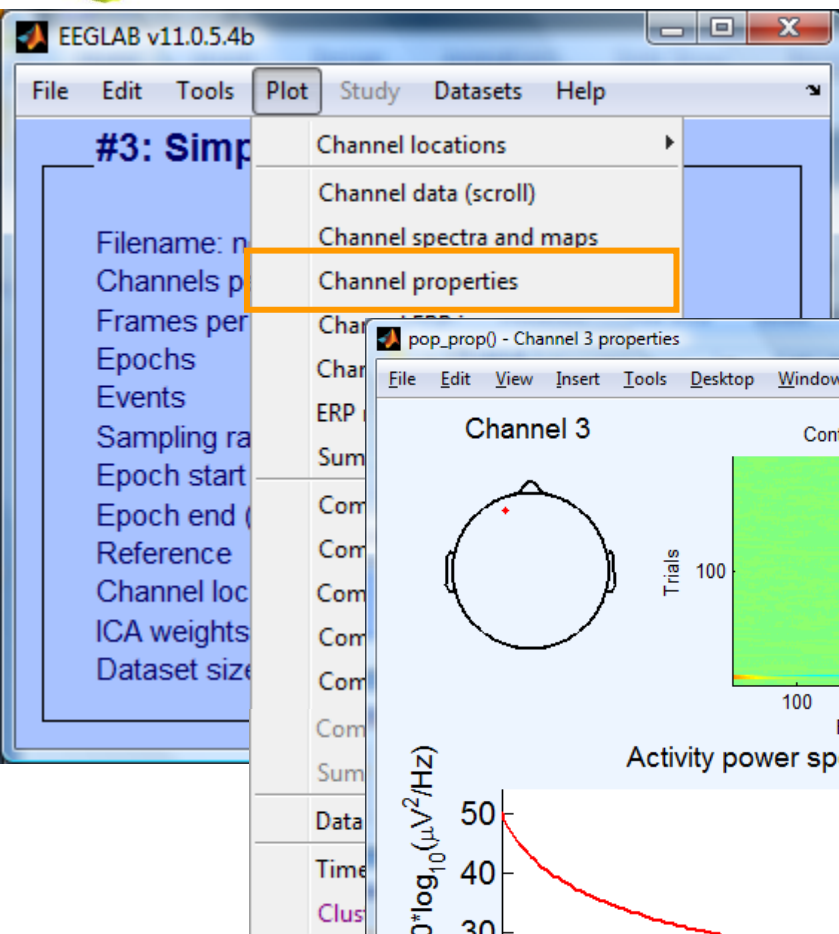
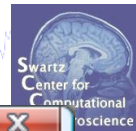
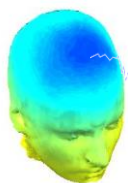


# Manually identifying bad channels



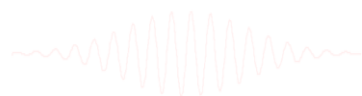
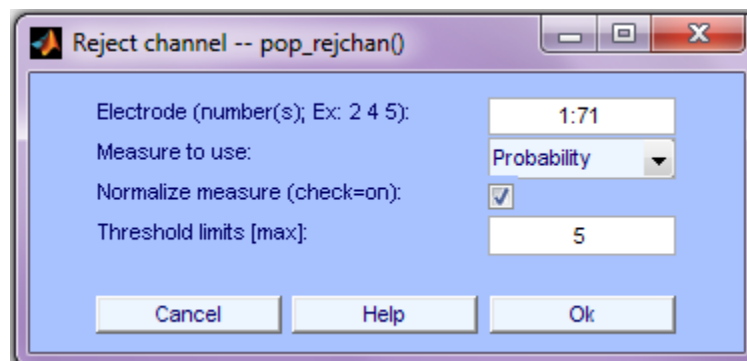
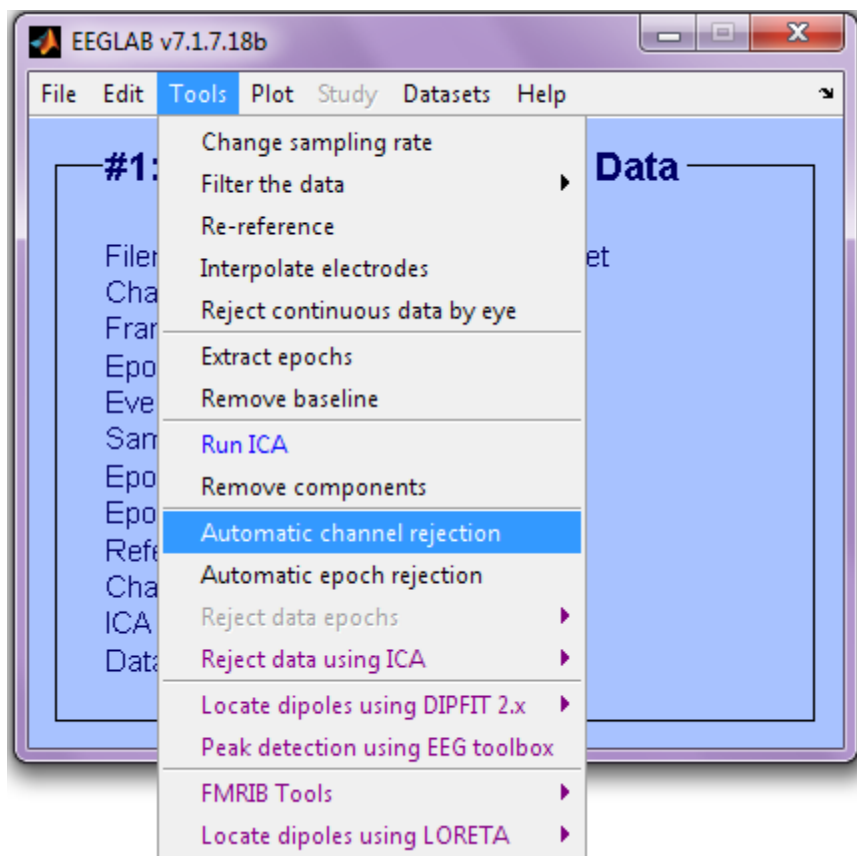


# Manually identifying bad channels





# Auto-detection of noisy channels



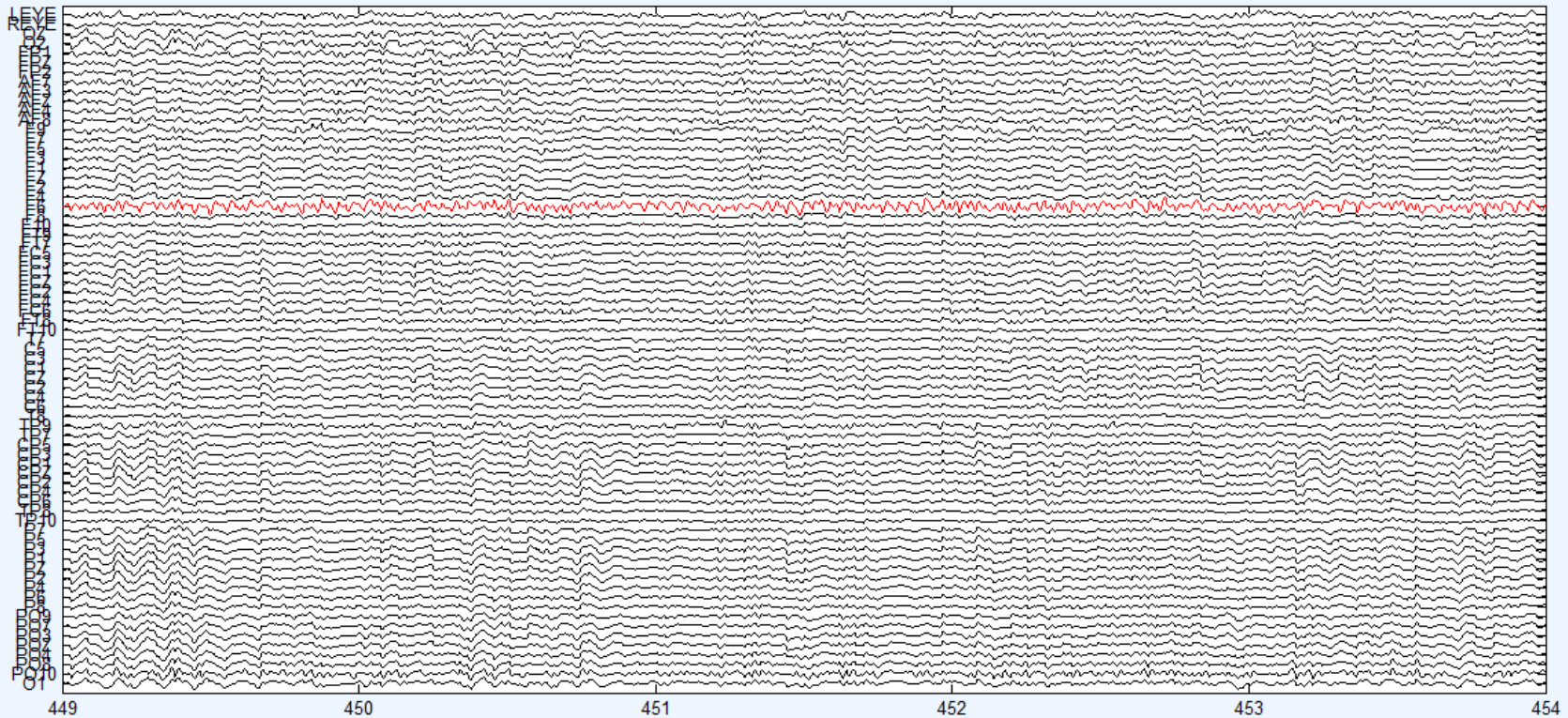


# Auto-detected noisy channel



Scroll component activities -- eegplot()

Figure Display Settings Help



Scale  
35  
↑

CANCEL

<<

<

449

>

>>

Chan.

Time

Value

TP8

452.1146

-2.6647

35

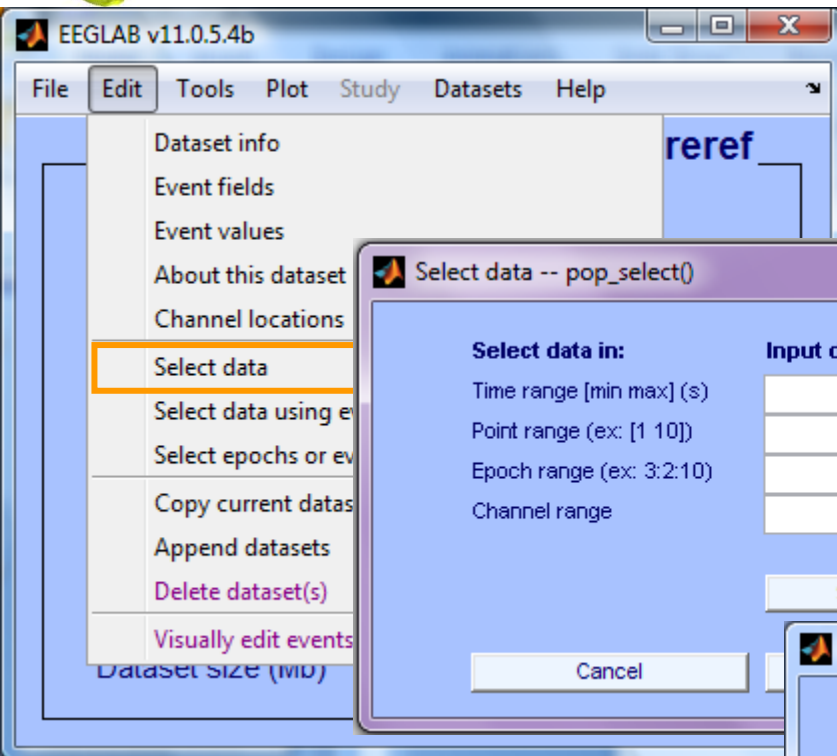
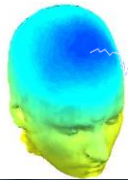
+

-

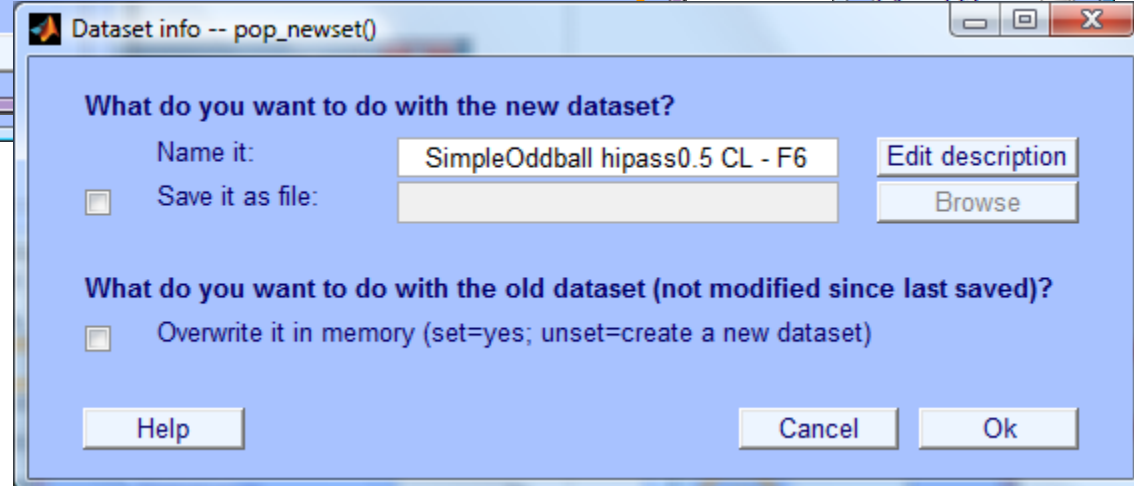
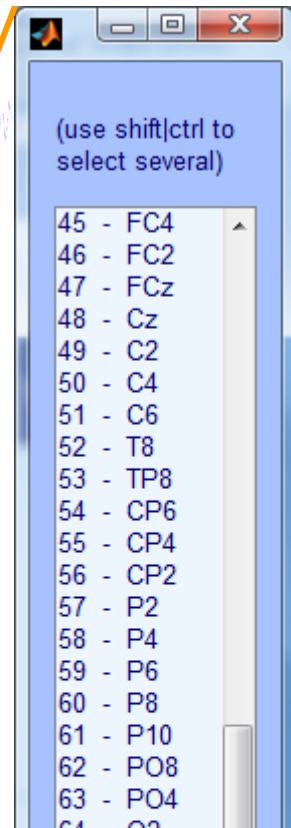
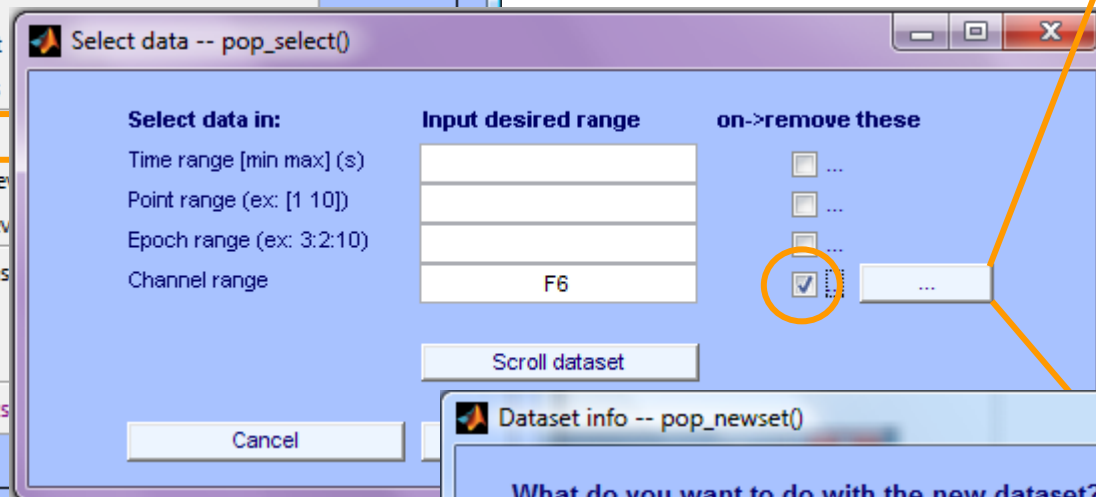
REJECT



# Removing channel(s)

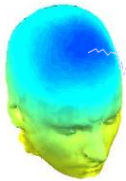


If not checked, will result  
in dataset with one channel





# Pre-processing pipeline



Collect high-density  
EEG data (>30 chan)

Import into EEGLAB

Import event markers  
and channel locations

Re-reference/  
down-sample  
(if necessary)

High pass filter  
( $\sim 5 - 1$  Hz)

Remove line noise  
(if necessary)

**STOP!**

Save your dataset  
here; you will import  
ICA weights to this  
dataset later

Identify/reject  
bad channels

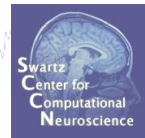
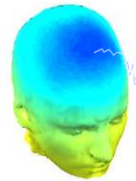
Reject large artifact  
time points

Plot data

Run ICA

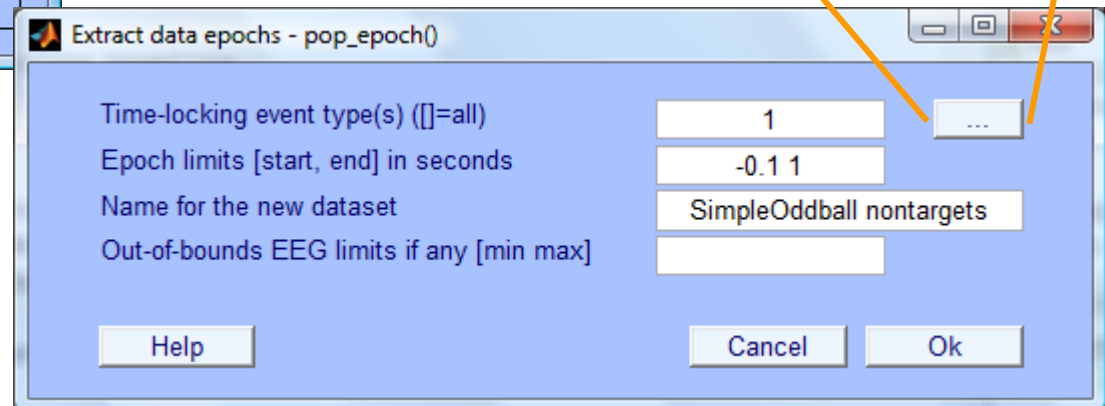
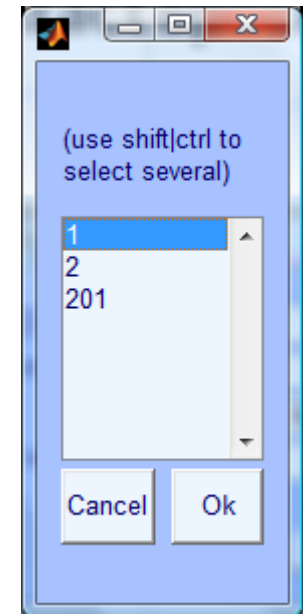


# Extract epochs



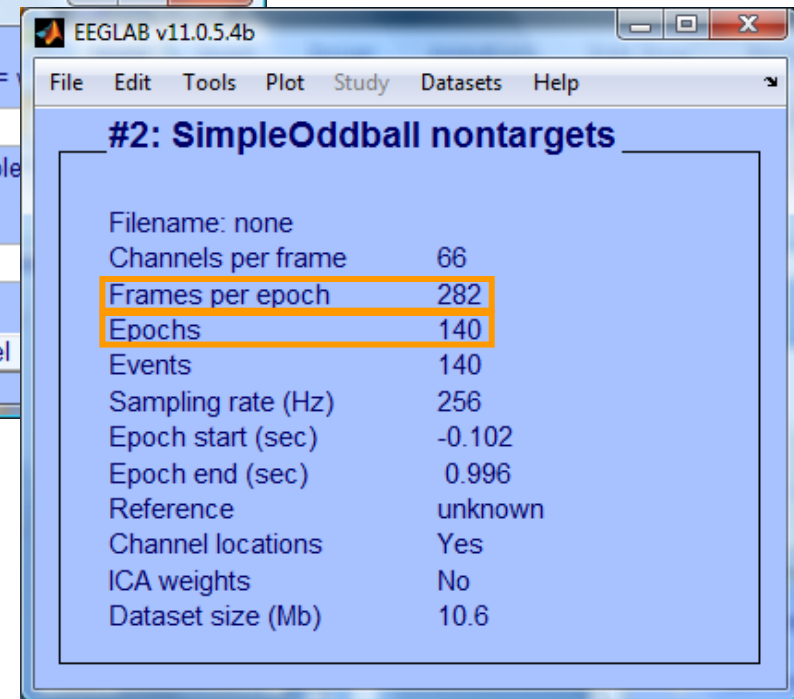
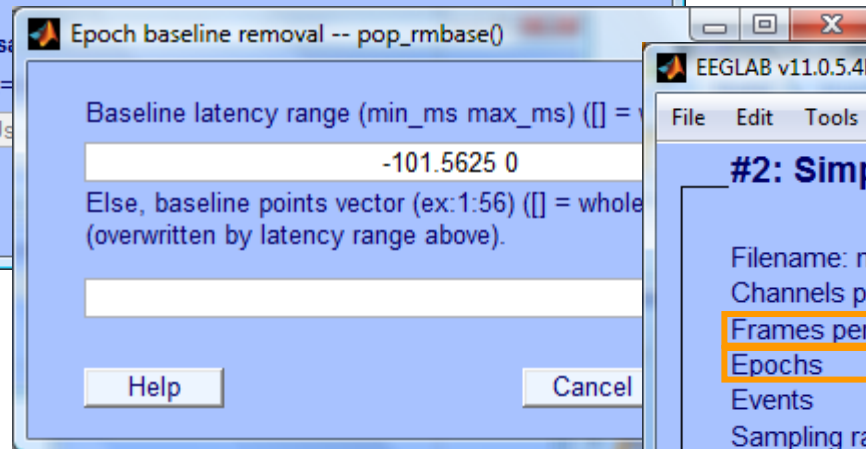
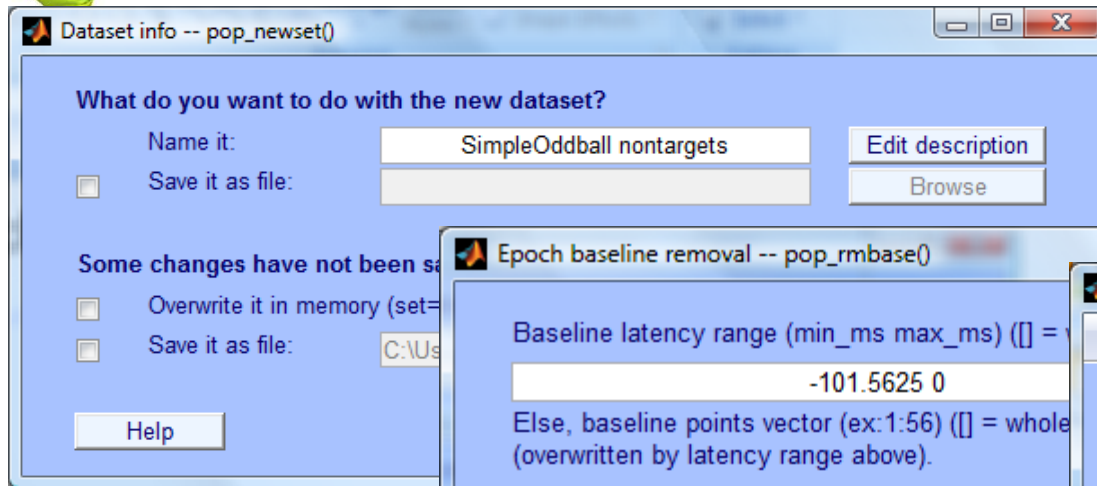
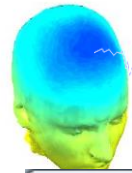
```
>> eeg_eventtypes (EEG)
```

1	140	star
2	60	circle
201	60	button press

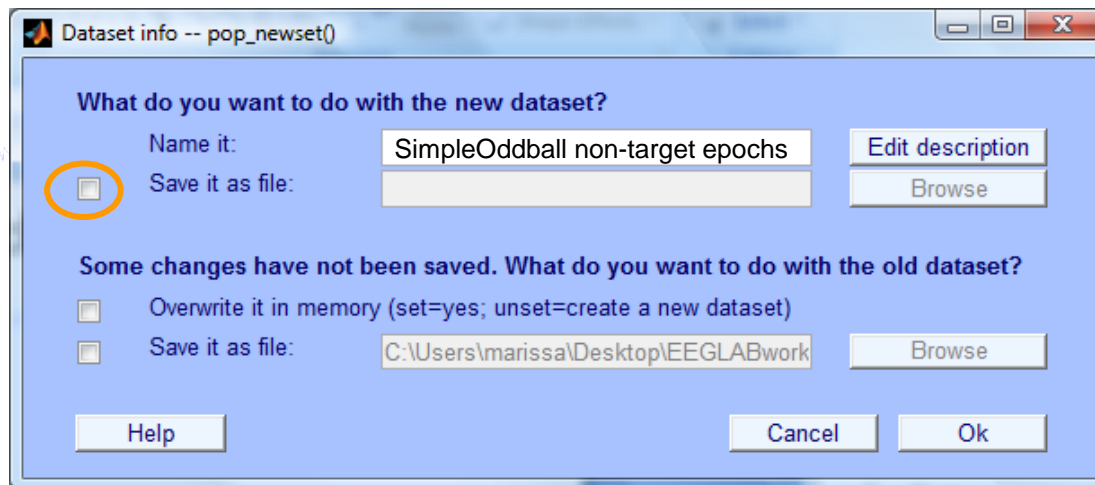
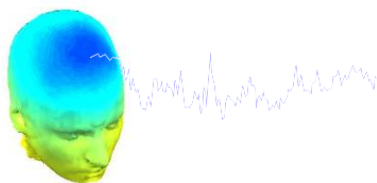




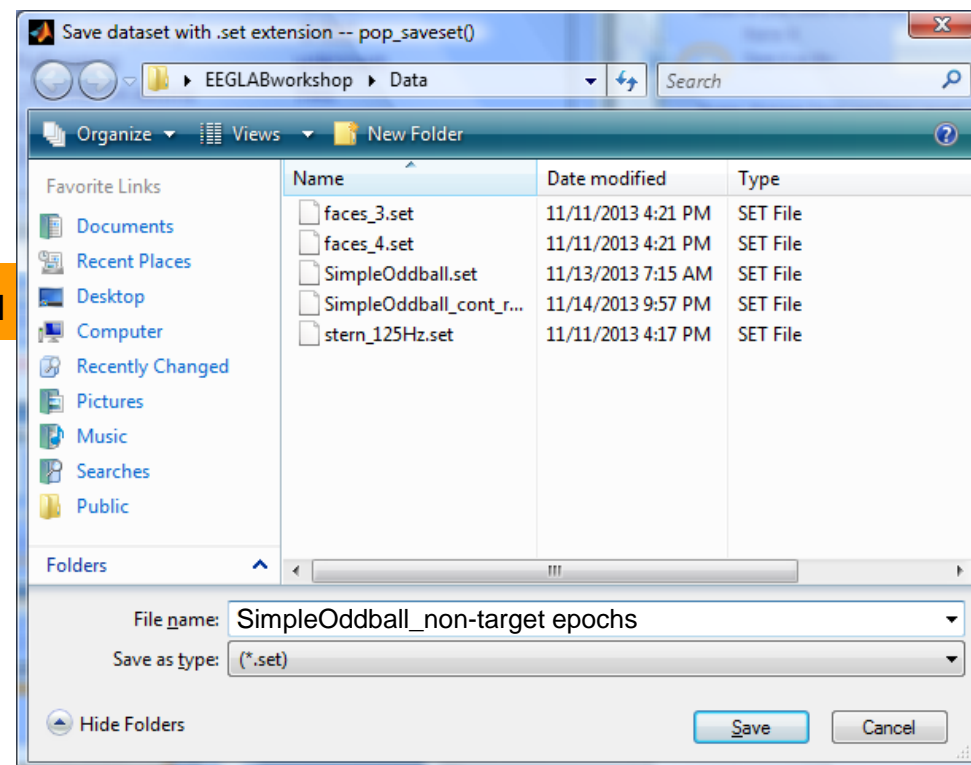
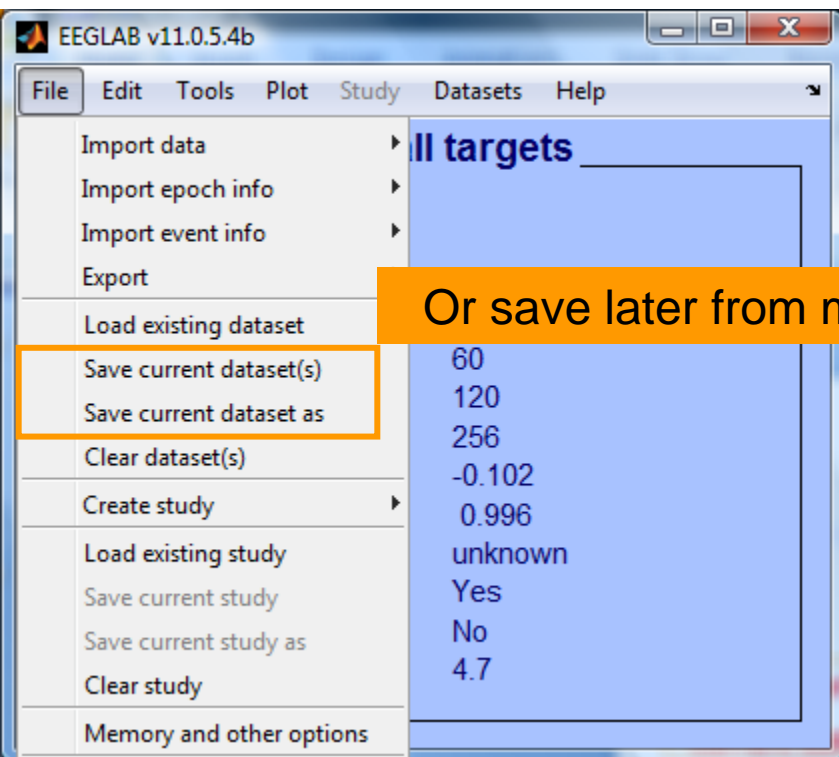
# Extract epochs





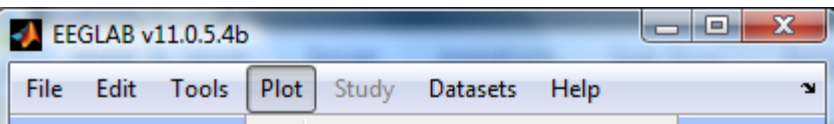
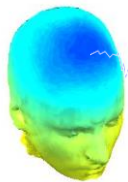


## Save dataset (optional)





# Scroll (epoched) channel data



## #3: Simple

Filename: n  
Channels per  
Frames per  
Epochs  
Events  
Sampling rate  
Epoch start  
Epoch end (s)  
Reference  
Channel location  
ICA weights  
Dataset size

Channel location

Channel data

Channel spectra

Channel properties

Channel ERP

Channel ERP

ERP map series

Sum/Component

Component

Component

Component

Component

Component

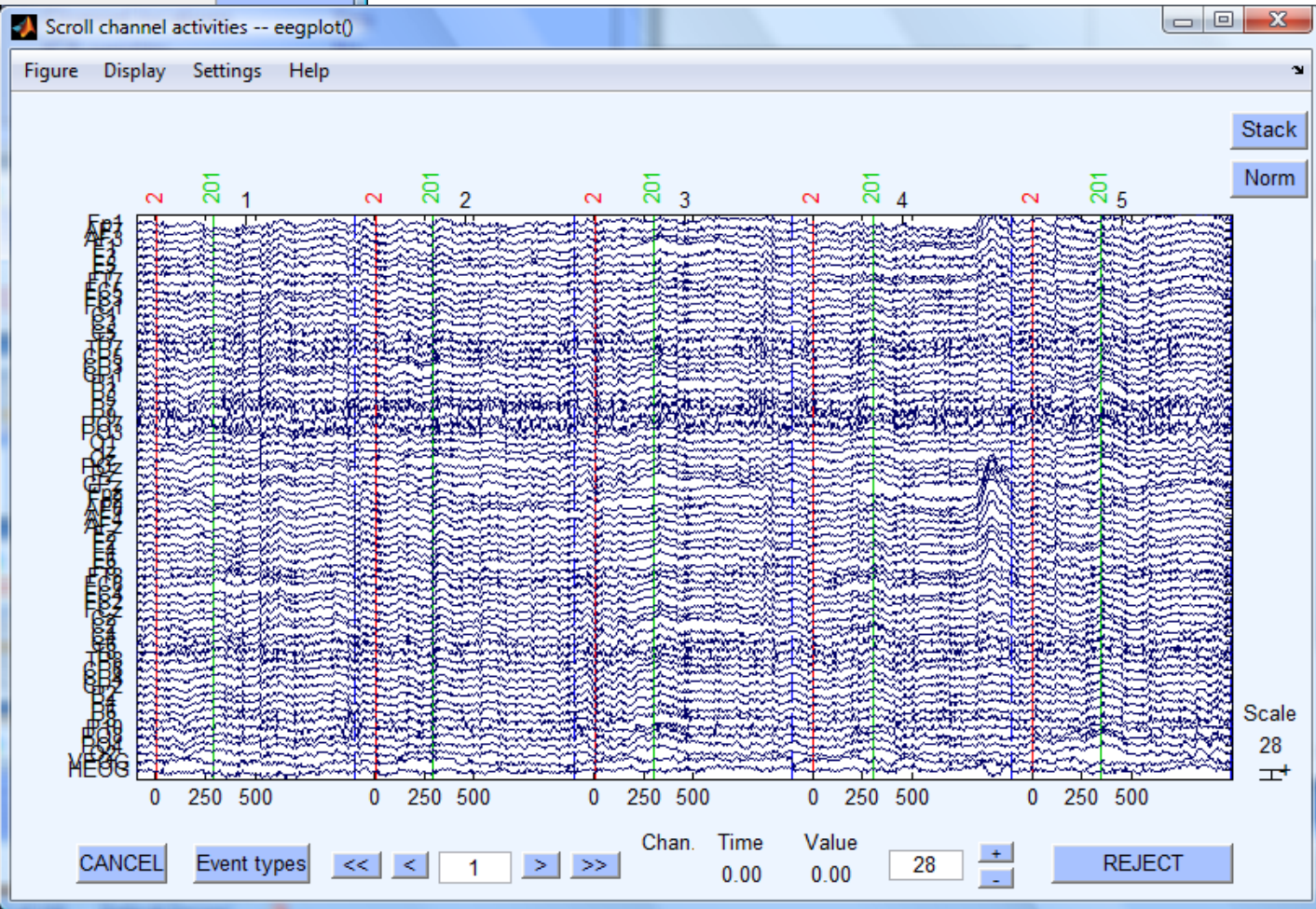
Component

Sum/Component

Data statistics

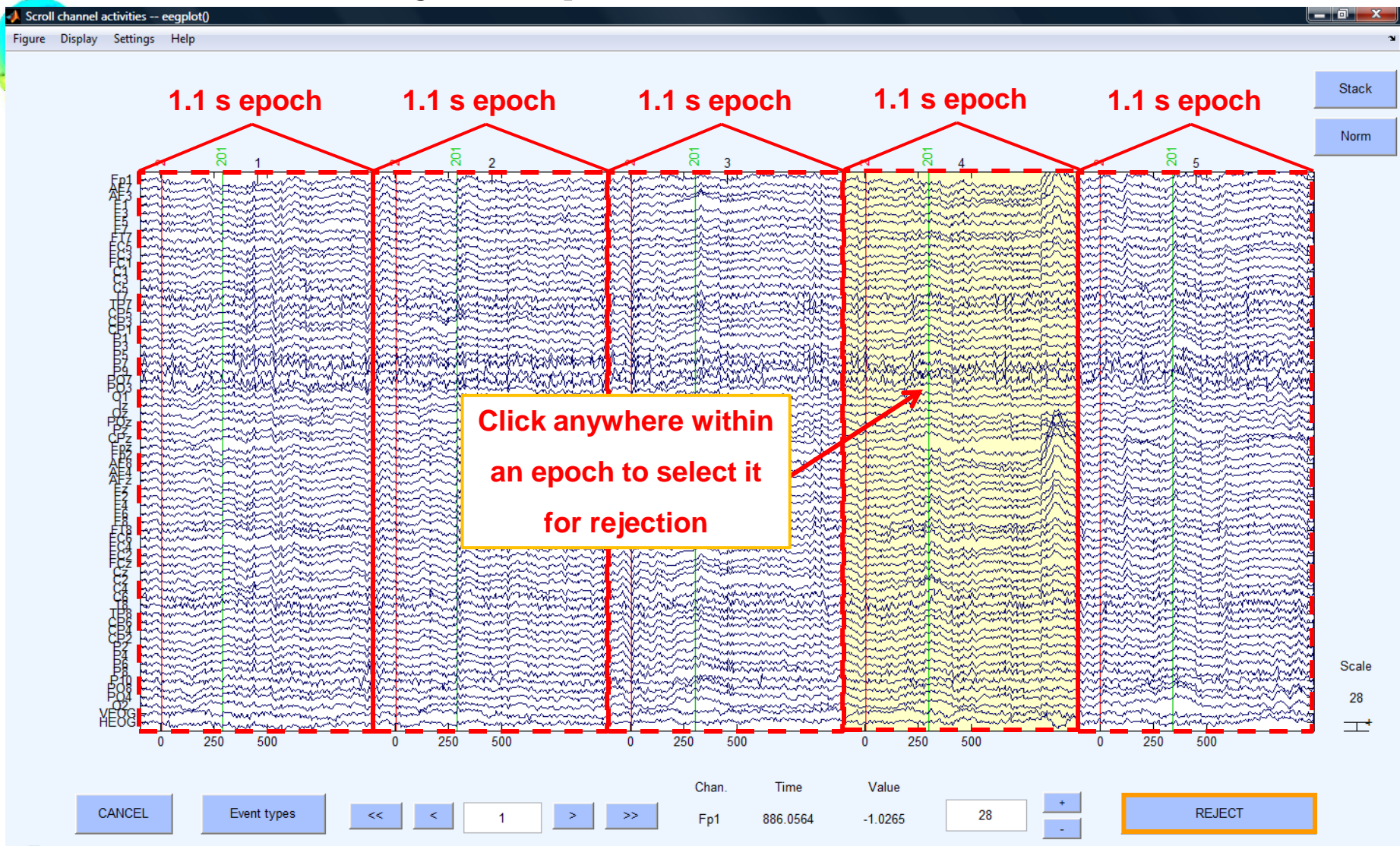
Time-frequency

Cluster data



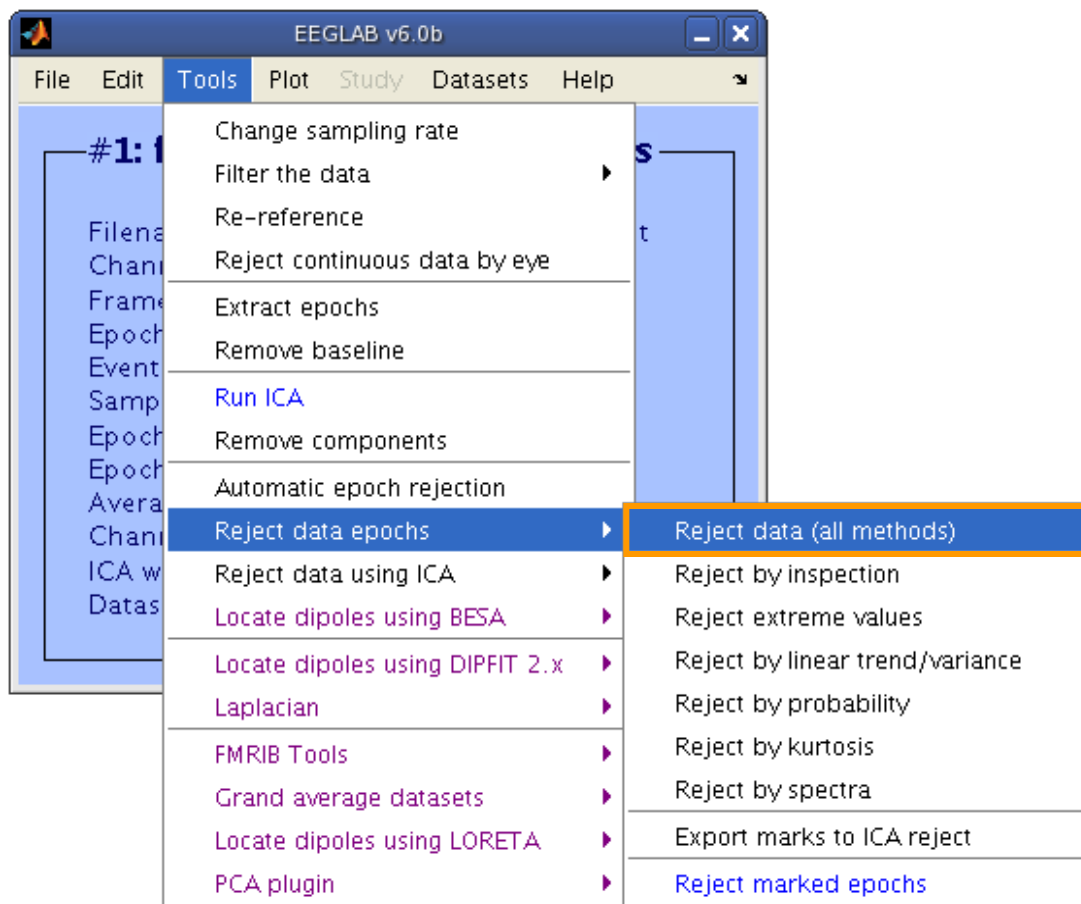


# Reject epochs with artifact



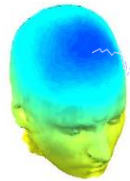


# Reject data epochs





# Reject data epochs



visual  
inspection

Reject trials using data statistics - pop\_rejmenu()

**Mark trials by appearance** ☐  Marked trials 0

**Find abnormal values** ☐

Upper limit(s) (uV)  Lower limit(s) (uV)   
Start time(s) (ms)  Ending time(s) (ms)   
Electrode(s)  Currently marked trials

**Find abnormal trends** ☐

Max slope (uV/epoch)  R-squared limit (0 to 1)   
Electrode(s)  Currently marked trials

**Find improbable data** ☐

Single-channel limit (std. dev.)  All channels limit (std. dev.)   
Electrode(s)  Currently marked trials

**Find abnormal distributions** ☐

Single-channel limit (std. dev.)  All channels limit (std. dev.)   
Electrode(s)  Currently marked trials

**Find abnormal spectra (slow)** ☐

Upper limit(s) (dB)  Lower limit(s) (dB)   
Low frequency(s) (Hz)  High frequency(s) (Hz)   
Electrode(s)  Currently marked trials

**Plotting options**

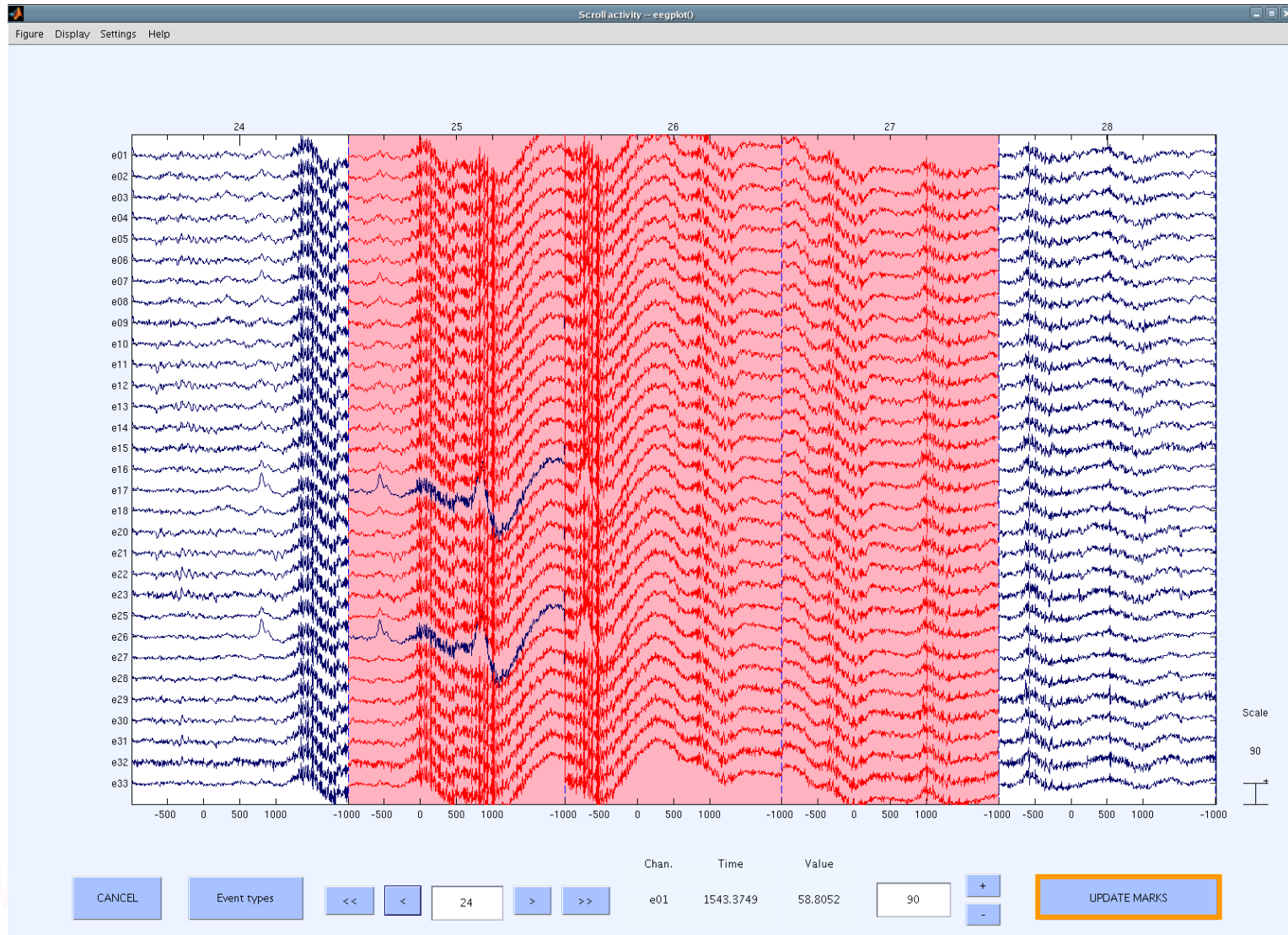
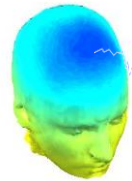
Show all trials marked for rejection by the measure selected above or checked below | /

☒ Abnormal appearance ☒ Abnormal values ☒ Abnormal trends  
☒ Improbable epochs ☒ Abnormal distributions ☒ Abnormal spectra

probability

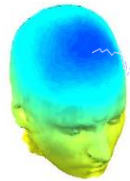


# Reject data epochs





# Reject data epochs



Reject trials using data statistics - pop\_rejmenu()

Mark trials by appearance ☐ Scroll Data Marked trials 0

**Find abnormal values** ☐

Upper limit(s) (uV)  Lower limit(s) (uV)   
 Start time(s) (ms)  Ending time(s) (ms)   
 Electrode(s)  Currently marked trials

**Find abnormal trends** ☐

Max slope (uV/epoch)  R-squared limit (0 to 1)   
 Electrode(s)  Currently marked trials

**Find improbable data** ☐

Single-channel limit (std. dev.)  All channels limit (std. dev.)   
 Electrode(s)  Currently marked trials

**Find abnormal distributions** ☐

Single-channel limit (std. dev.)  All channels limit (std. dev.)   
 Electrode(s)  Currently marked trials

**Find abnormal spectra (slow)** ☐

Upper limit(s) (dB)  Lower limit(s) (dB)   
 Low frequency(s) (Hz)  High frequency(s) (Hz)   
 Electrode(s)  Currently marked trials

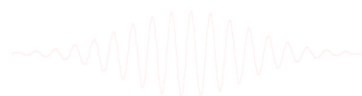
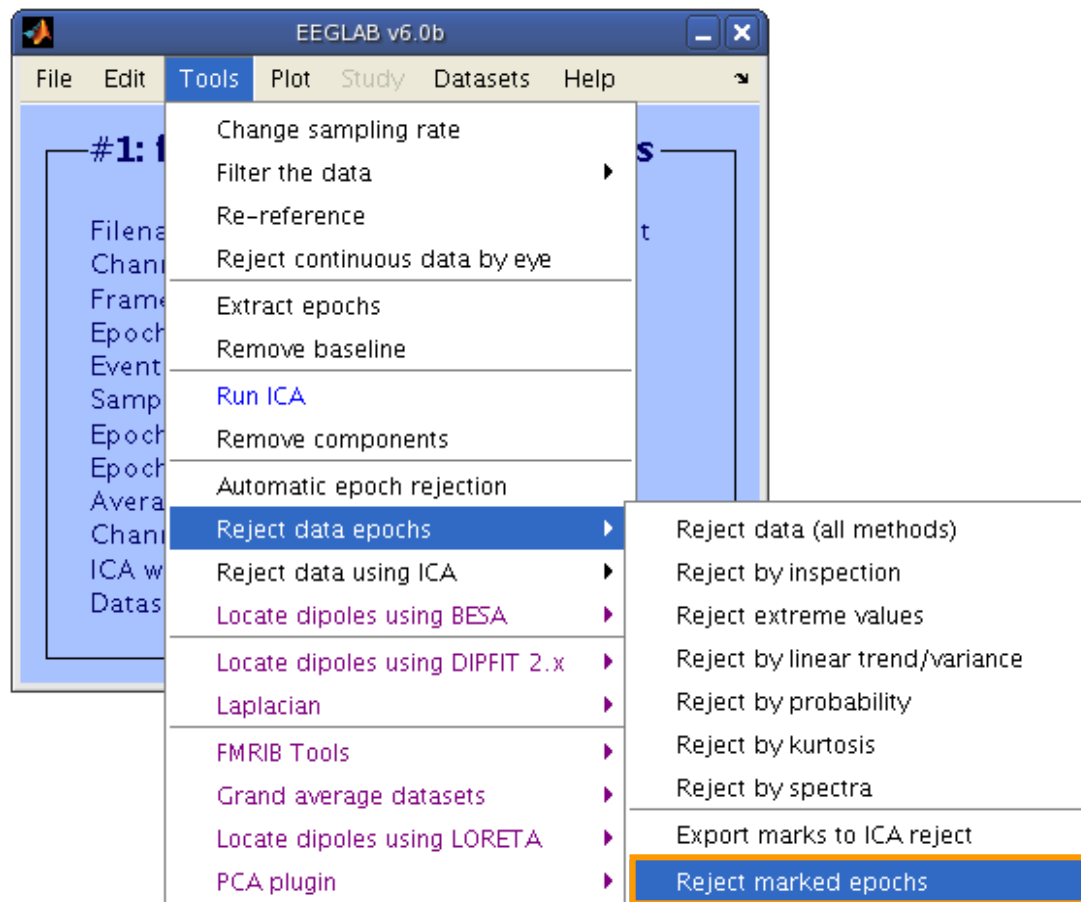
**Plotting options**

Show all trials marked for rejection by the measure selected above or checked below | /

☒ Abnormal appearance ☒ Abnormal values ☒ Abnormal trends  
☒ Improbable epochs ☒ Abnormal distributions ☒ Abnormal spectra

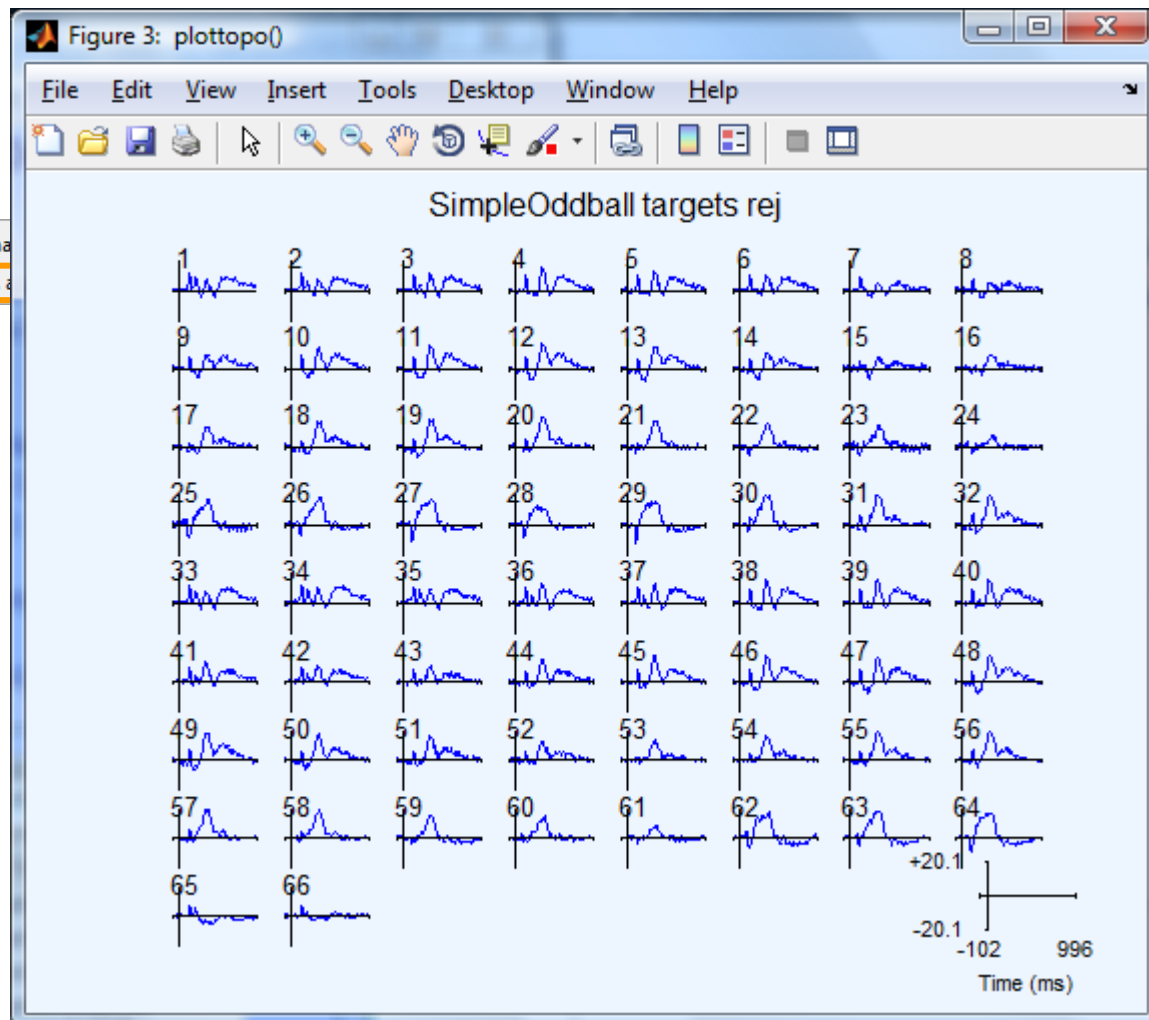
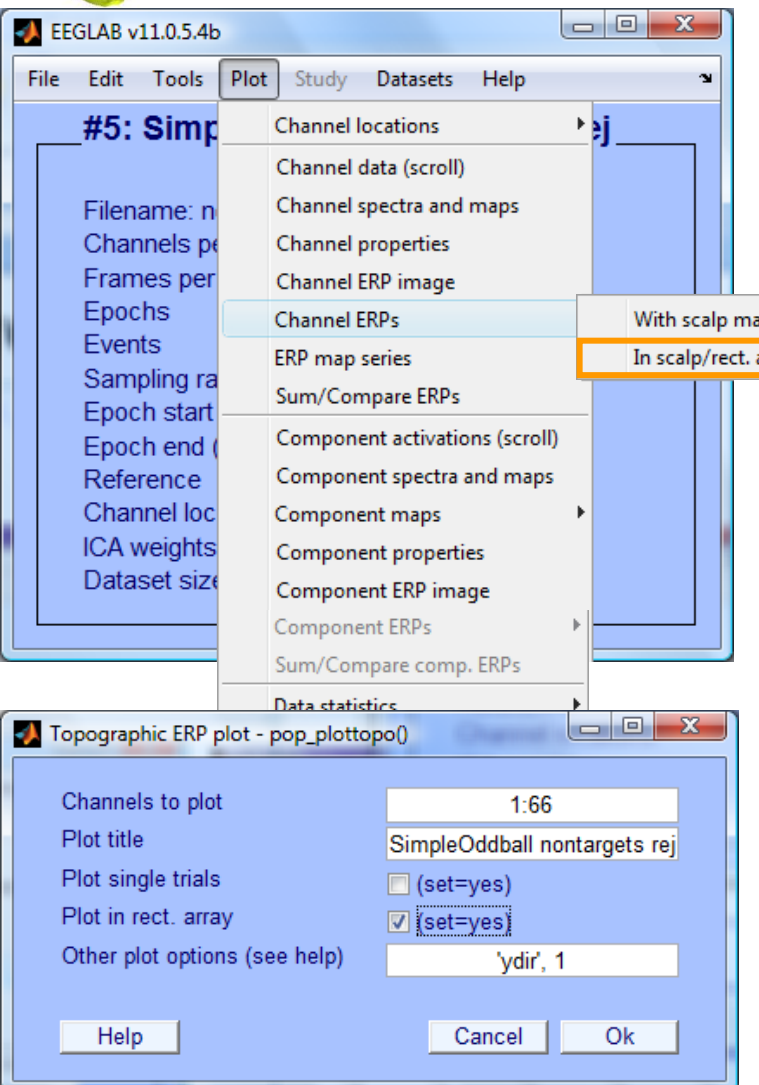
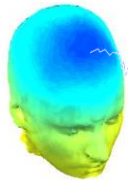


# Reject data epochs



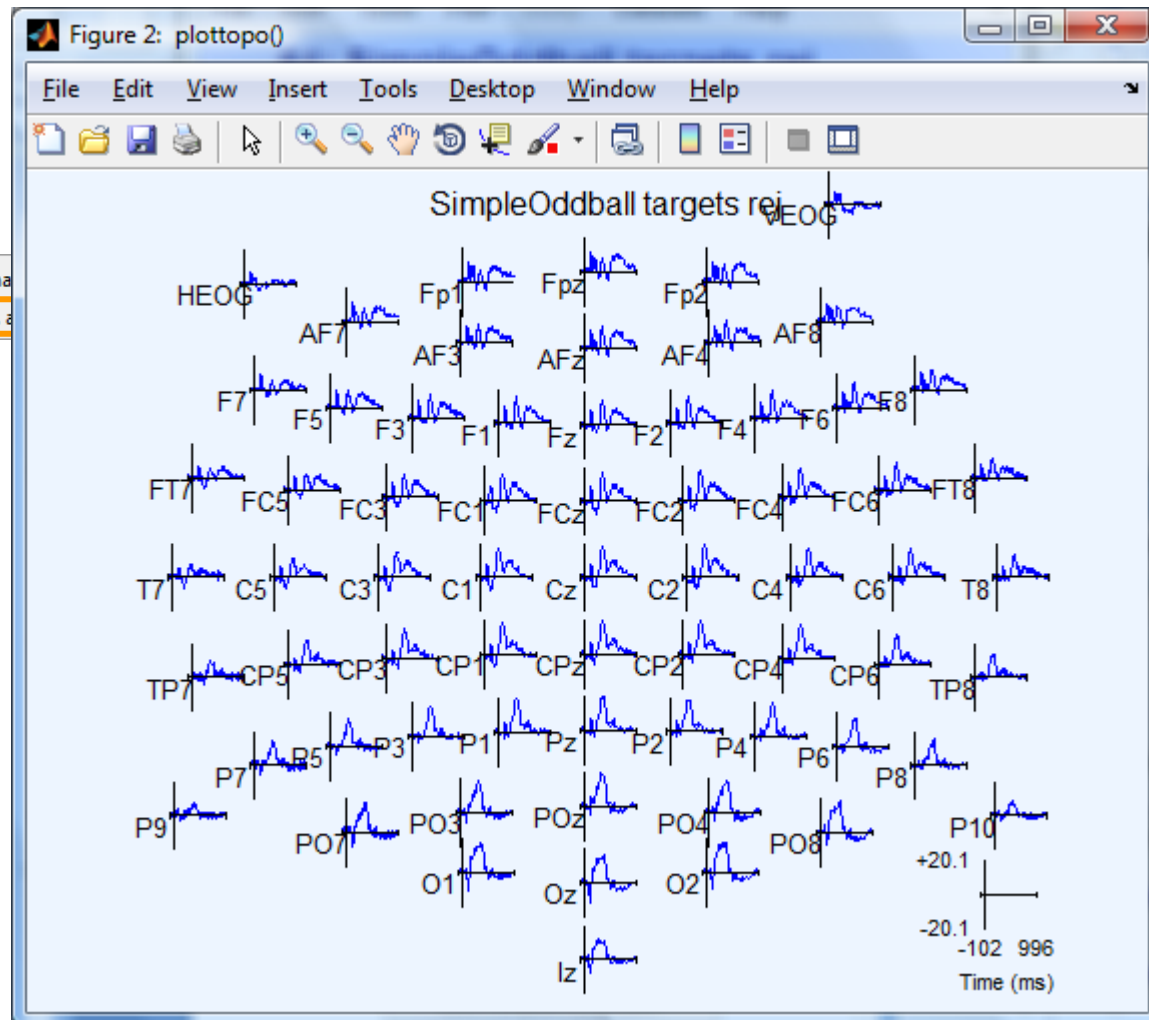
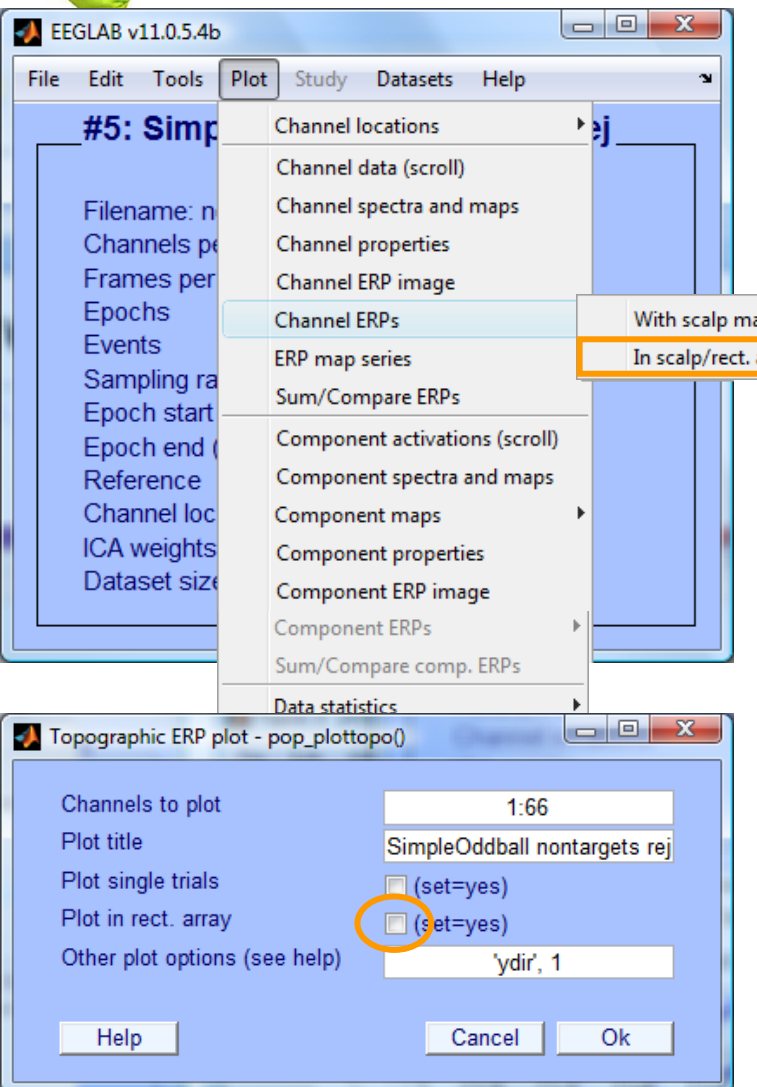
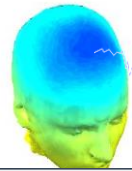


# Visualize ERP in rectangular array



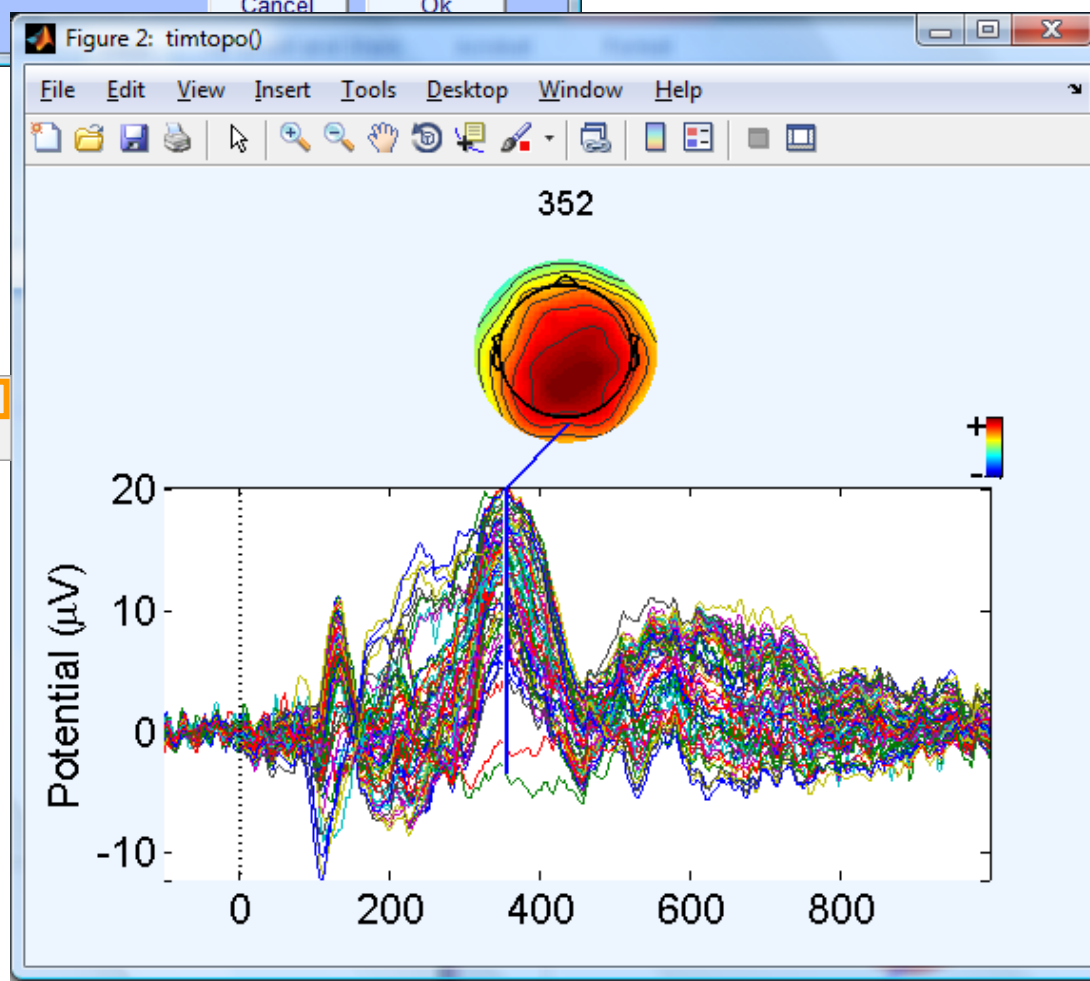
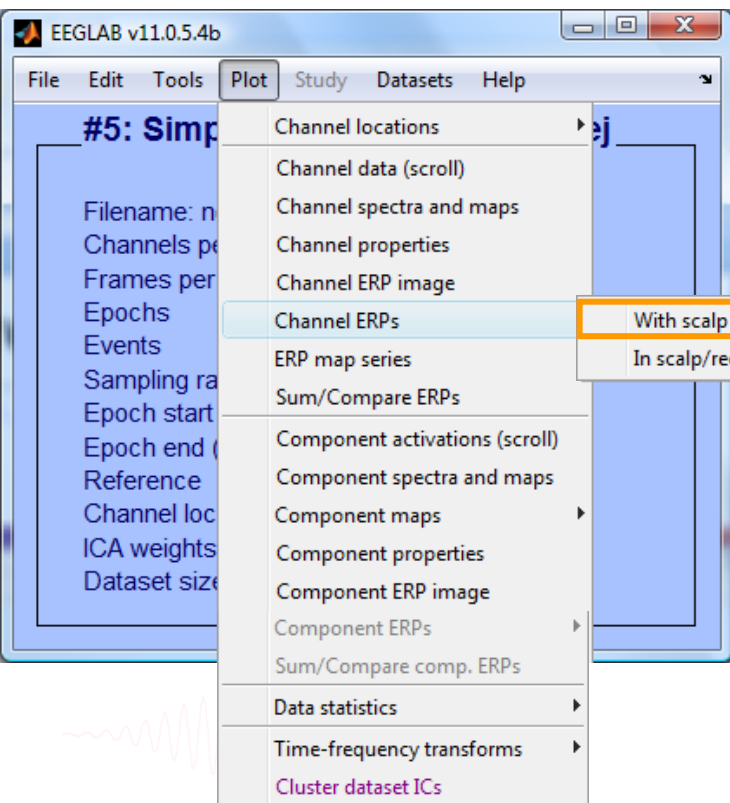
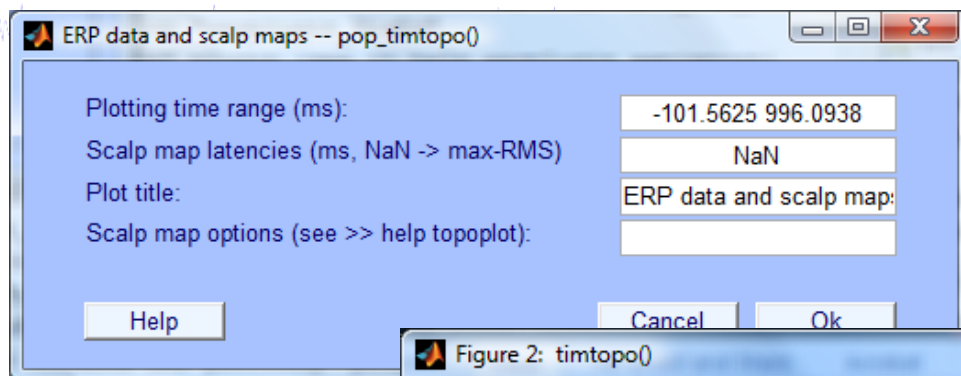


# Visualize ERP in topographic array



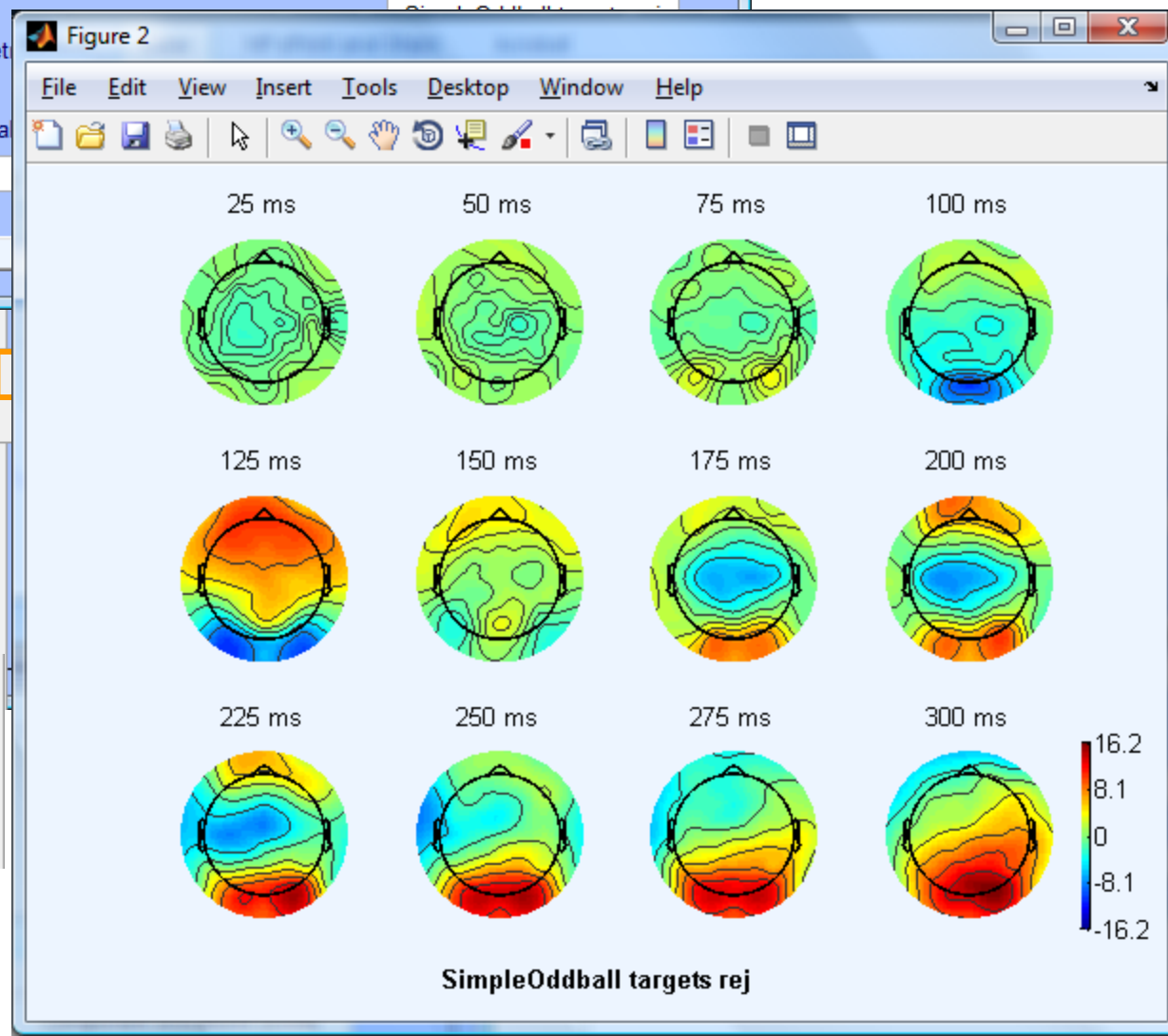
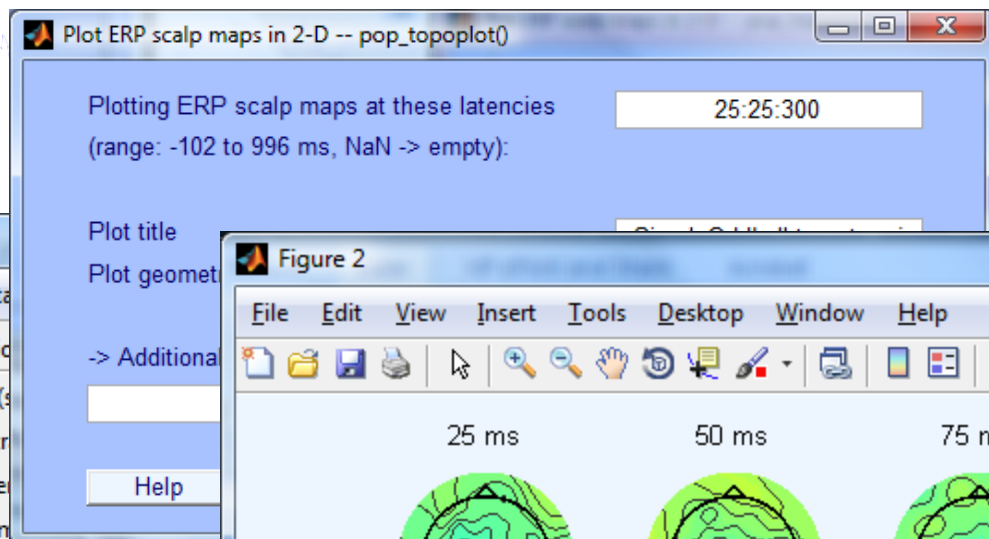
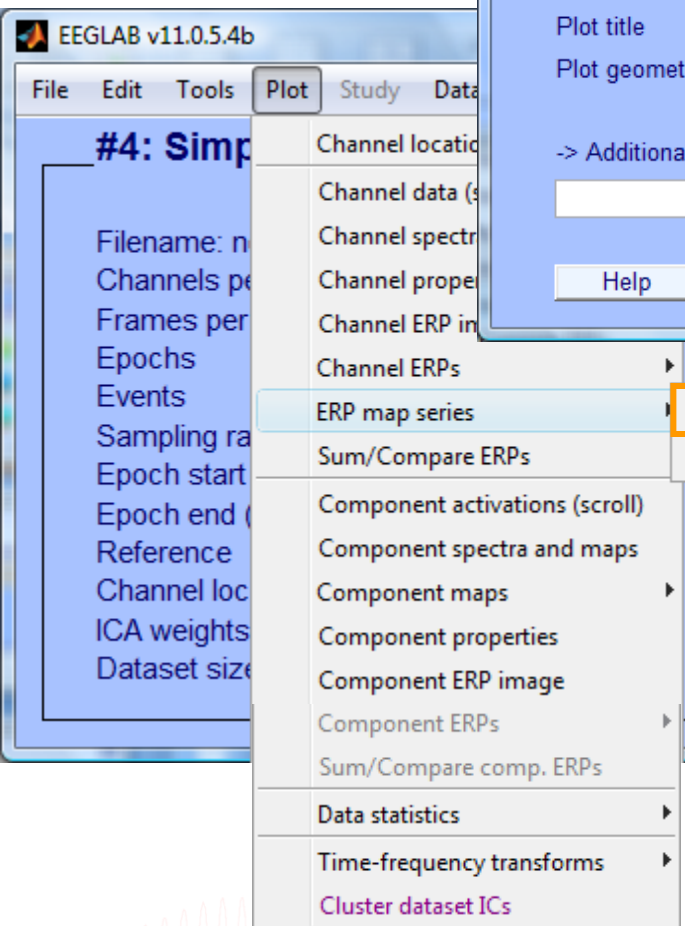


# Visualize ERP scalp distribution



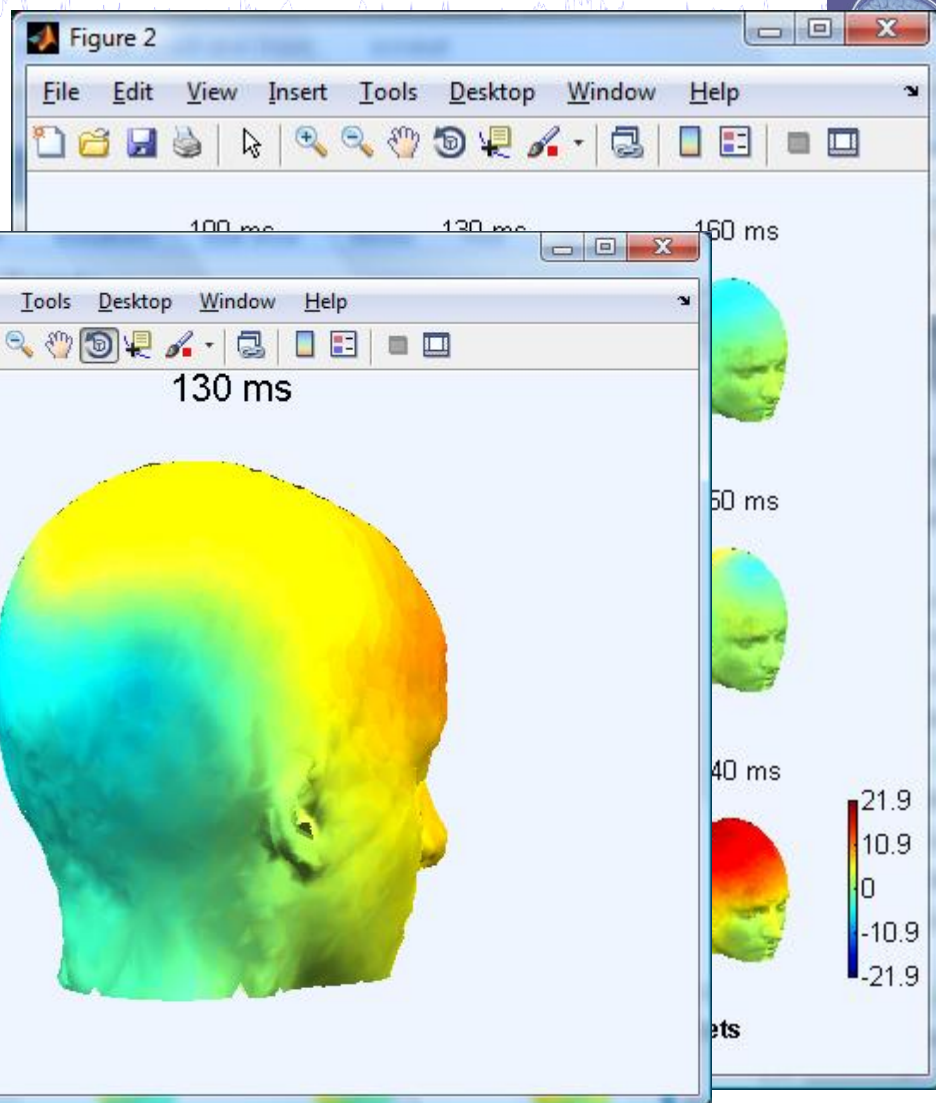
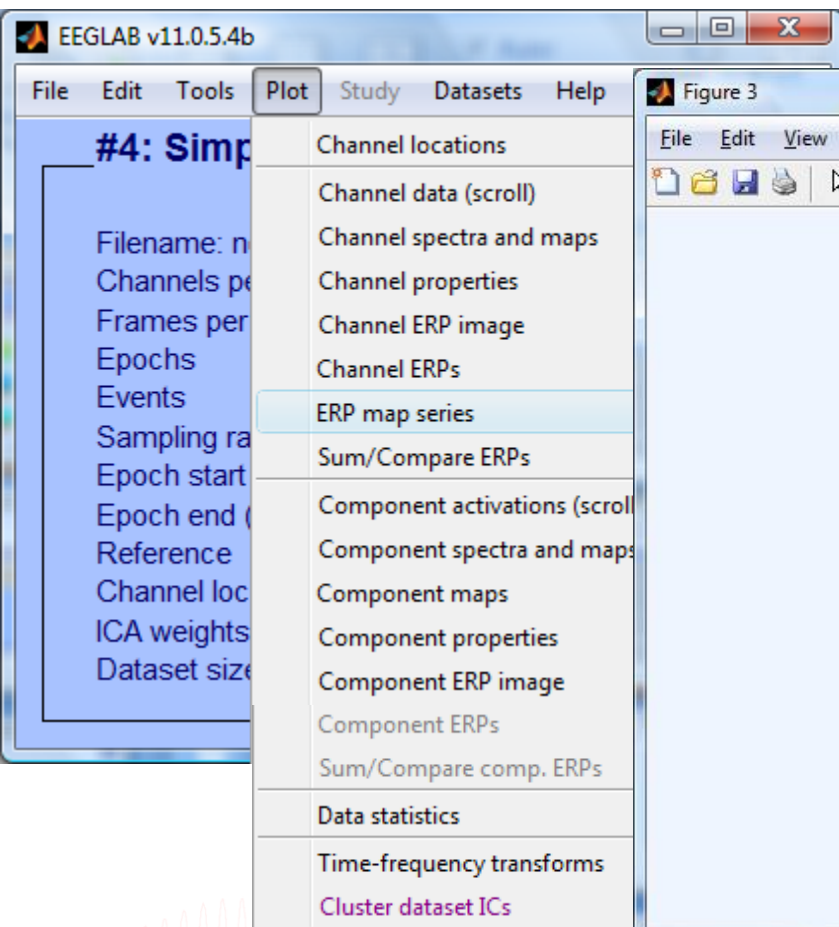
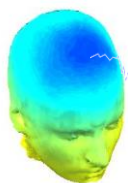


# Visualize channel ERPs in 2D





# Visualize channel ERPs in 3D

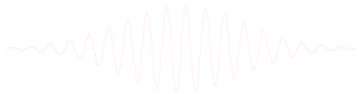




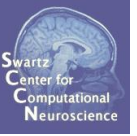
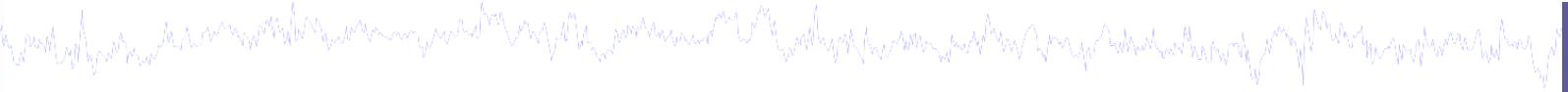
# Exercises (continuous data)



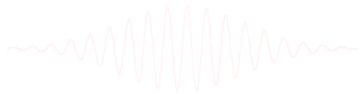
- Load SimpleOddball.set
- Re-reference data to average reference
- Hi-pass filter the continuous data, then save
- Epoch the data on circles (event type 1)
- Scroll the epoched data and perform visual rejection of epochs
- Explore the automated artifact rejection tools
- Save 'clean' epoched datasets for circles (targets)





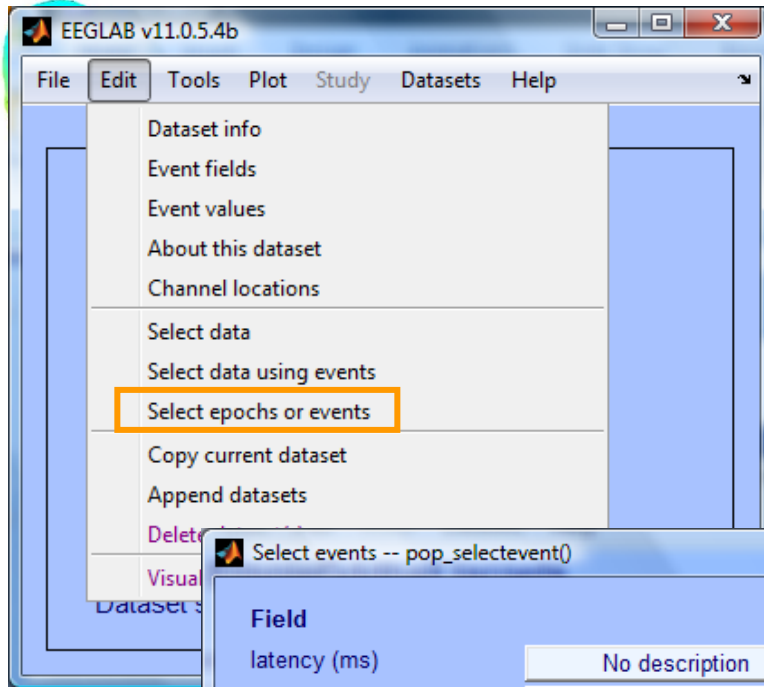


# Supplementary material

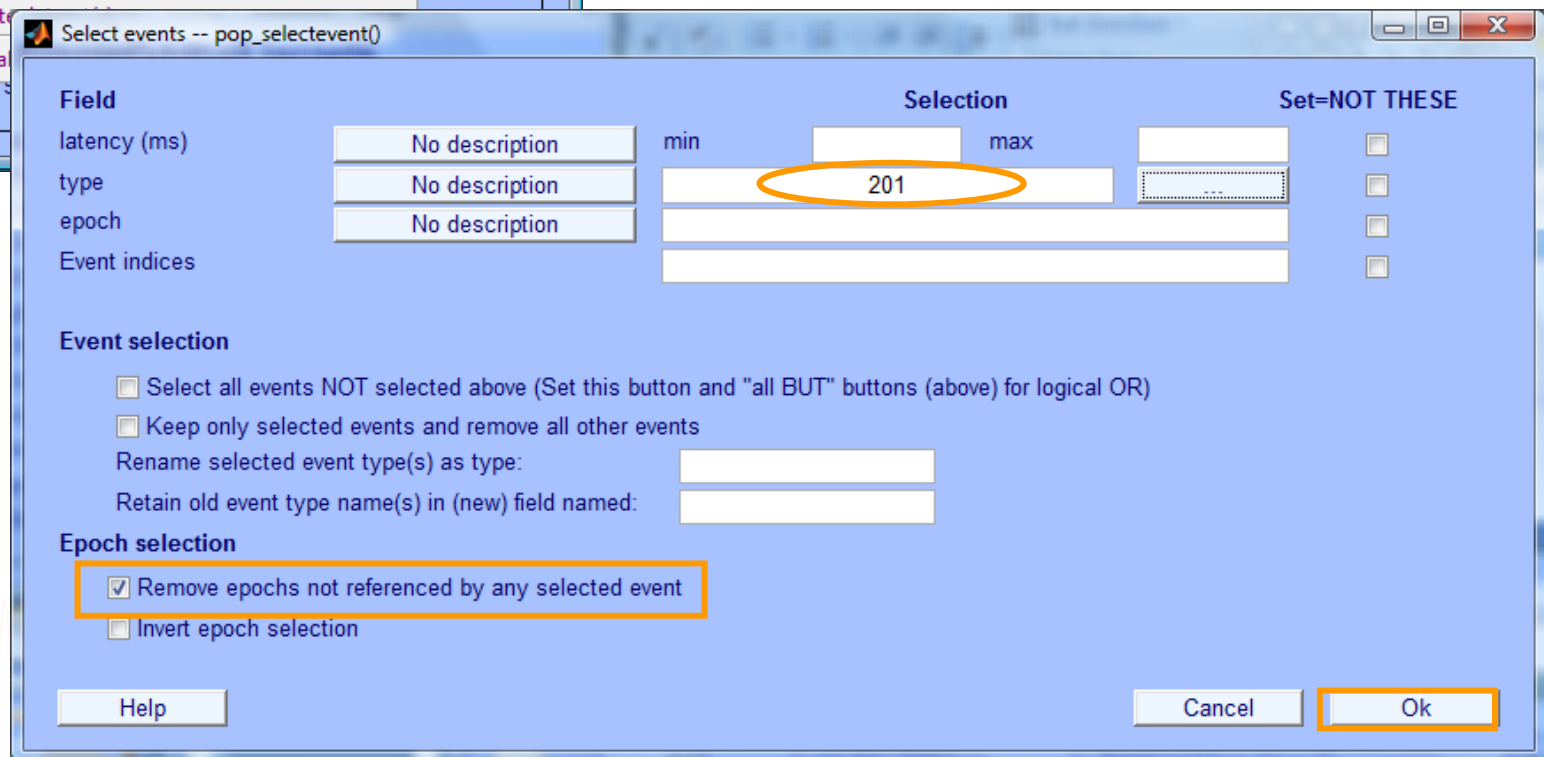




# Select a subset of epochs

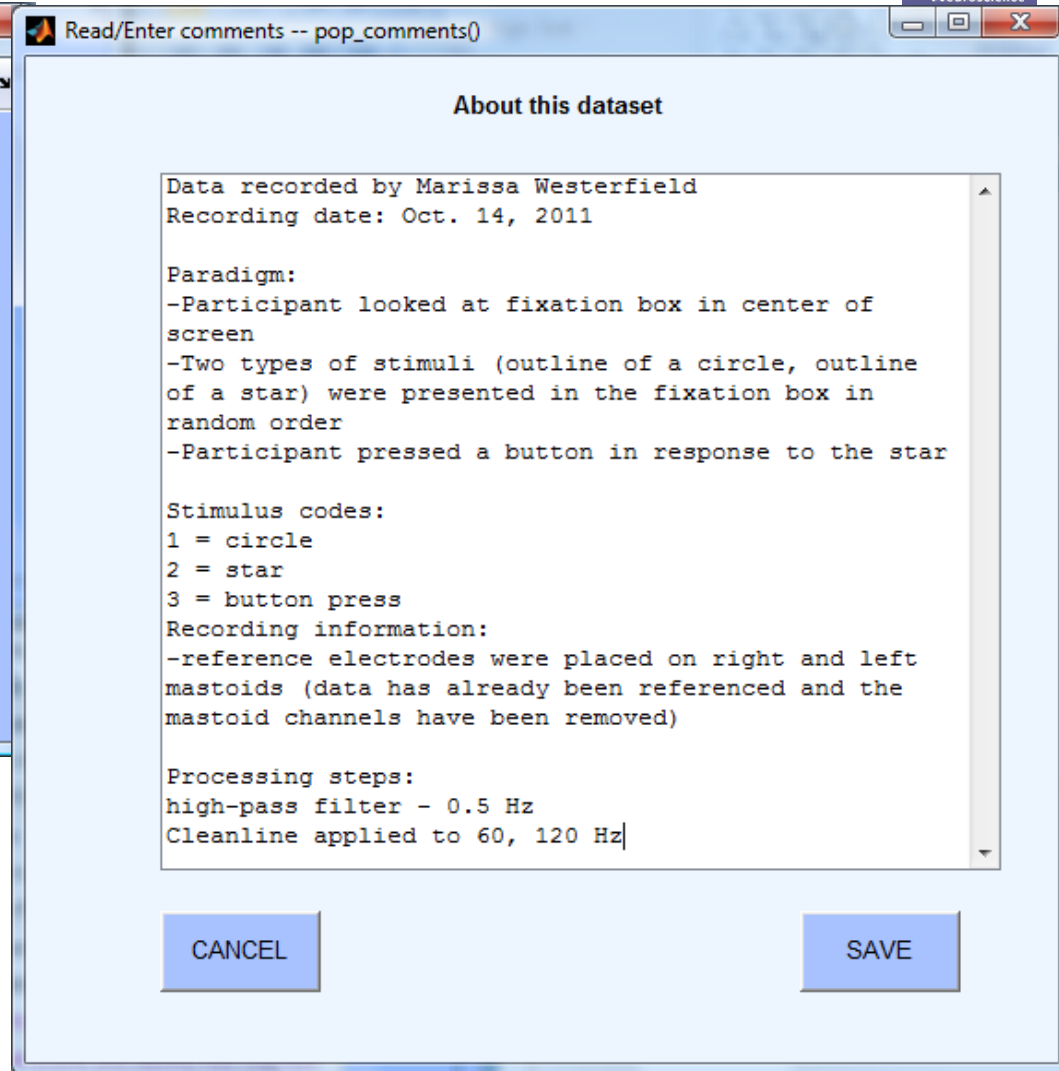
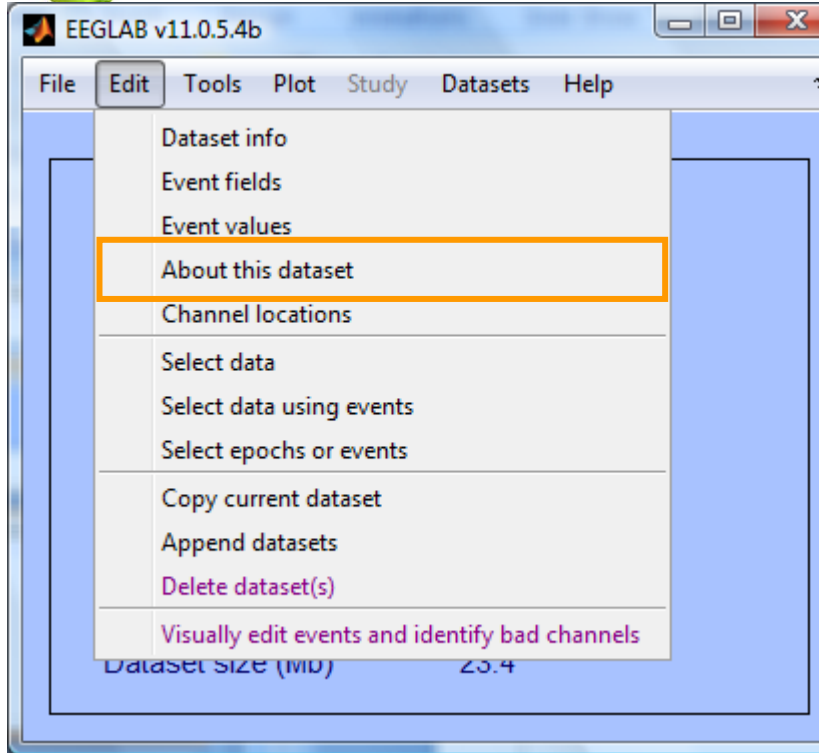
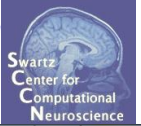
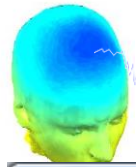


Keep only epochs with a 201 (button press) event





# Comments and dataset history



Also:

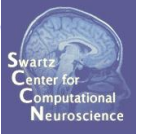
>> EEG.comments

or

>> EEG.history



# Removing channel(s)



- You may prefer to interpolate bad channels rather than remove them altogether
- The loss in dimensionality will affect the ICA decomposition
- Usual solution:
  - Delete the bad channels before running ICA
  - STUDY tools will do much of this automatically (interpolate missing channels, etc)

