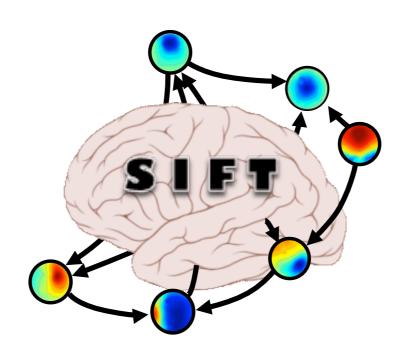
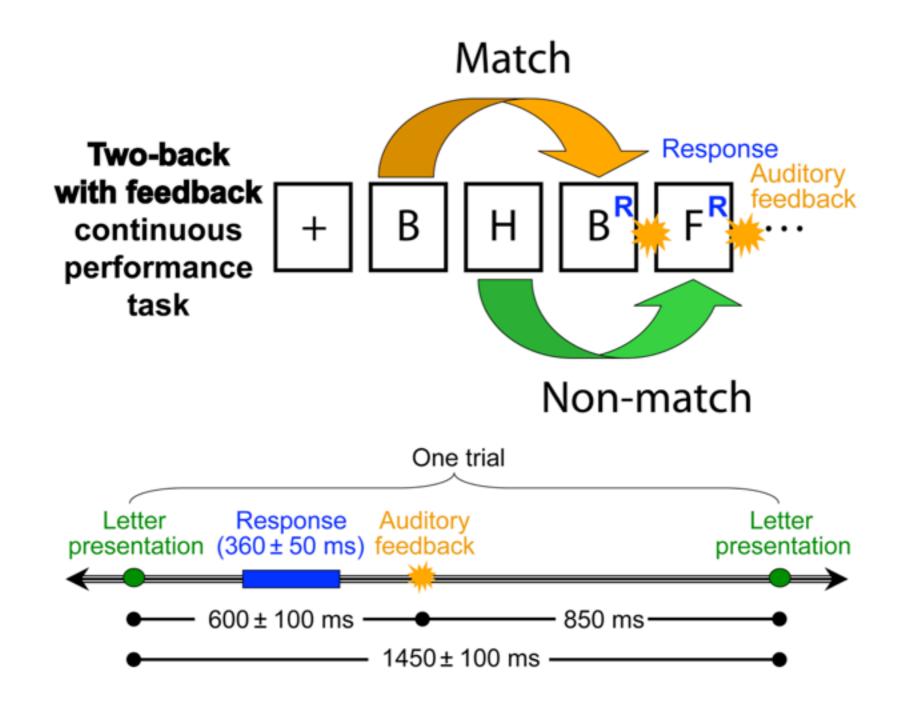
# The Source Information Flow Toolbox

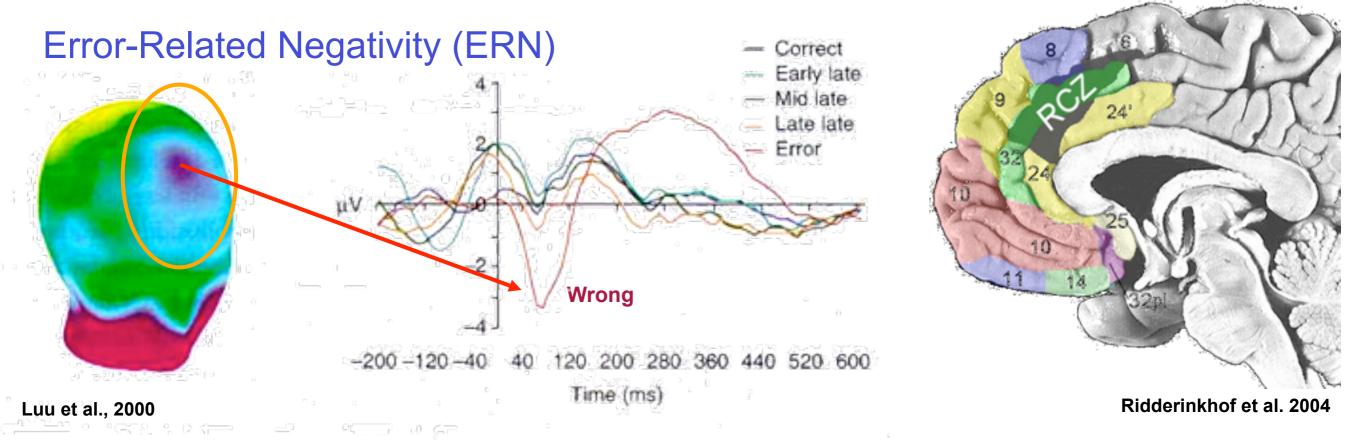


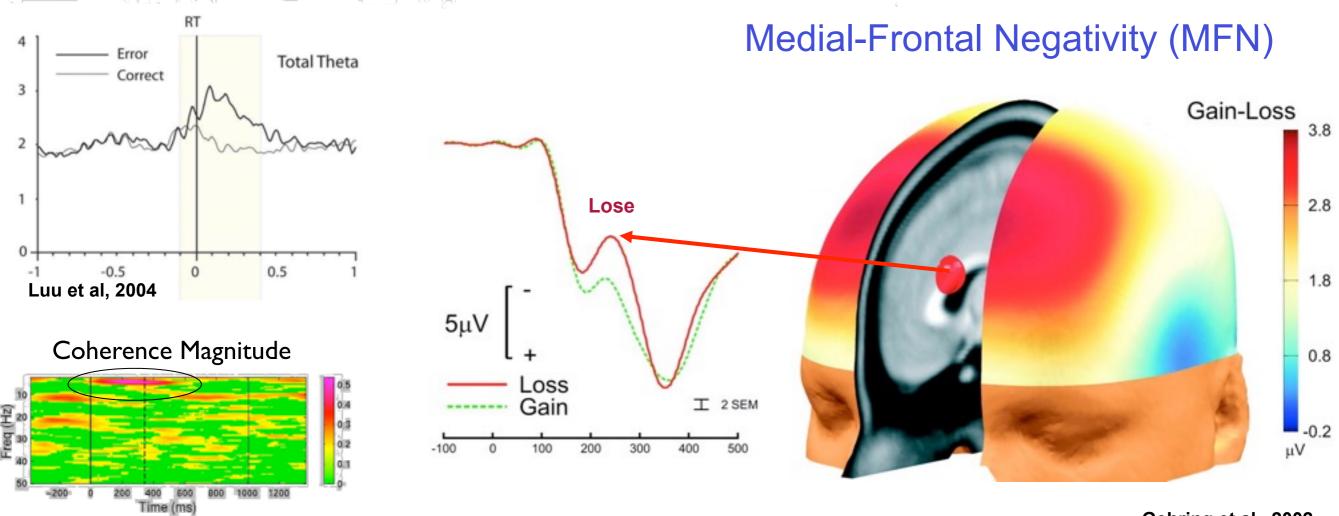
**Practicum** 

Tim Mullen

## Experimental Data

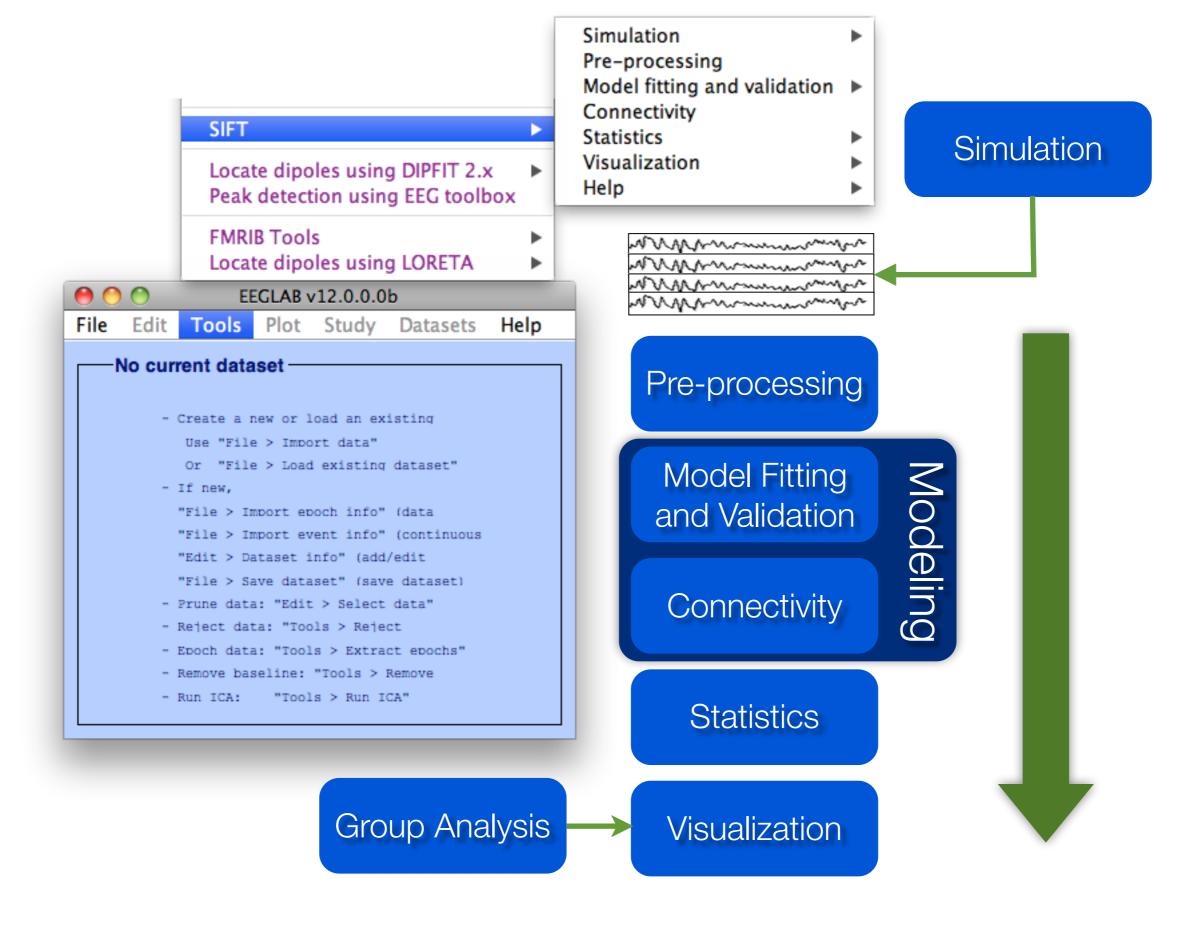






Makeig et al, 2001

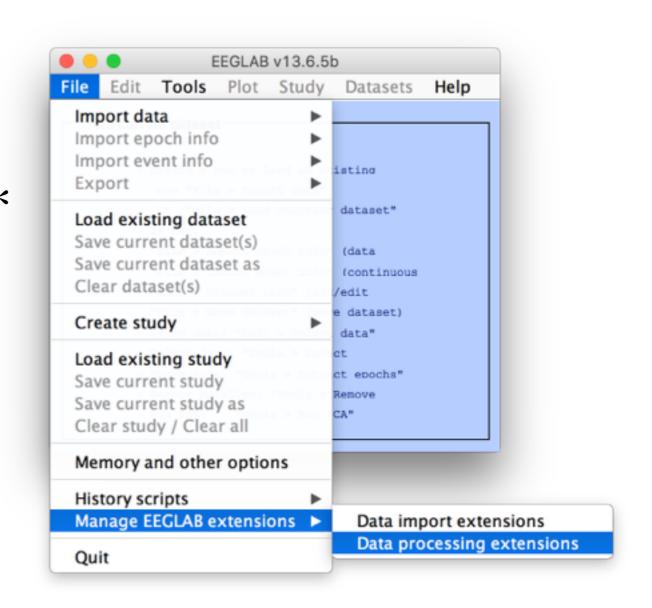
Gehring et al., 2002



# Starting EEGLAB/SIFT

#### SIFT Requirements:

- Matlab 2008b (or greater) \*
- Signal Processing Toolbox
- Statistics Toolbox
- EEGLAB
- SIFT 1.5



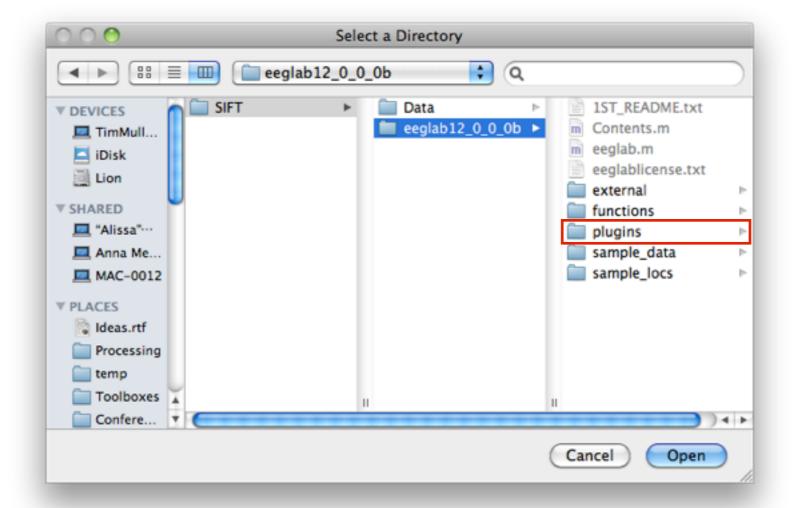
\* SIFT 1.5+ required for compatibility with Matlab 2014a or greater

# Starting EEGLAB/SIFT

If you can't install SIFT via plugin manager...

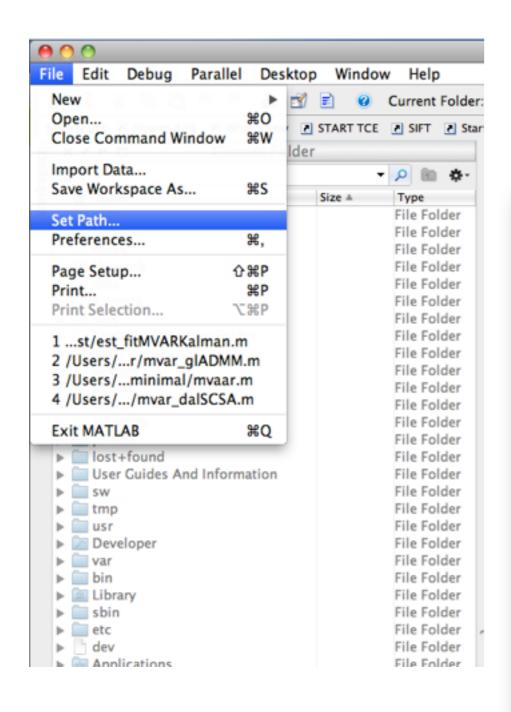
Copy <USB-key>/.../SIFT\_I.5 to EEGLAB plugins folder

(unzipped)

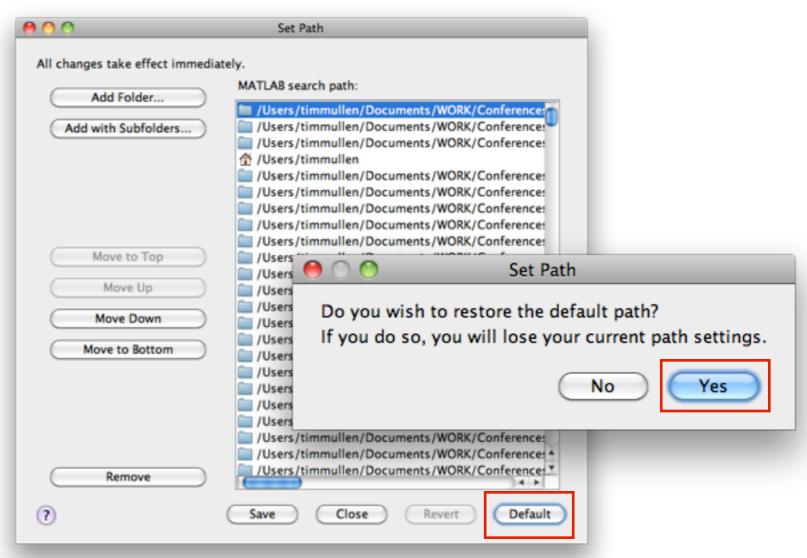




### If you don't have SIFT Installed

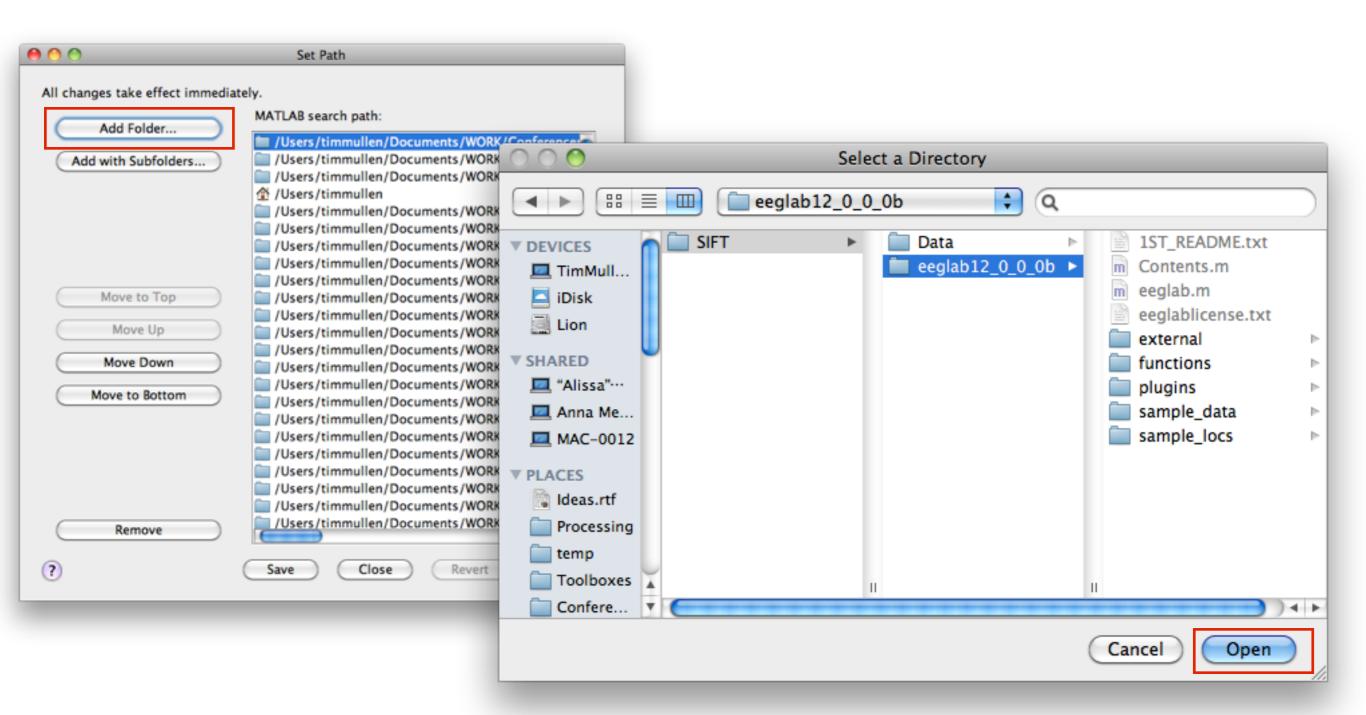


#### I. Clear the Matlab Path





### 2. Add EEGLAB+SIFT to path



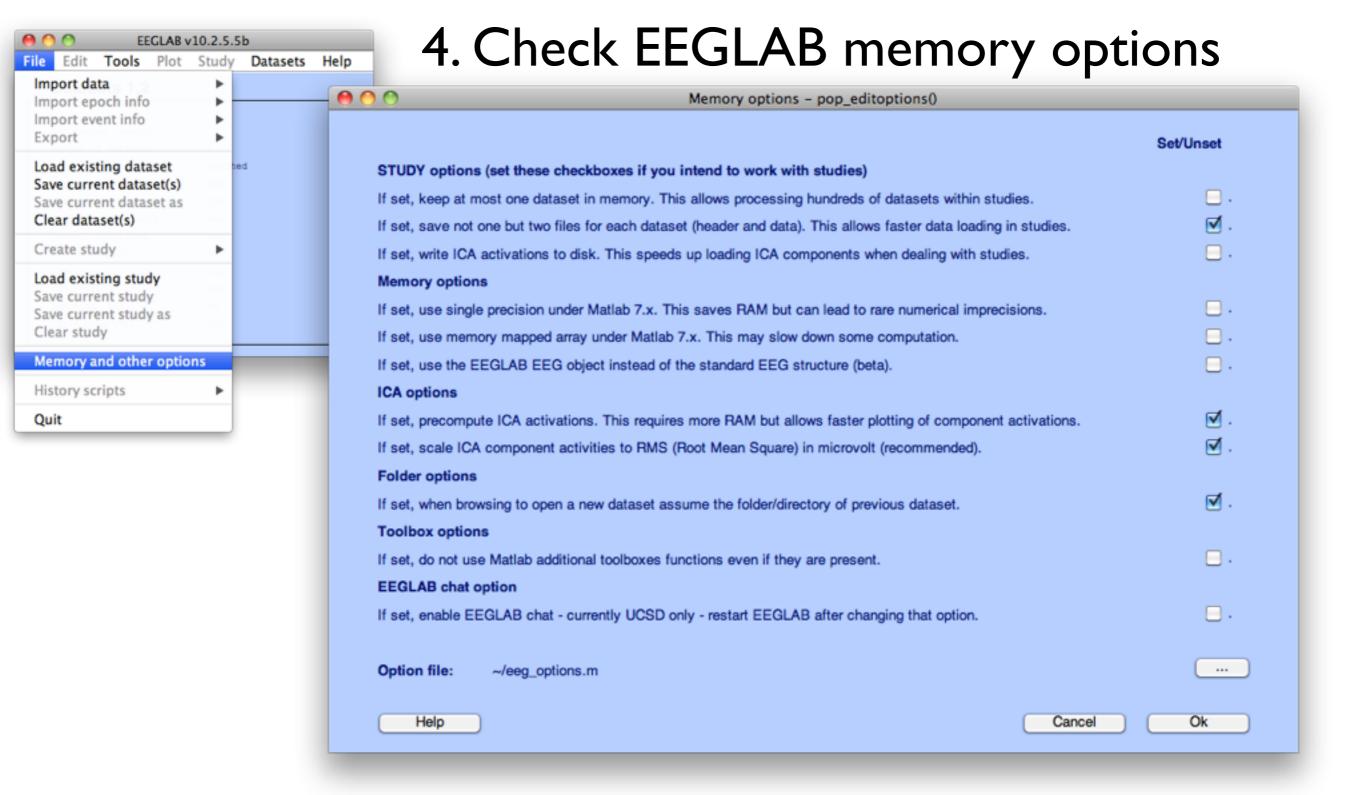
# Starting EEGLAB/SIFT

#### 3. Start EEGLAB

```
× ₹ → □ Command Window

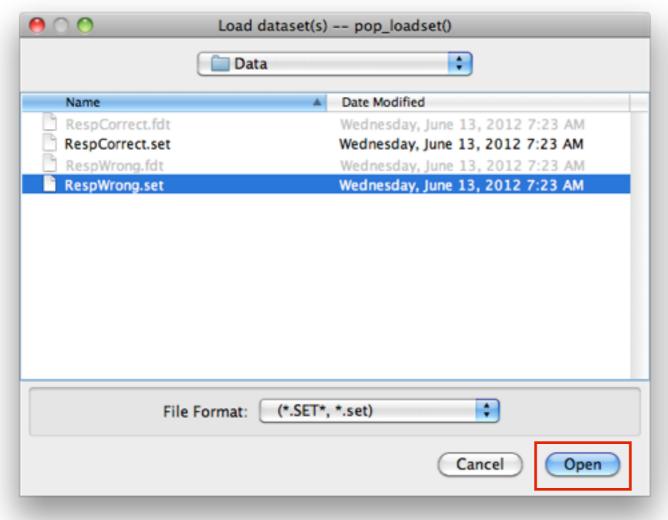
fx >> eeglab
```

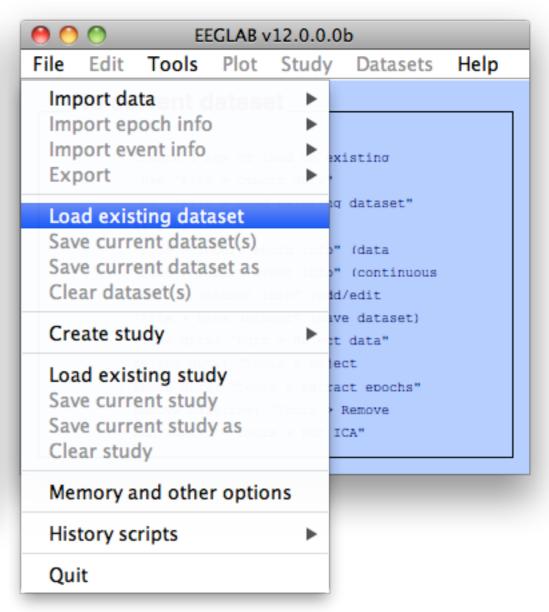




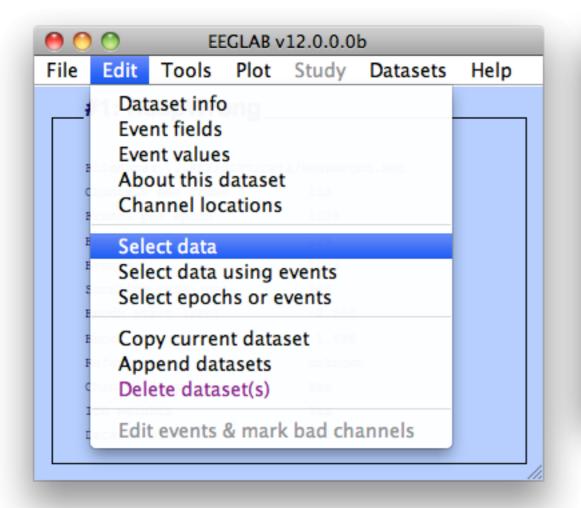
# 2 Loading Data

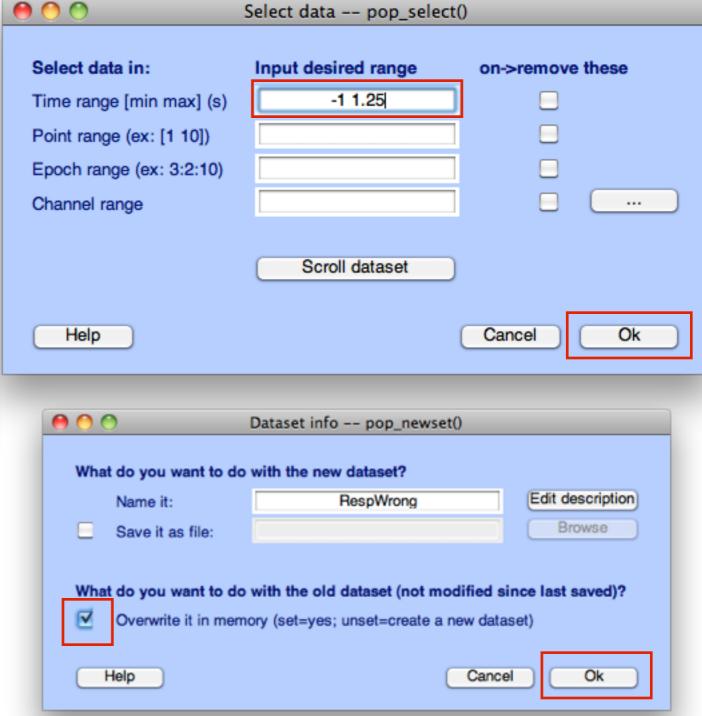
## <USB Key>/SIFT/Data/



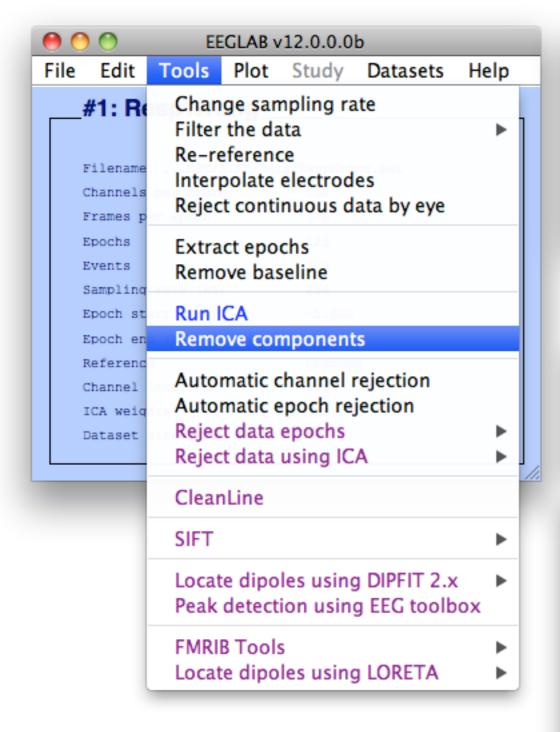


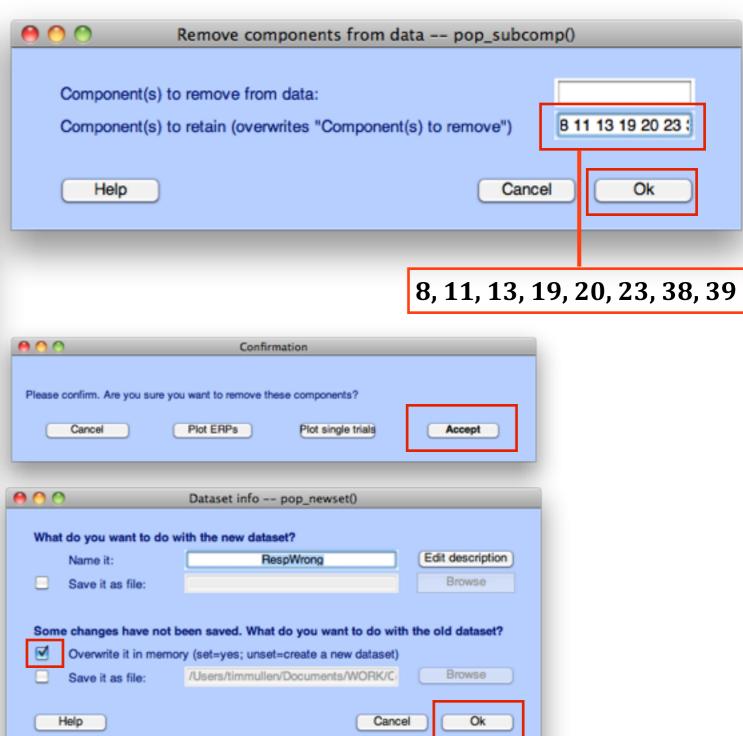
### Preprocessing: Select Data Range





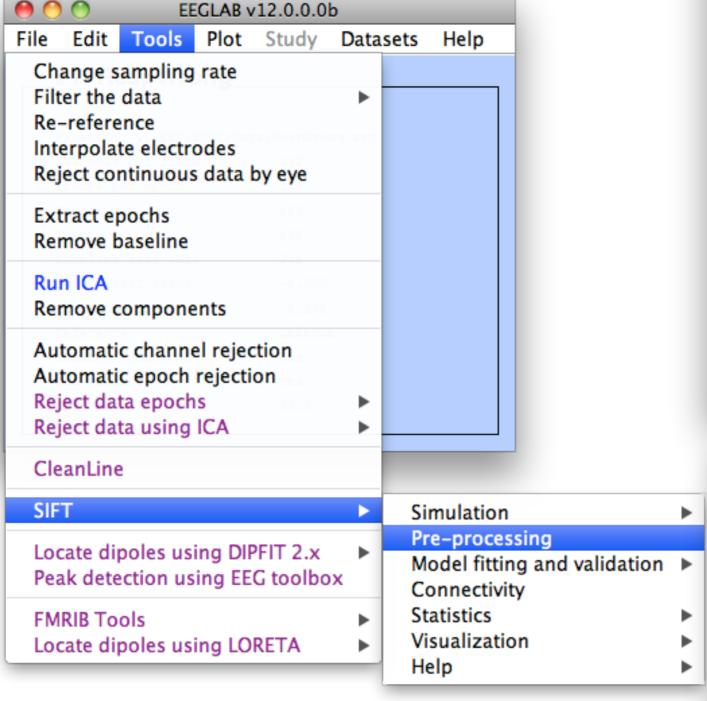
### Preprocessing: Select Components

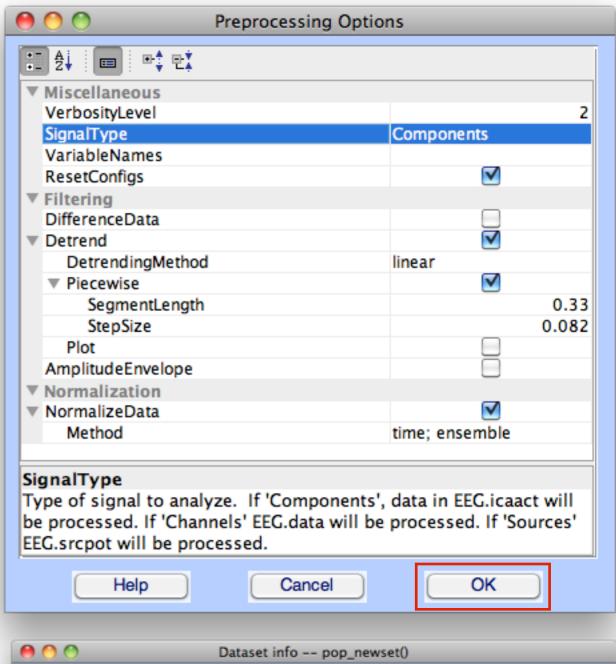




## Preprocessing: SIFT

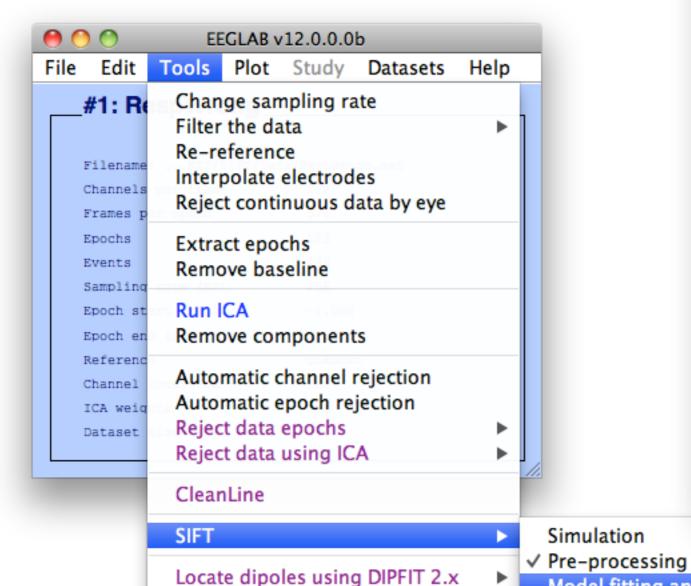
3





	Name it:	RespWrong	Edit description
3	Save it as file:		Browse
	1	been saved. What do you want to do with th	ne old dataset?
Som	1	been saved. What do you want to do with the	ne old dataset?

## Model Order Selection



Peak detection using EEG toolbox

Locate dipoles using LORETA

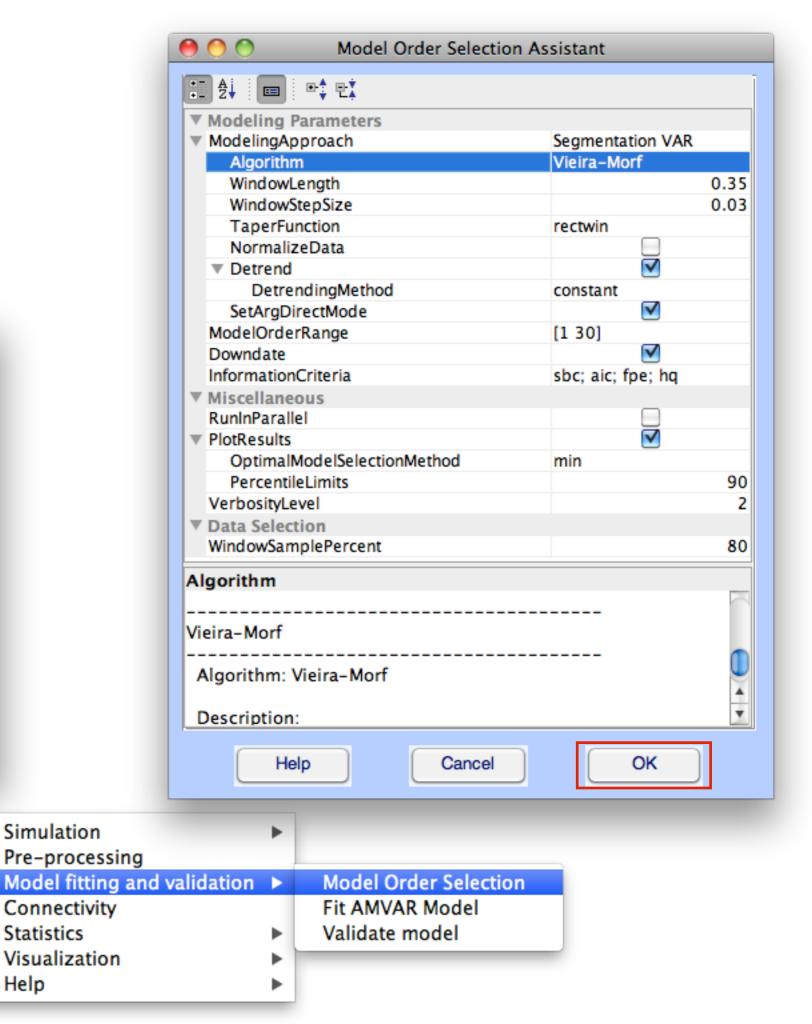
**FMRIB Tools** 

Connectivity

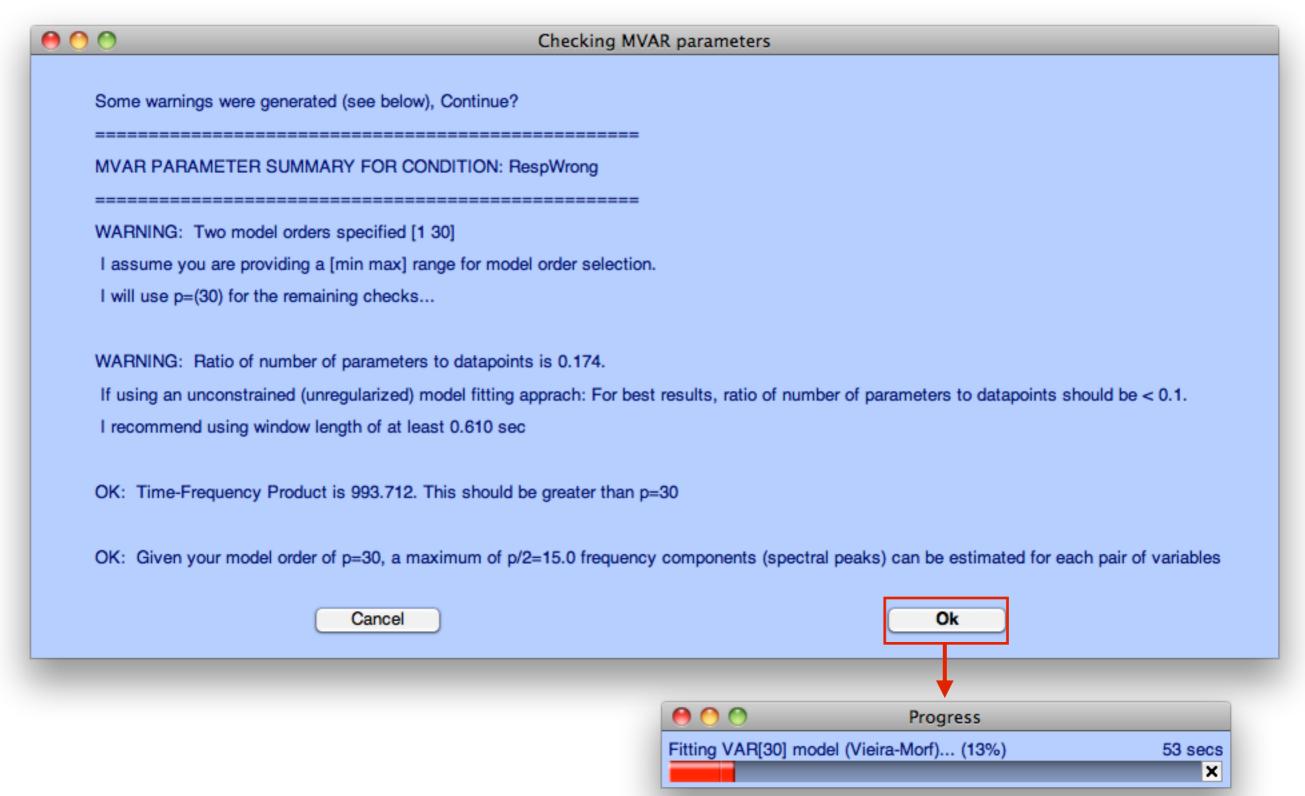
Visualization

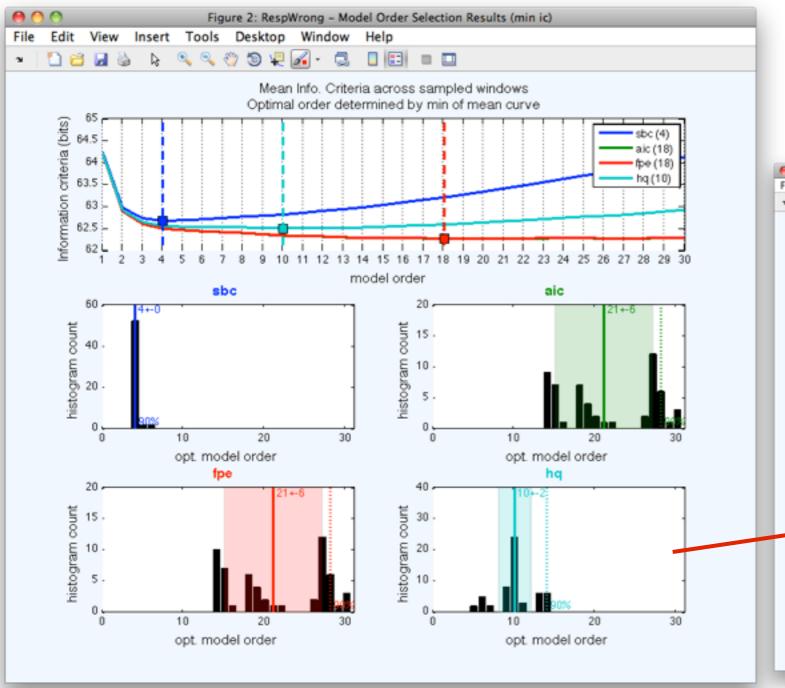
Statistics

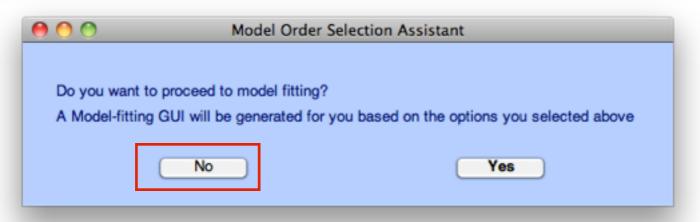
Help

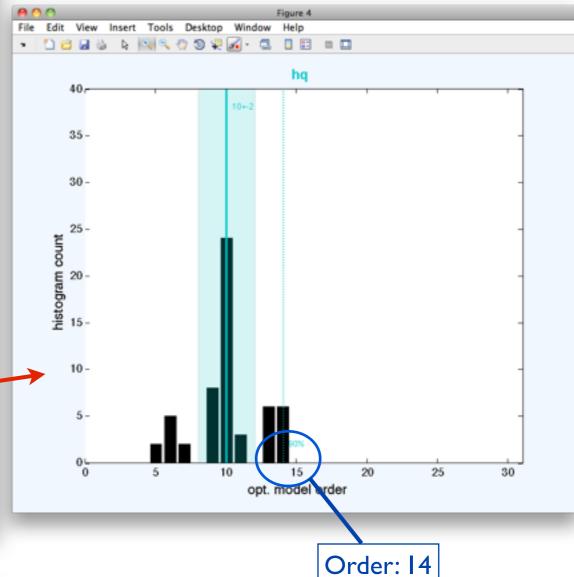


# 4 Model Order Selection

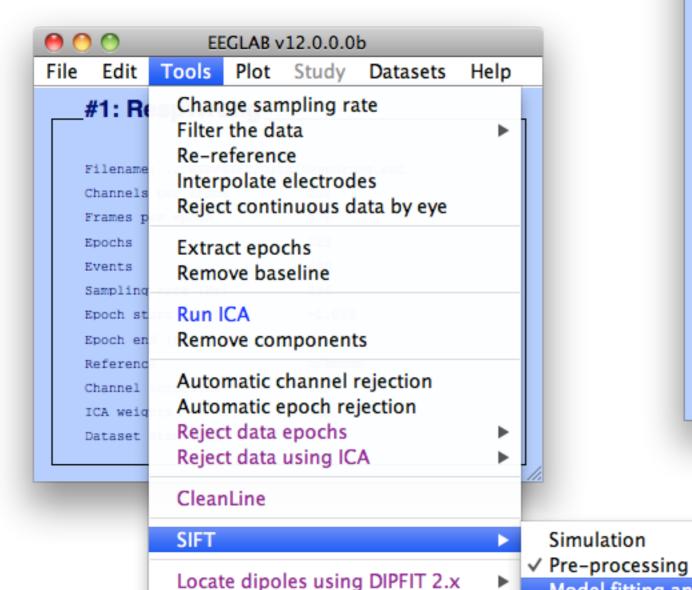








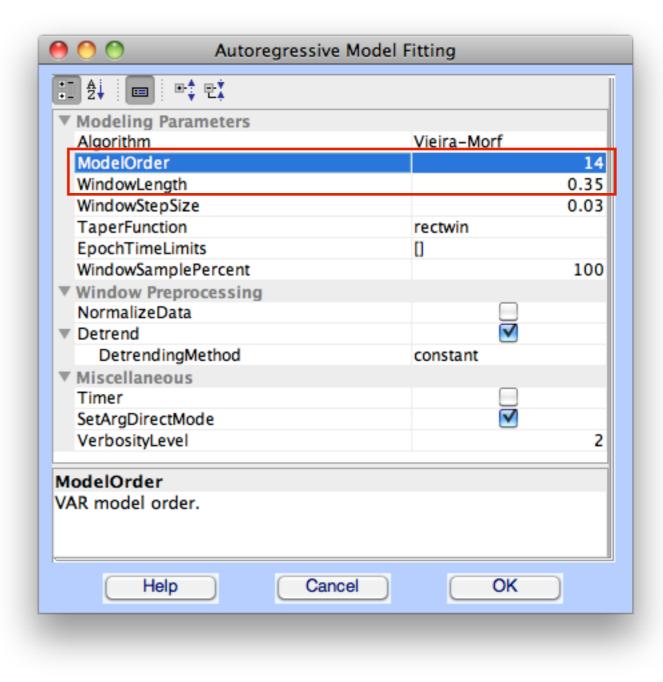
## 5 Model Fitting



Peak detection using EEG toolbox

Locate dipoles using LORETA

**FMRIB Tools** 



✓ Model Order Selection

Fit AMVAR Model

Validate model

Model fitting and validation ▶

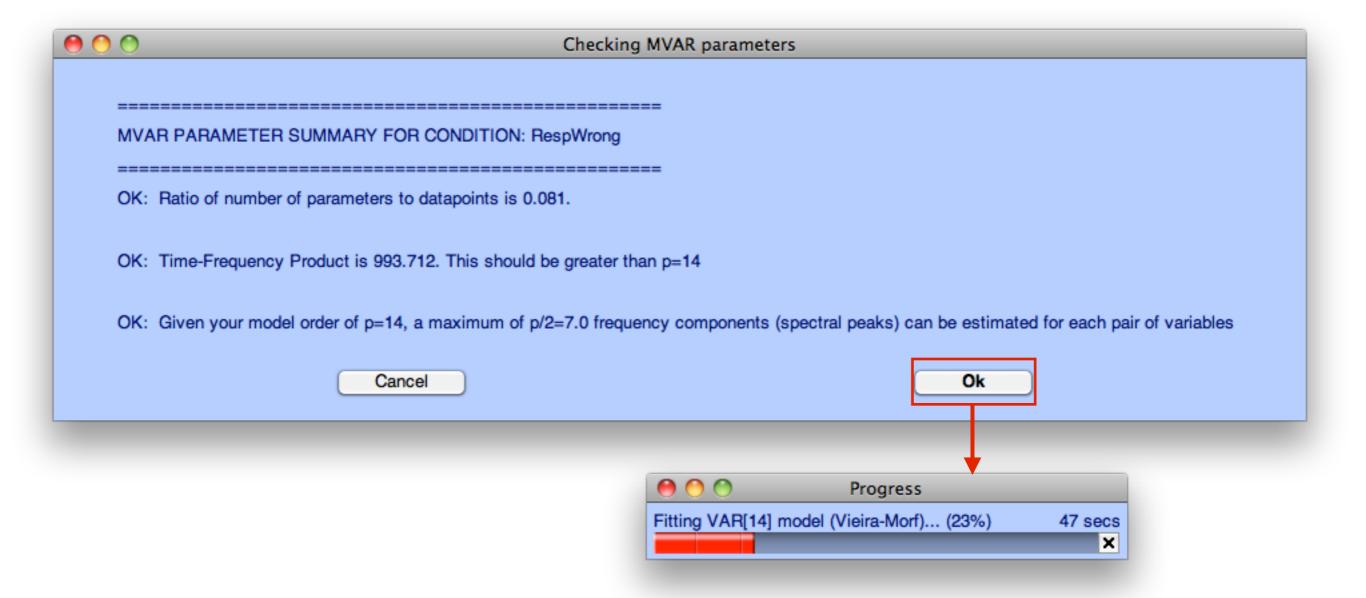
Connectivity

Visualization

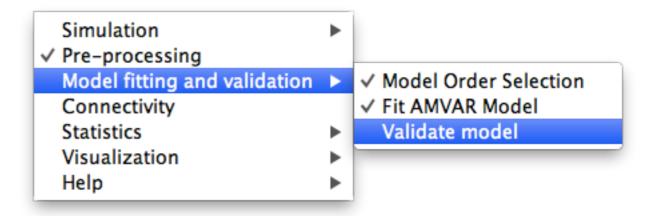
Statistics

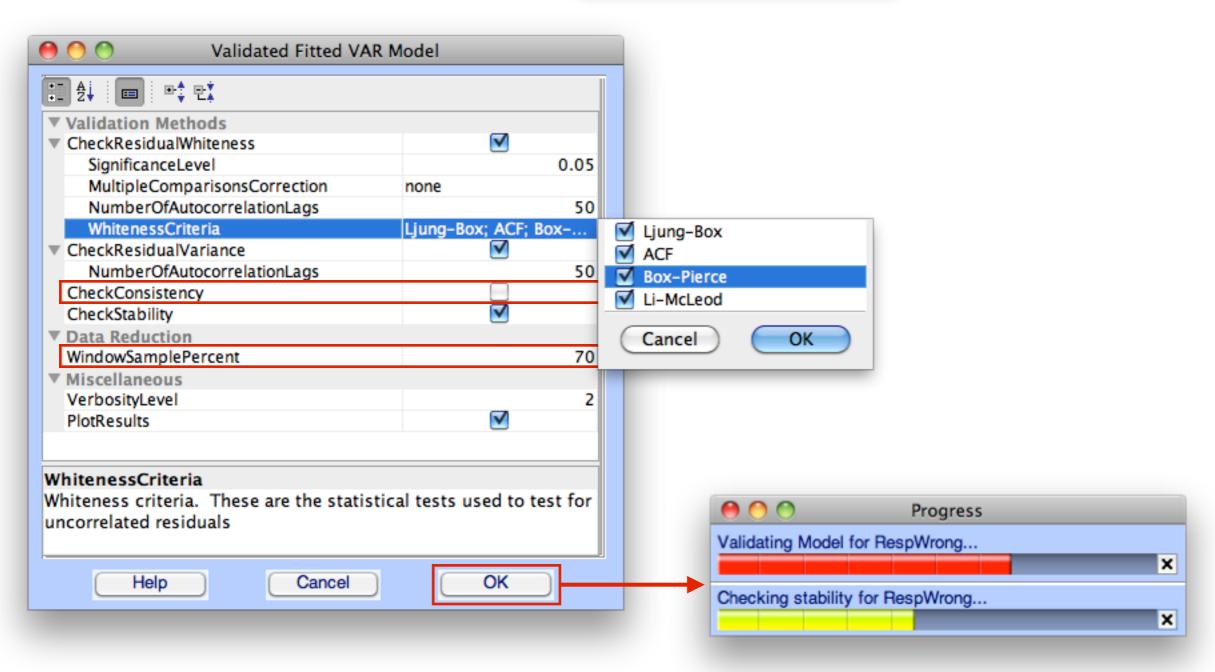
Help

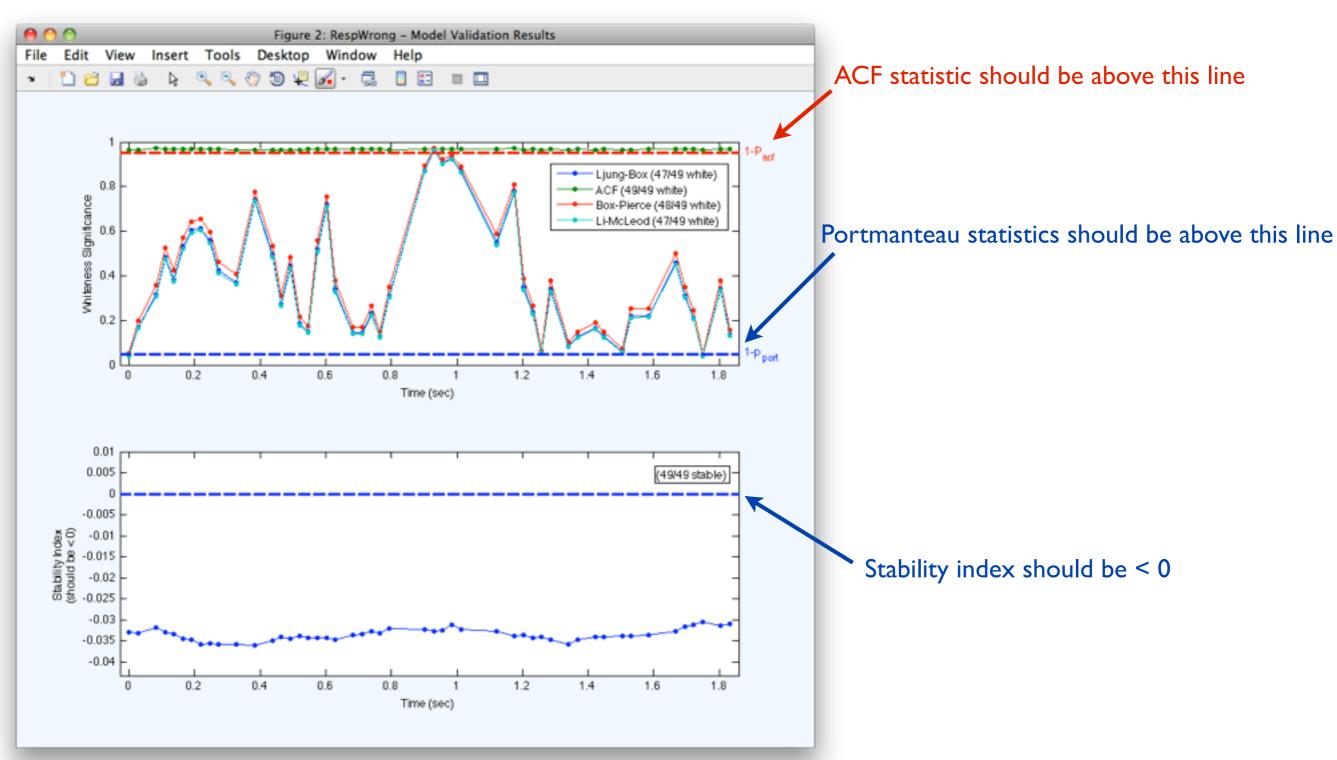
## 5 Model Fitting



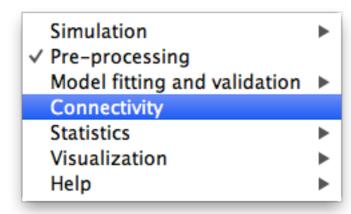
## 6 Model Validation

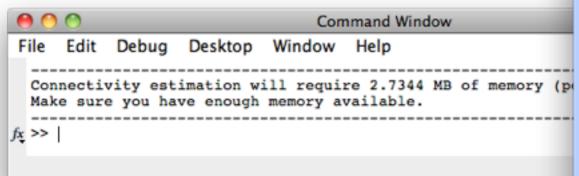


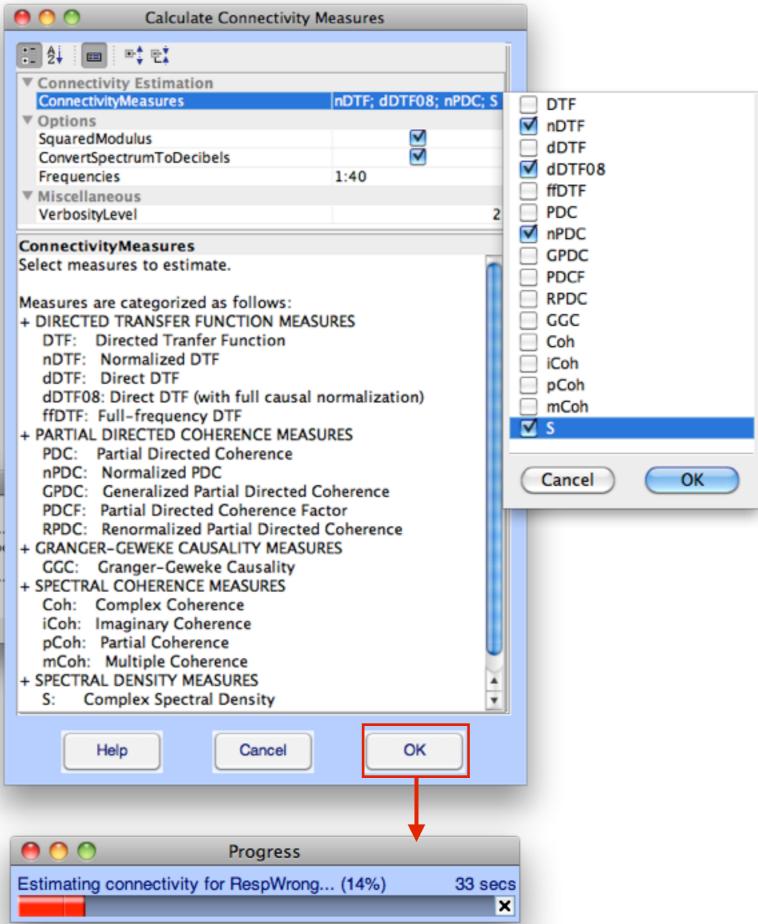




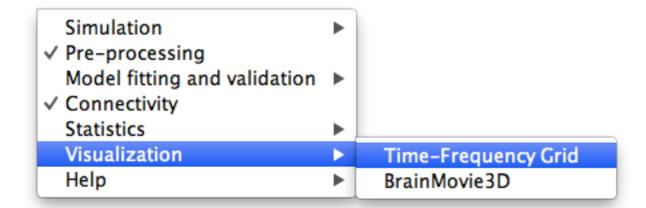
## 7 Connectivity



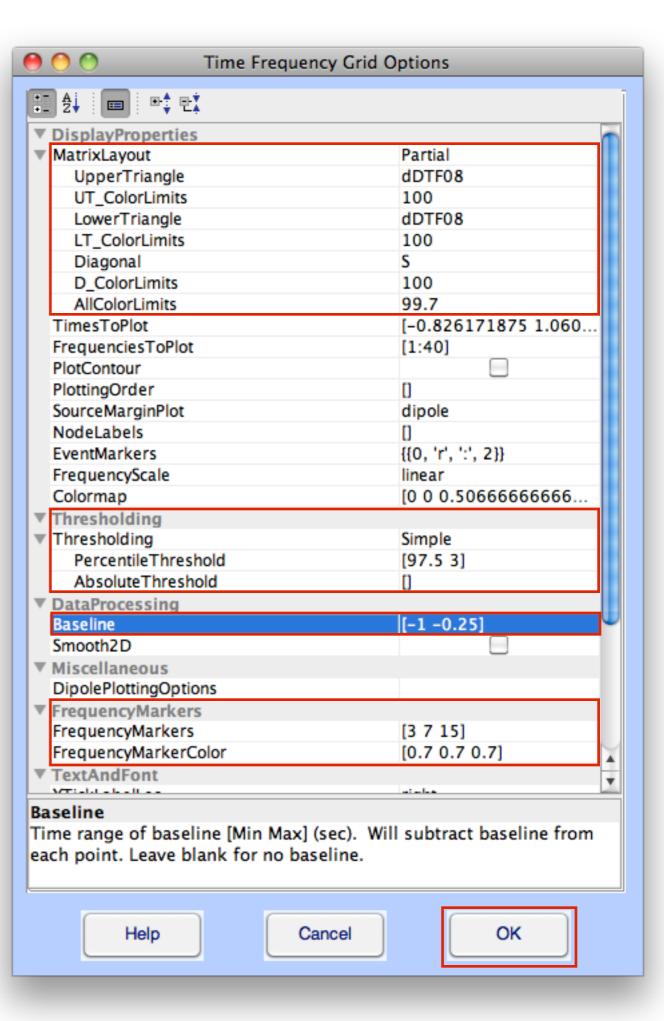




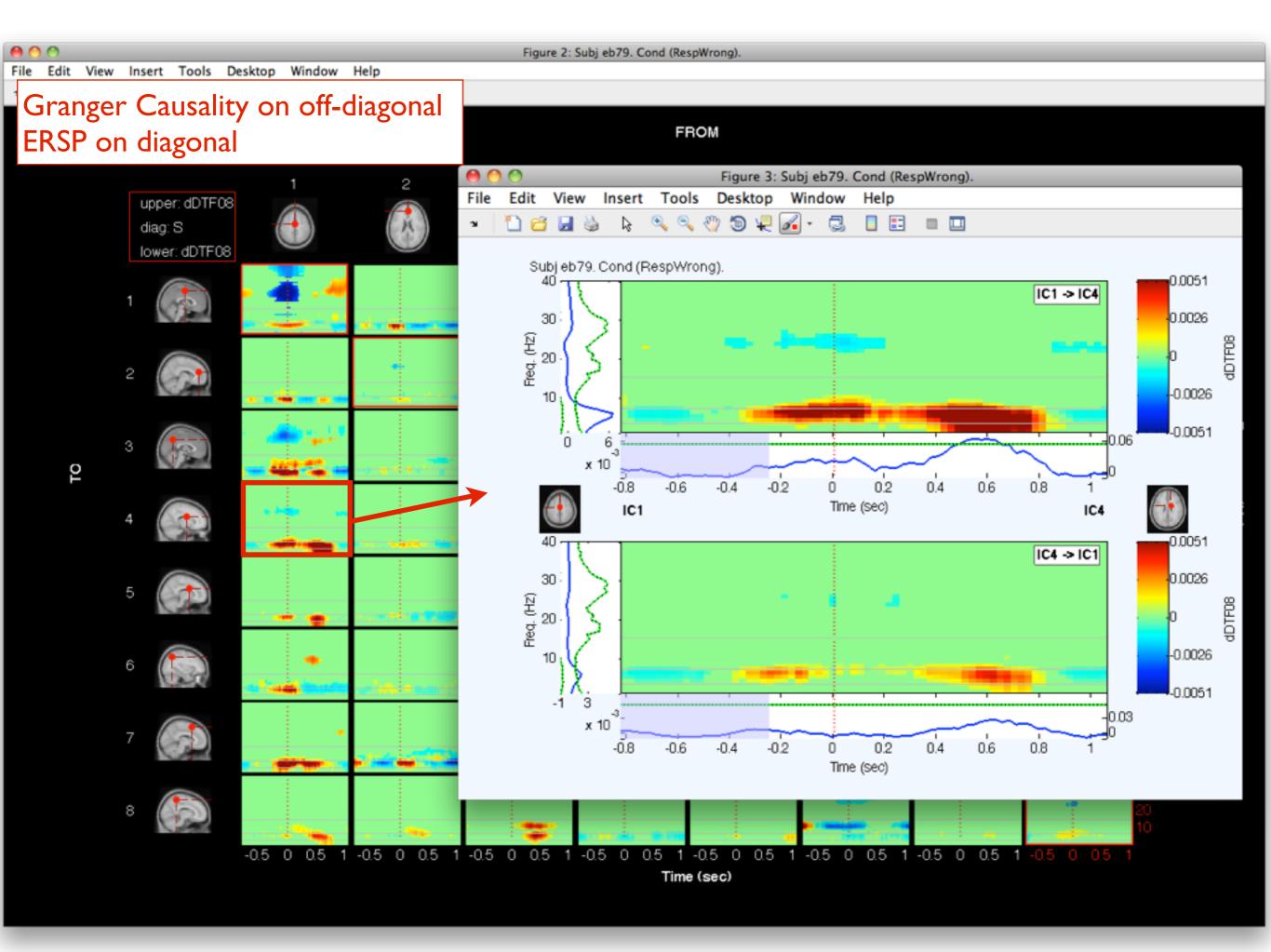
## Visualization: Time-Frequency Grid

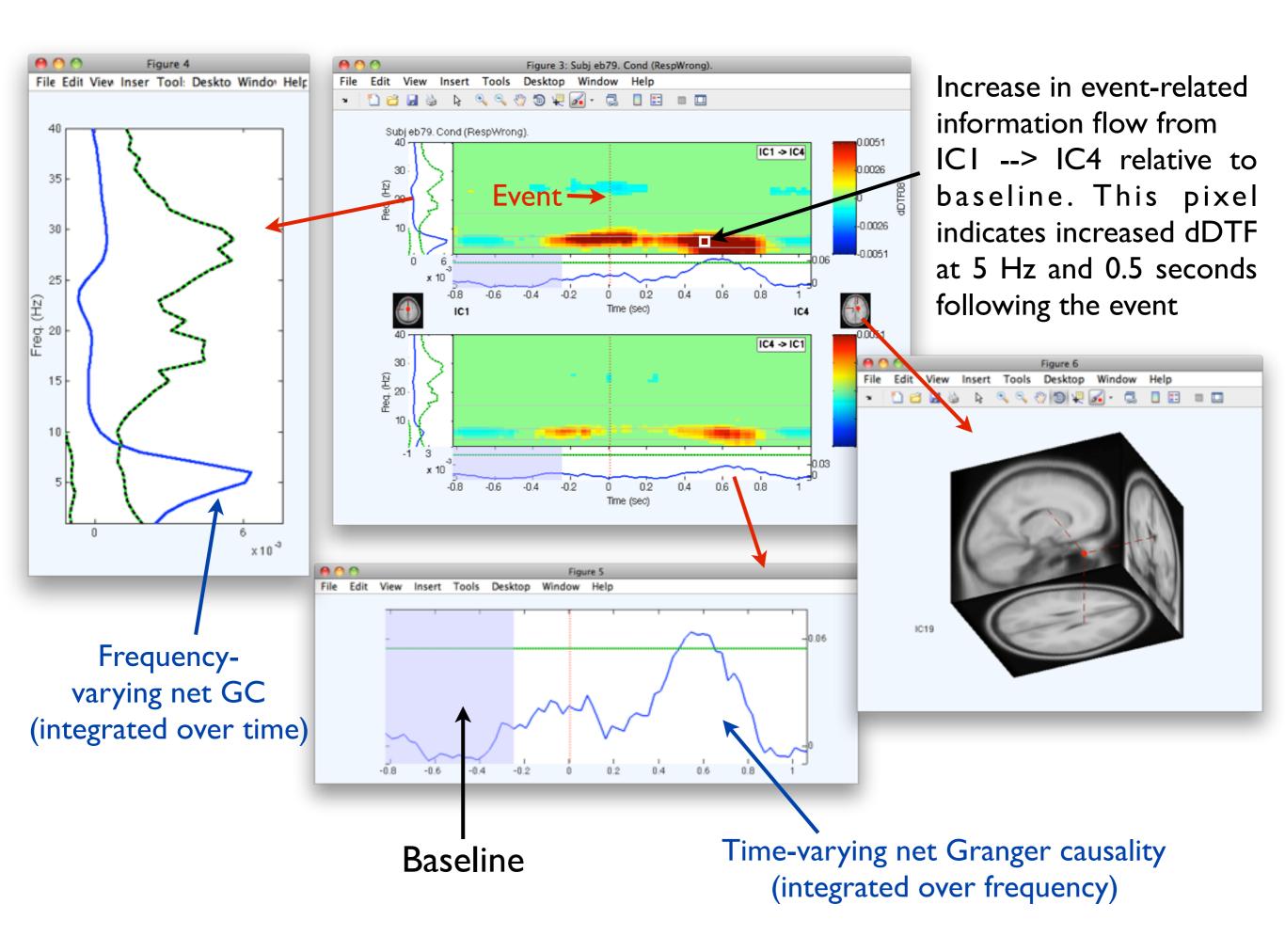


8



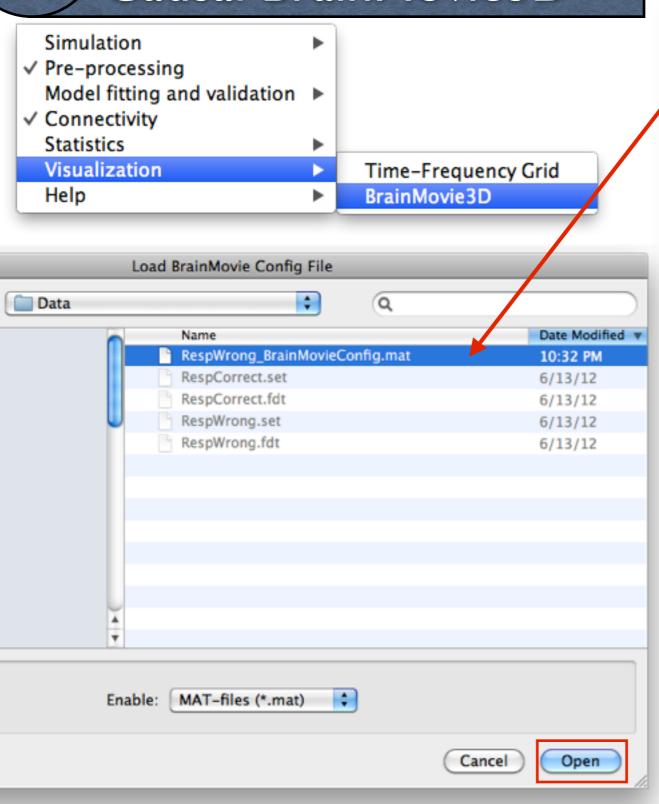


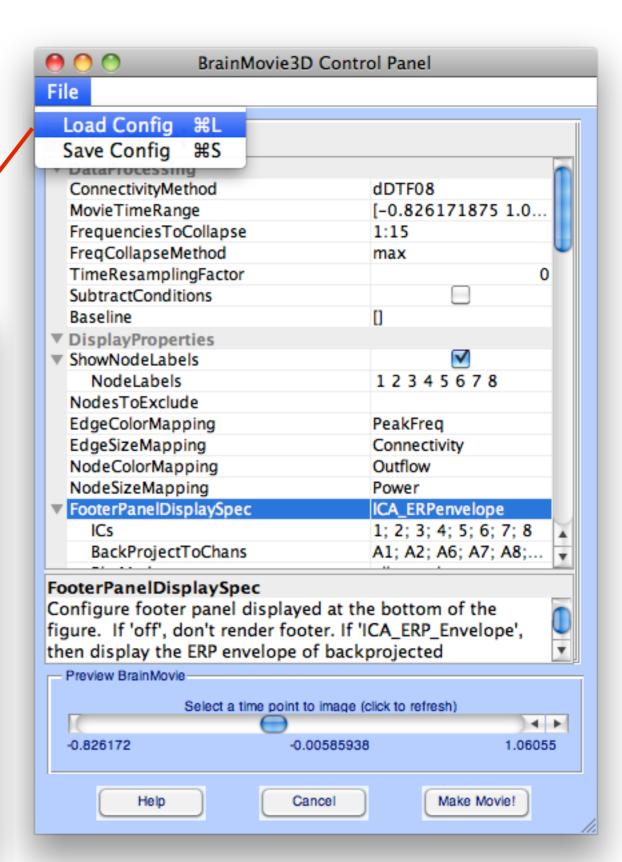




## Visualization: Causal BrainMovie3D

9





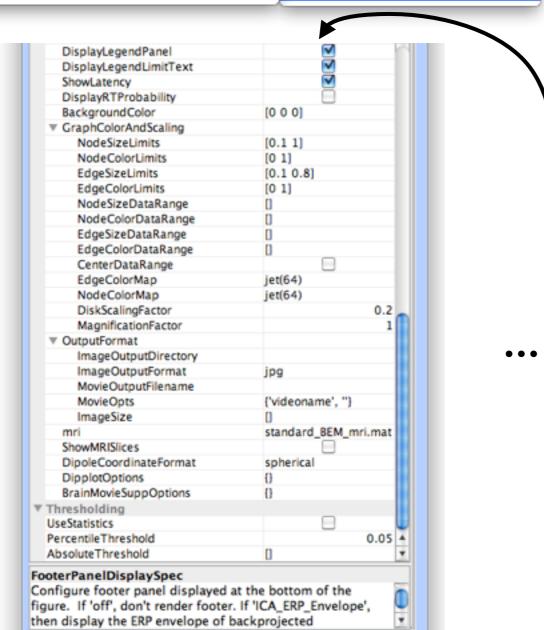
### Visualization: Causal BrainMovie3D

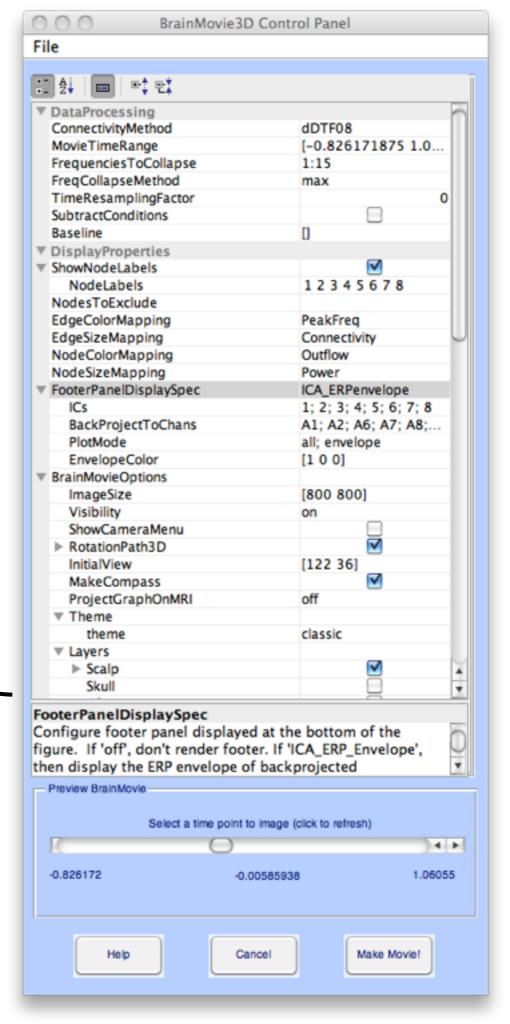
Simulation √ Pre-processing Model fitting and validation ▶ √ Connectivity Statistics Visualization Time-Frequency Grid

9

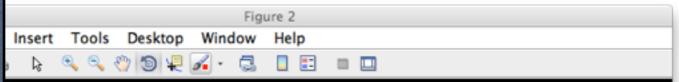
BrainMovie3D

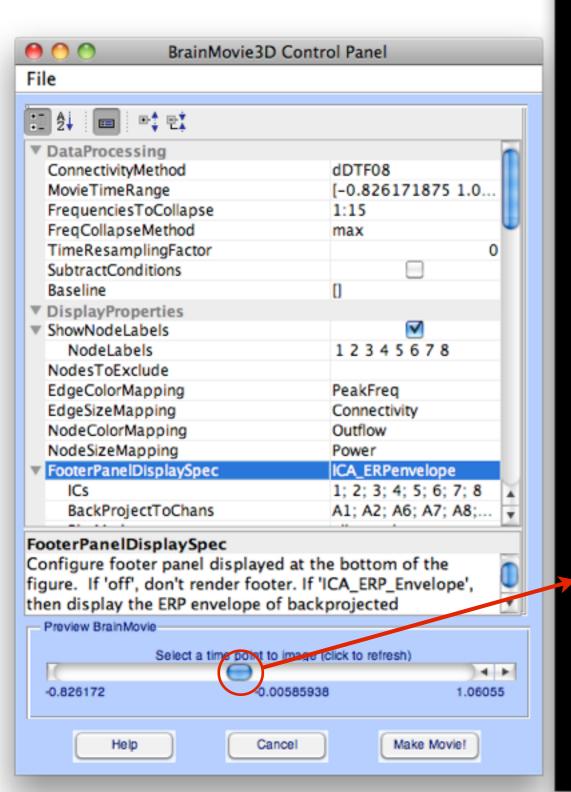
Help

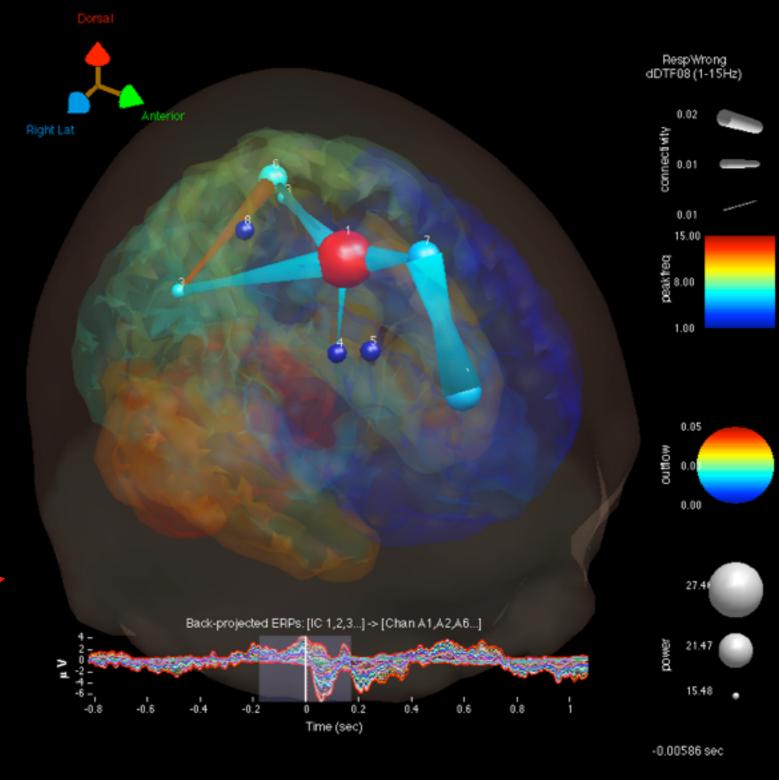




### 9 Visualization: Causal BrainMovie3D





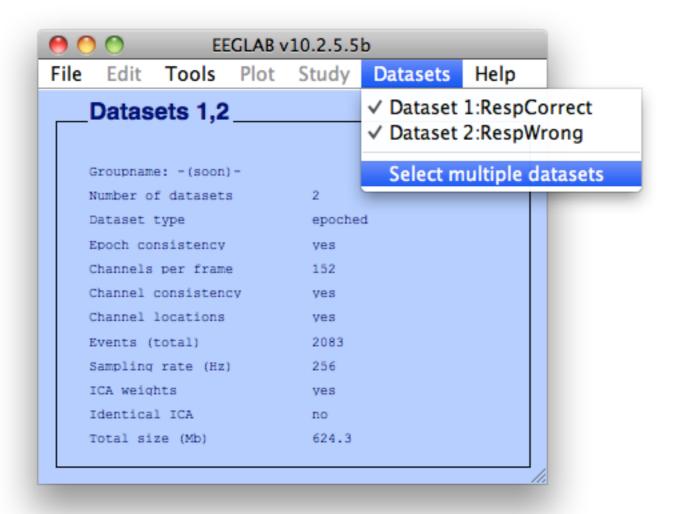


## 10 Additional Exercises

- 1. Explore changing some of the Time-Frequency Grid parameters. Try plotting the TF-Grid with logarithmic frequency spacing (option: FrequencyScale). Change the SourceMarginPlot to "topoplot" to see your ICA topographic plots.
- 2. Explore different parameters for the BrainMovie3D. What is different between delta (1-3 Hz) and theta (3-7 Hz) band connectivity?
- 3. Recompute connectivity for the RespWrong condition, selecting the Coherence (Coh) and Partial Coherence (pCoh) methods in addition to the original nPDC, nDTF, dDTF08, and S methods. Create a Time-Frequency Grid laying out Coherence (Coh) on the UpperTriangle, Partial Coherence (pCoh), on the LowerTriangle and the ERSP (S) on the diagonal. Use a baseline of [-1 -0.25]. What is different between coherence and partial coherence? Create another Time-Frequency Grid with dDTF08 on the Upper and Lower Triangles. What is different between coherence and dDTF (Granger-Causality)?
- 4.Redo the entire pipeline (Steps 1-9) for the RespCorrect condition (located in same /Data folder as RespWrong). Select both conditions in EEGLAB (Datasets-->Select Multiple Datasets). Create a Time-Frequency Grid. Choose to the plot the difference RespWrong-RespCorrect (option: PlotConditionDifference->ConditionOrder) with the dDTF08 on the Upper and Lower Triangle and ERSP on the diagonal.
- 5. Advanced Users: Try executing the previous pipeline entirely from the command line. Consult <sift\_root>/scripts/ScriptingExample\_1\_0b.m for guidance.

## Visualization of condition differences

## Select RespWrong and RespCorrect datasets



## Visualization of condition differences

