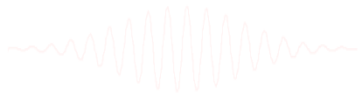
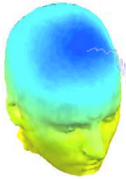


# Clustering of ICA components

Arnaud Delorme

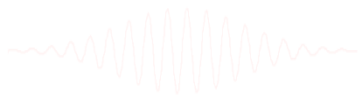
(with Julie Onton, Romain Grandchamp, Nima Bigdely Shamlo, Scott Makeig)

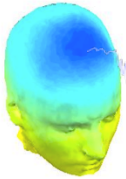




## Outline

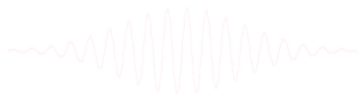
- ICA clusters and reliability within subjects
- ICA clusters and reliability across subjects
- Clustering in EEGLAB theory & Practice

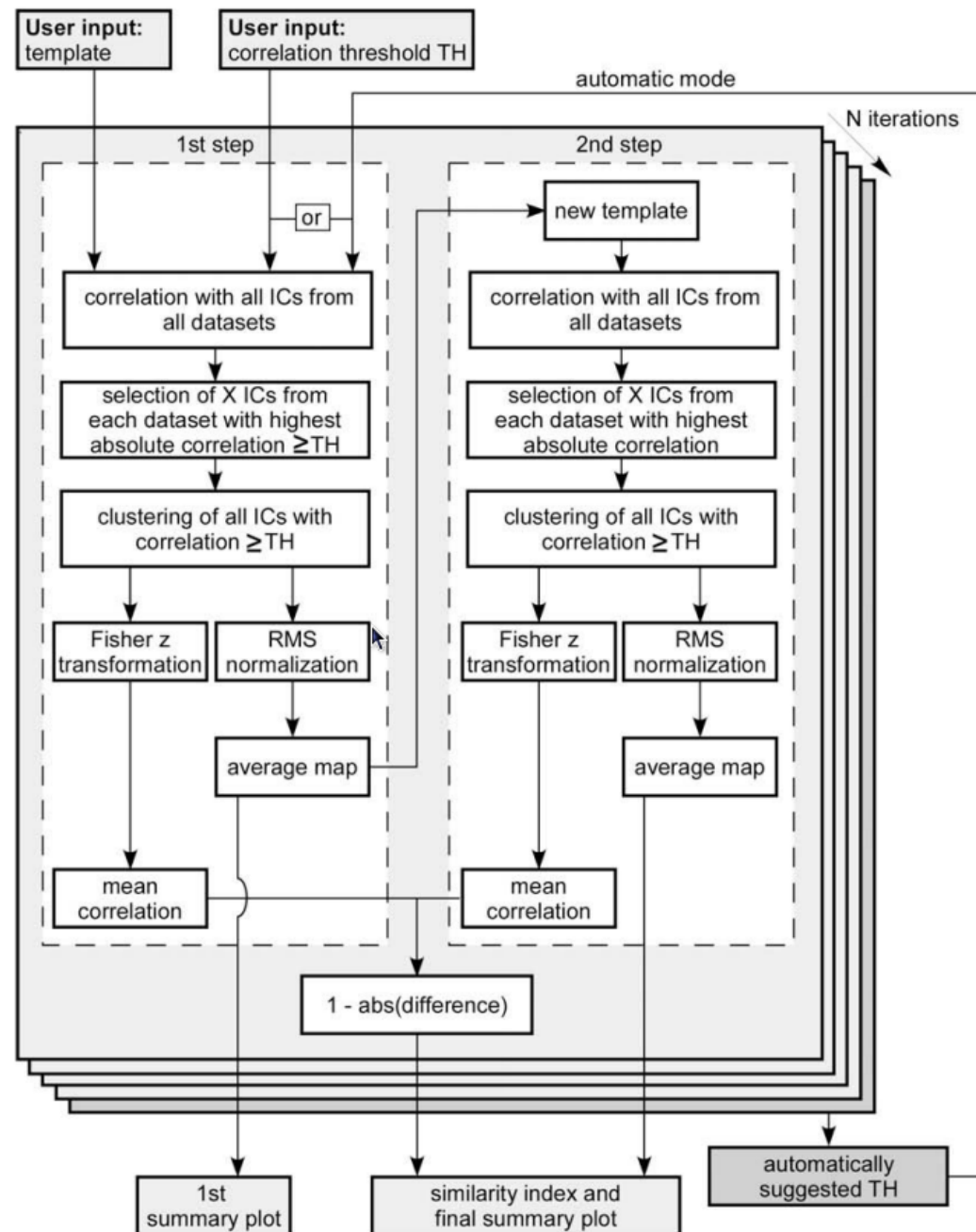




## ICA decomposition of multiple data sets from the same individuals

- Experimental protocol
  - Mind wandering experiment
  - 2 subjects
  - 11 x 30 min. sessions
  - 2 sessions per week
  - EEG from Biosemi 64 channels
  - $F_s=1024$  Hz

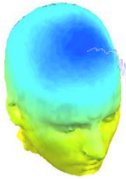




F. Campos Viola et al., "Semi-automatic identification of independent components representing EEG artifact," Clinical Neurophysiology 120, no. 5 (2009): 868–877.

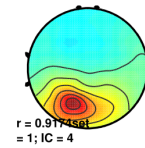
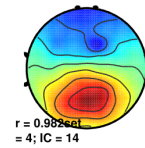
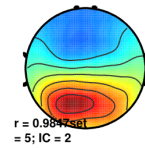
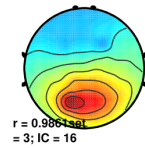
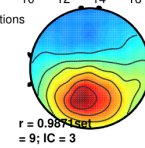
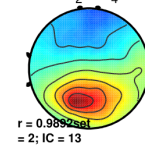
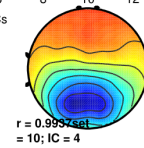
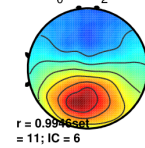
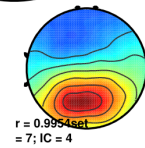
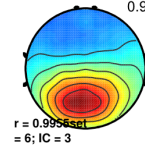
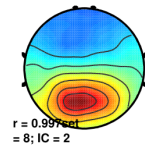
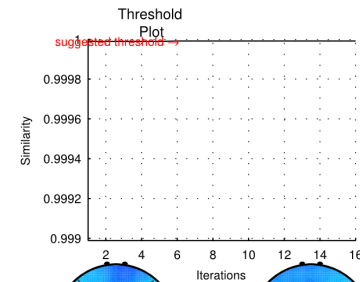
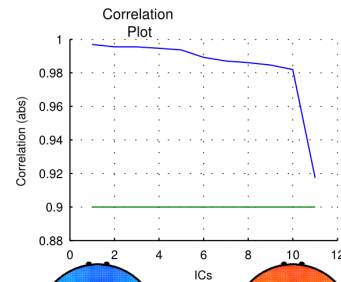
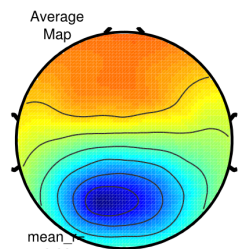


# Results (Cluster 1)

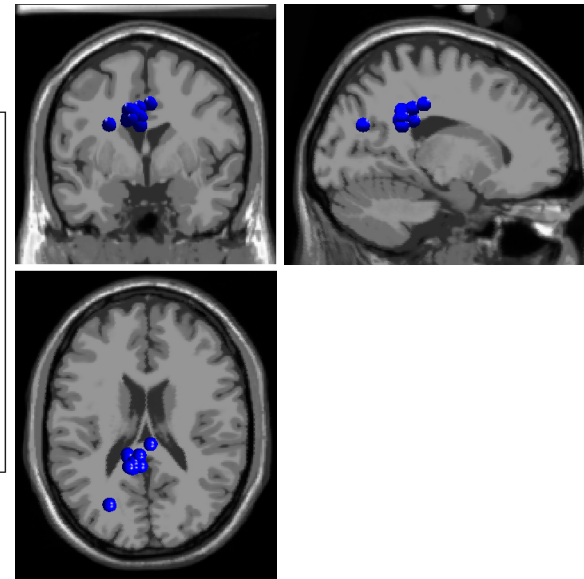
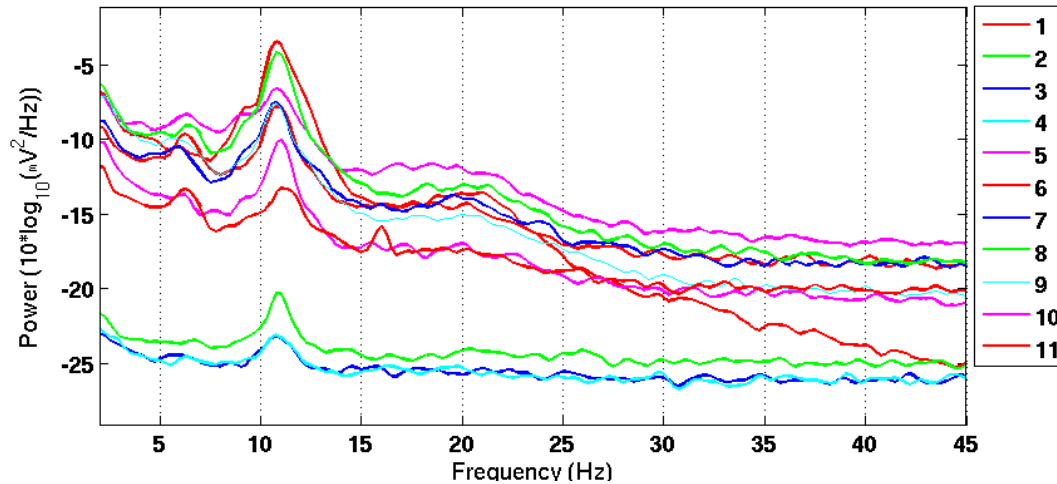


100 % Sessions contribute

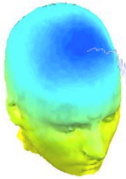
INFO:  
Template: CB Session 7 PREPROC:STEP 2; Set 7; IC 3;  
Number of datasets: 11  
Correlation threshold: 0.9 (green line)  
Max ICs from each dataset: 1  
Cluster: 11 ICs from 11 sets  
All datasets contribute.  
Similarity = 1.0000



Cls 3 Spectrum

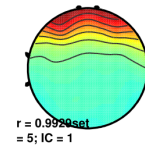
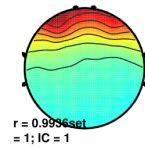
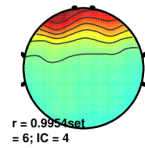
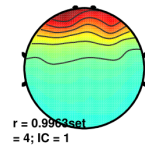
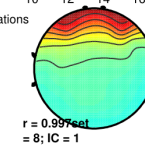
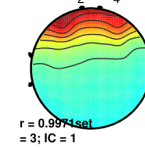
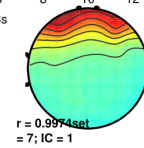
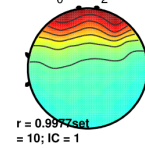
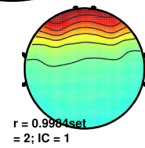
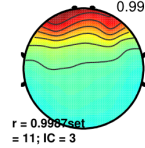
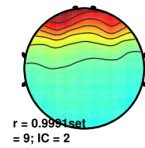
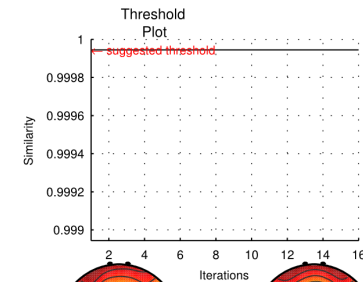
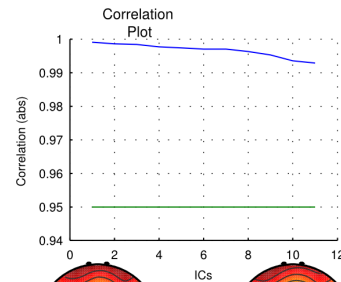
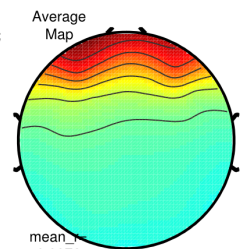


# Results (Cluster 2)

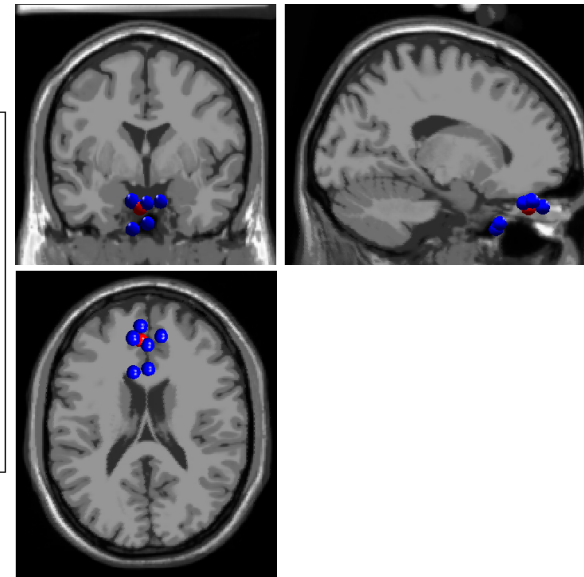
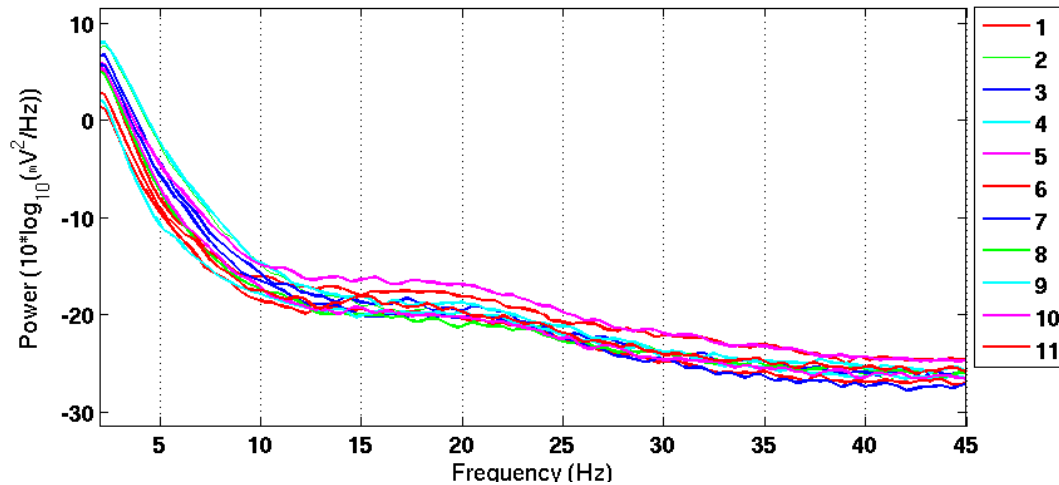


100 % Sessions contribute

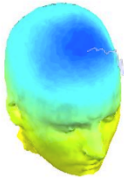
INFO:  
Template: CB Session 5 PREPROC:STEP 2; Set 5; IC 1;  
Number of datasets: 11  
Correlation threshold: 0.95 (green line)  
Max ICs from each dataset: 1  
Cluster: 11 ICs from 11 sets  
All datasets contribute.  
Similarity = 0.9999



Cls 4 Spectrum

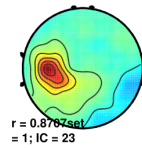
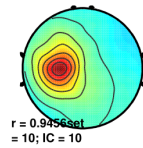
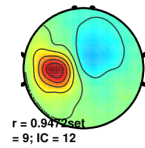
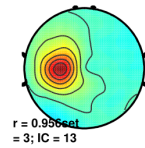
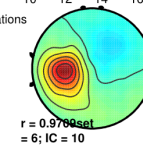
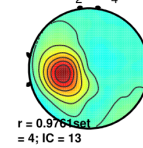
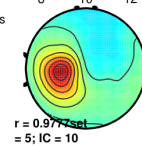
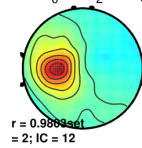
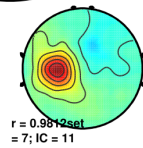
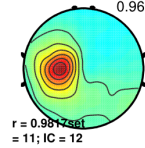
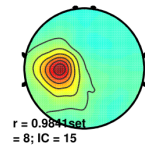
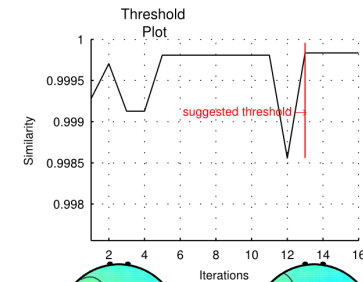
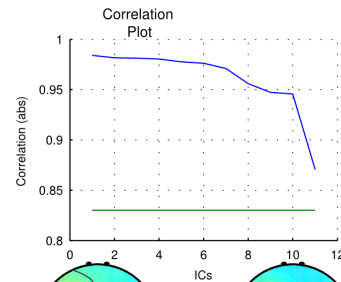
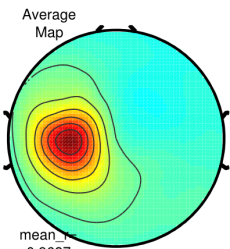


# Results (Cluster 8)

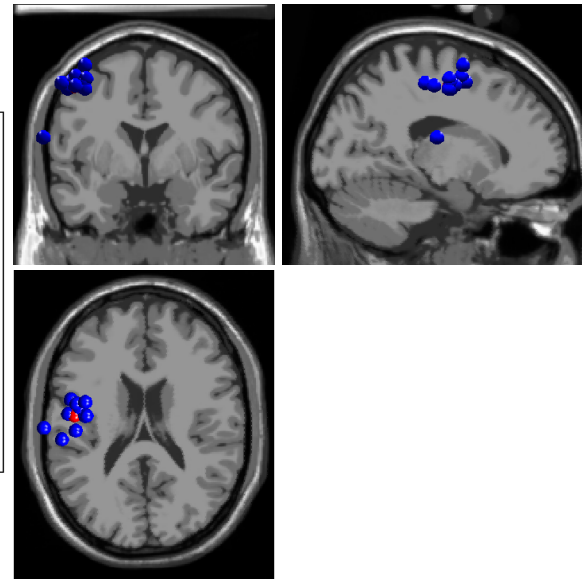
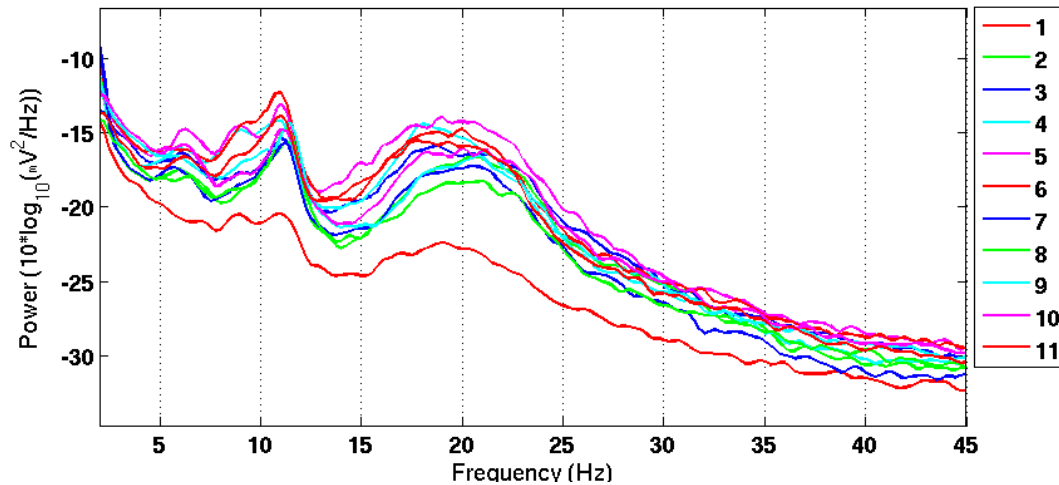


100 % Sessions contribute

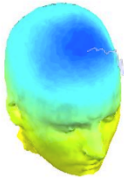
INFO:  
Template: CB Session 7 PREPROC:STEP 2; Set 7; IC 11;  
Number of datasets: 11  
Correlation threshold: 0.83 (green line)  
Max ICs from each dataset: 1  
Cluster: 11 ICs from 11 sets  
All datasets contribute.  
Similarity = 0.9998



Cls 8 Spectrum

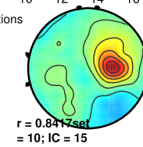
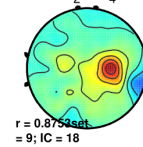
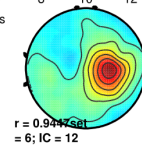
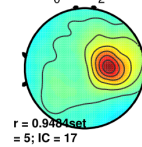
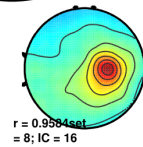
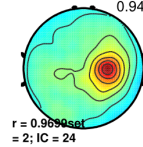
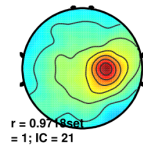
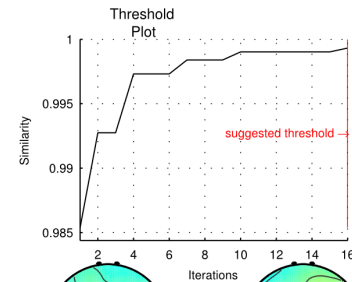
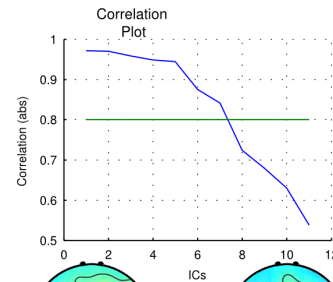
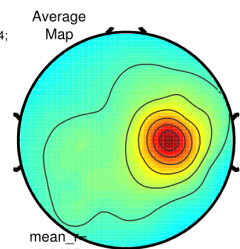


# Results (Cluster 13)

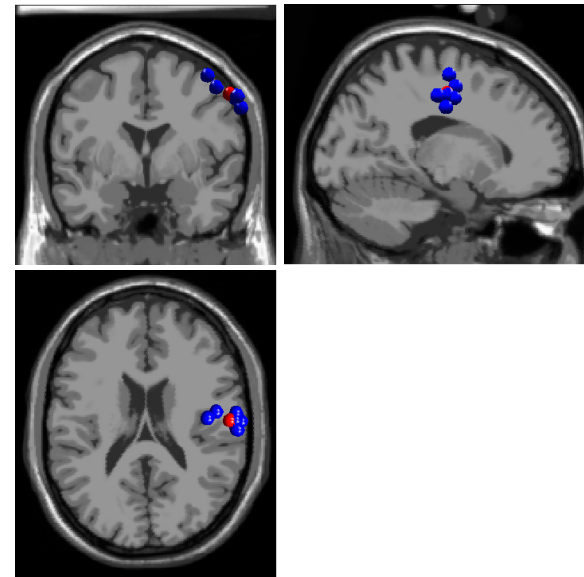
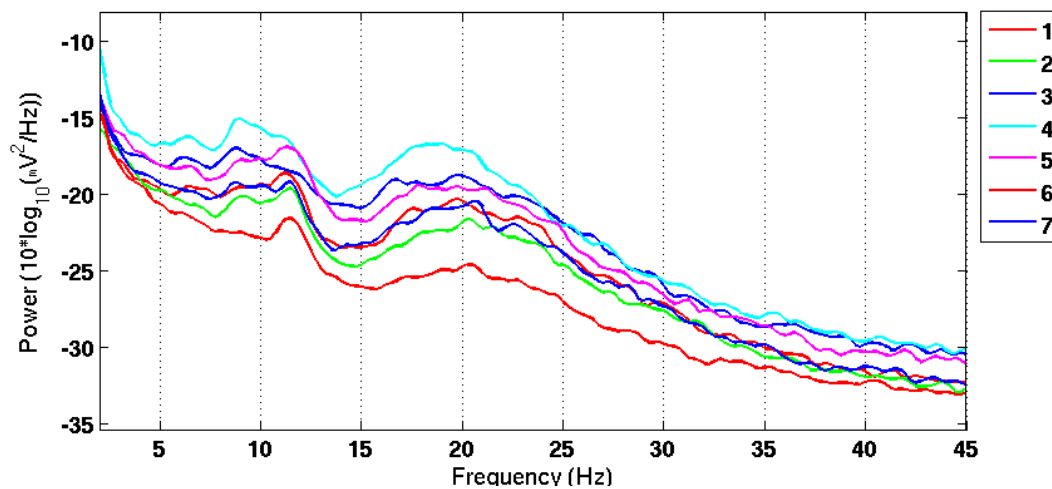


63.64% Sessions contribute

INFO:  
Template: CB Session 2 PREPROC:STEP 2; Set 2; IC 24;  
Number of datasets: 11  
Correlation threshold: 0.8 (green line)  
Max ICs from each dataset: 1  
Cluster: 7 ICs from 7 sets  
Sets not contributing:  
#3; #4; #7; #11;  
Similarity = 0.9993

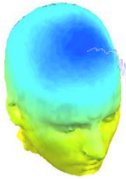


CIs 13 Spectrum



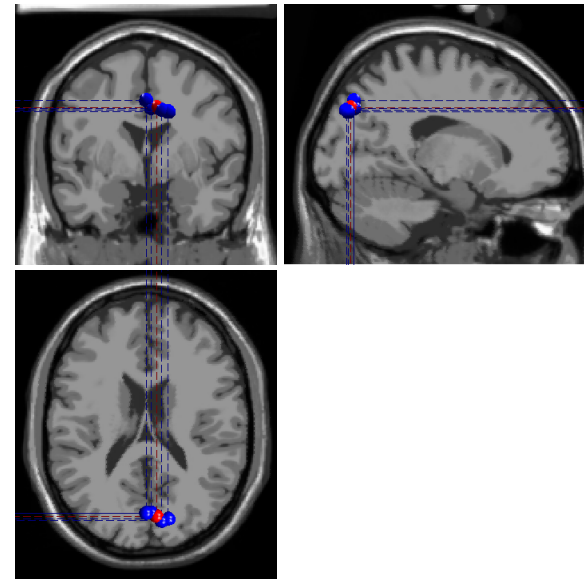
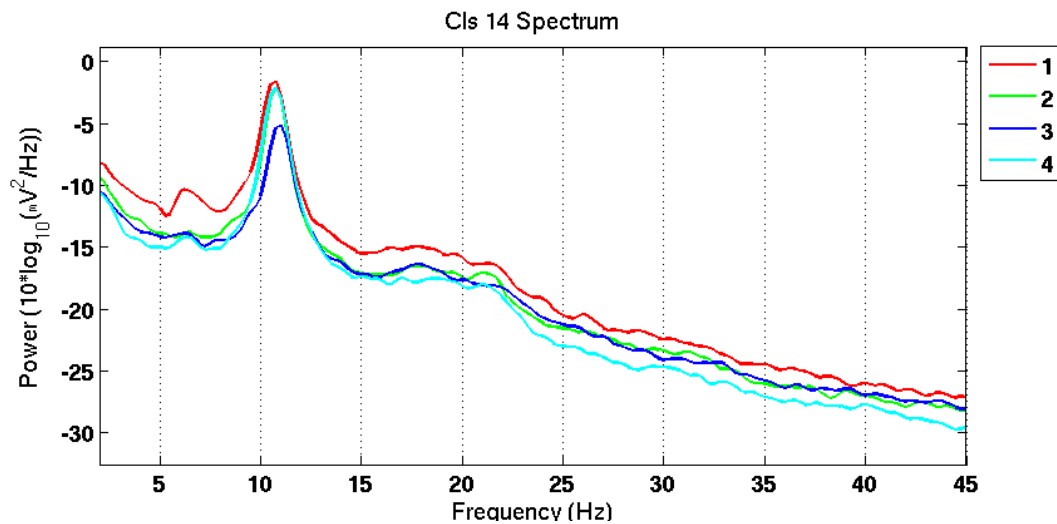
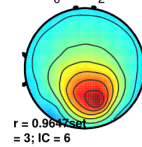
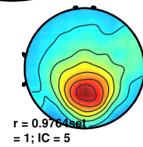
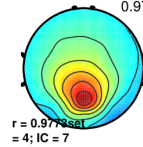
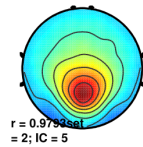
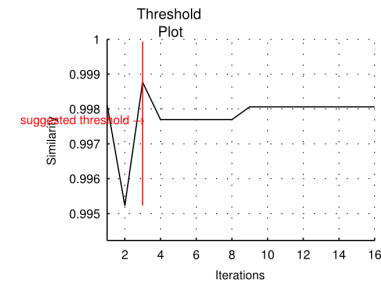
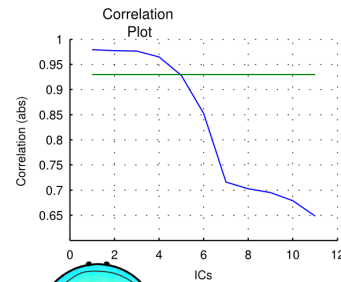
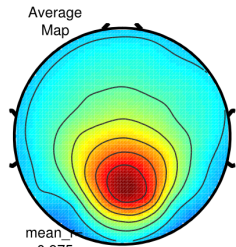


# Results (Cluster 14)

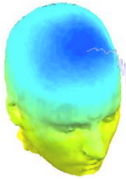


36.36% Sessions contribute

INFO:  
Template: CB Session 1 PREPROC:STEP 2; Set 1; IC 5;  
Number of datasets: 11  
Correlation threshold: 0.93 (green line)  
Max ICs from each dataset: 1  
Cluster: 4 ICs from 4 sets  
Sets not contributing:  
#5; #6; #7; #8; #9; #10#11;  
Similarity = 0.9988



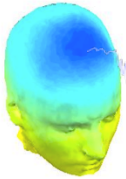
# Inter iteration Cluster Consistency



Iterations

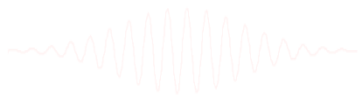
Clusters

	1	2	3	4	5	6	7	8	9	10	Mean
3	100	100	100	100	100	100	100	100	100	100	<b>100</b>
4	100	100	100	100	100	100	90	100	100	100	<b>99</b>
5	90	40	10	90	90	60	100	10	60	90	<b>64</b>
6	60	0	100	60	100	90	60	60	90	60	<b>68</b>
7	90	100	90	90	60	90	90	100	90	90	<b>89</b>
8	80	80	60	80	40	80	80	80	80	100	<b>76</b>
9	60	90	50	60	80	60	0	10	60	50	<b>52</b>
10	40	90	10	40	0	50	50	0	50	60	<b>39</b>
11	60	20	0	0	10	60	10	90	60	60	<b>37</b>
12	100	50	50	100	50	100	100	50	100	50	<b>75</b>
13	50	10	20	50	90	50	50	10	50	20	<b>40</b>
14	20	10	10	20	20	30	20	20	30	30	<b>21</b>

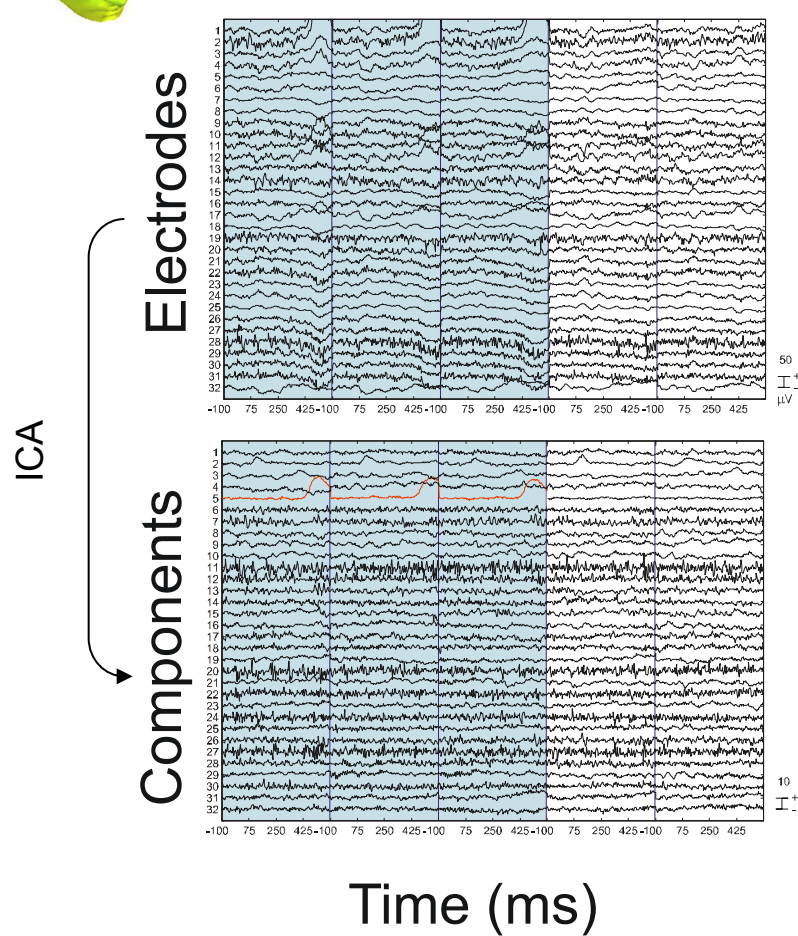
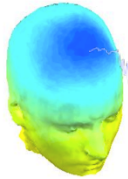


## Outline

- ICA clusters and reliability within subjects
- **ICA clusters and reliability across subjects**
- Clustering in EEGLAB theory & Practice

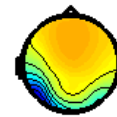
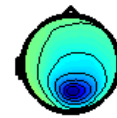
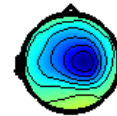
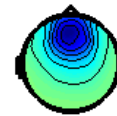


# Localization

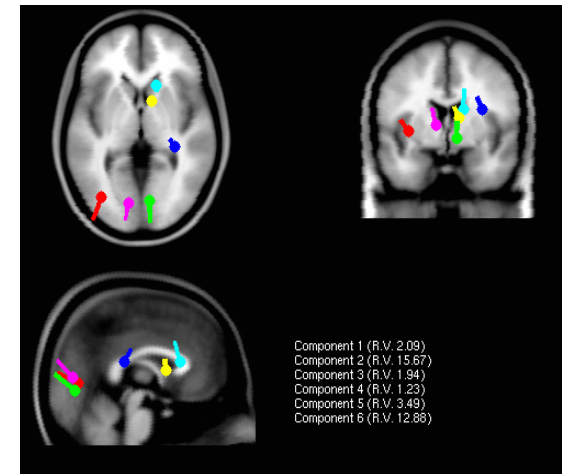


ICA component

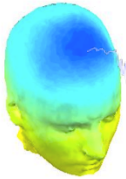
scalp maps



Localization



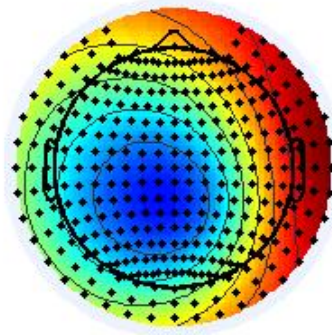




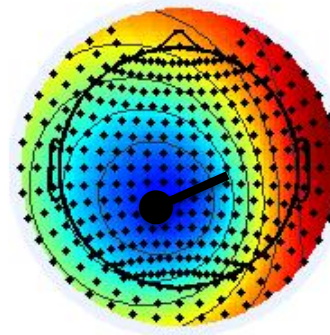
# Computing residual variance (%)



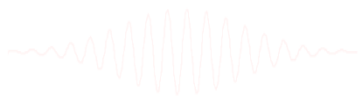
Actual

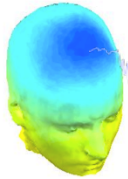


Dipole projection



$$r = \frac{\sum (x_i - \tilde{x}_i)^2}{\sum x_i^2}$$





# Validation of the ICA algorithm for EEG



## Data

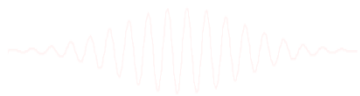
- 13 subjects performing a memory task
- 71 electrodes including EOGs
- more than 300,000 data points/subject

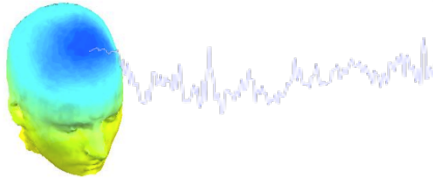
## Decomposition

- 23 ICA algorithms plus PCA and Promax

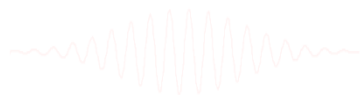
## Analysis

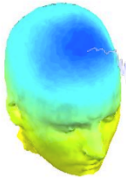
- Localization of all components with a single dipole (4-shell spherical model)



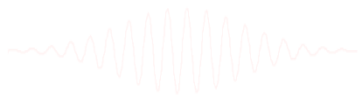
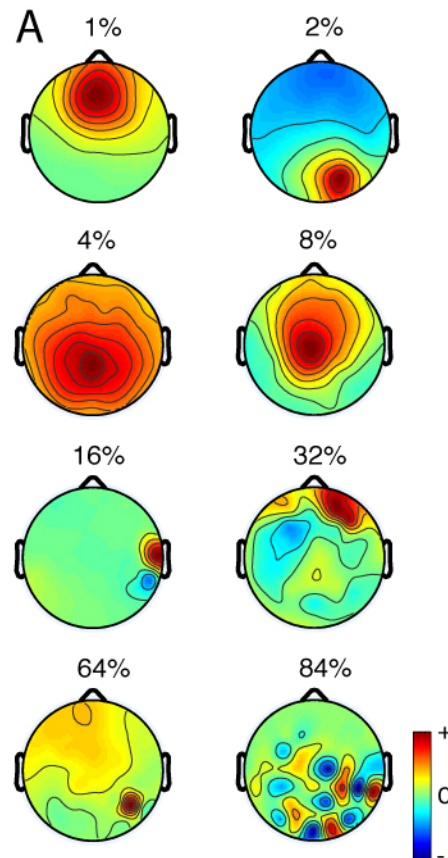


Algorithm (Matlab func.)	D%	LL	Origin
Extended Infomax (runica)	29.9	178	EEGLAB 4.515
Pearson	29.1	169	ICACentral (6)
Infomax (runica)	28.2	160	EEGLAB 4.515
ERICA	26.9	184	ICALAB 1.5.2
SONS	25.4	183	ICALAB 1.5.2
SHIBBS	23.7	169	ICACentral (5)
FastICA*	23.5	169	ICACentral (2)
JADE (jader)	23.4	169	EEGLAB 4.515
TICA	23.4	169	ICALAB 1.5.2
JADE optimized (jade_op)	21.4	169	ICALAB 1.5.2
JADE w/ time delay (jade_td)	20.2	169	ICALAB 1.5.2
eeA	19.0	305	ICACentral (8)
Infomax (icaML) †	18.8	212	ICA DTU Tbox
FOBI	18.6	169	ICALAB 1.5.2
SOBIRO (acsobiro)	17.9	167	EEGLAB 4.515
EVD 24	17.7	169	ICALAB 1.5.2
EVD	17.0	169	ICALAB 1.5.2
SOBI	16.1	583	EEGLAB 4.515
icaMS†	10.6	169	ICA DTU Tbox
AMUSE	8.5	169	ICALAB 1.5.2
PCA	3.1	583	EEGLAB 4.515
Promax	33.7	467	EEGLAB 4.515
Whitening/Sphering	57.6	164	EEGLAB 4.515

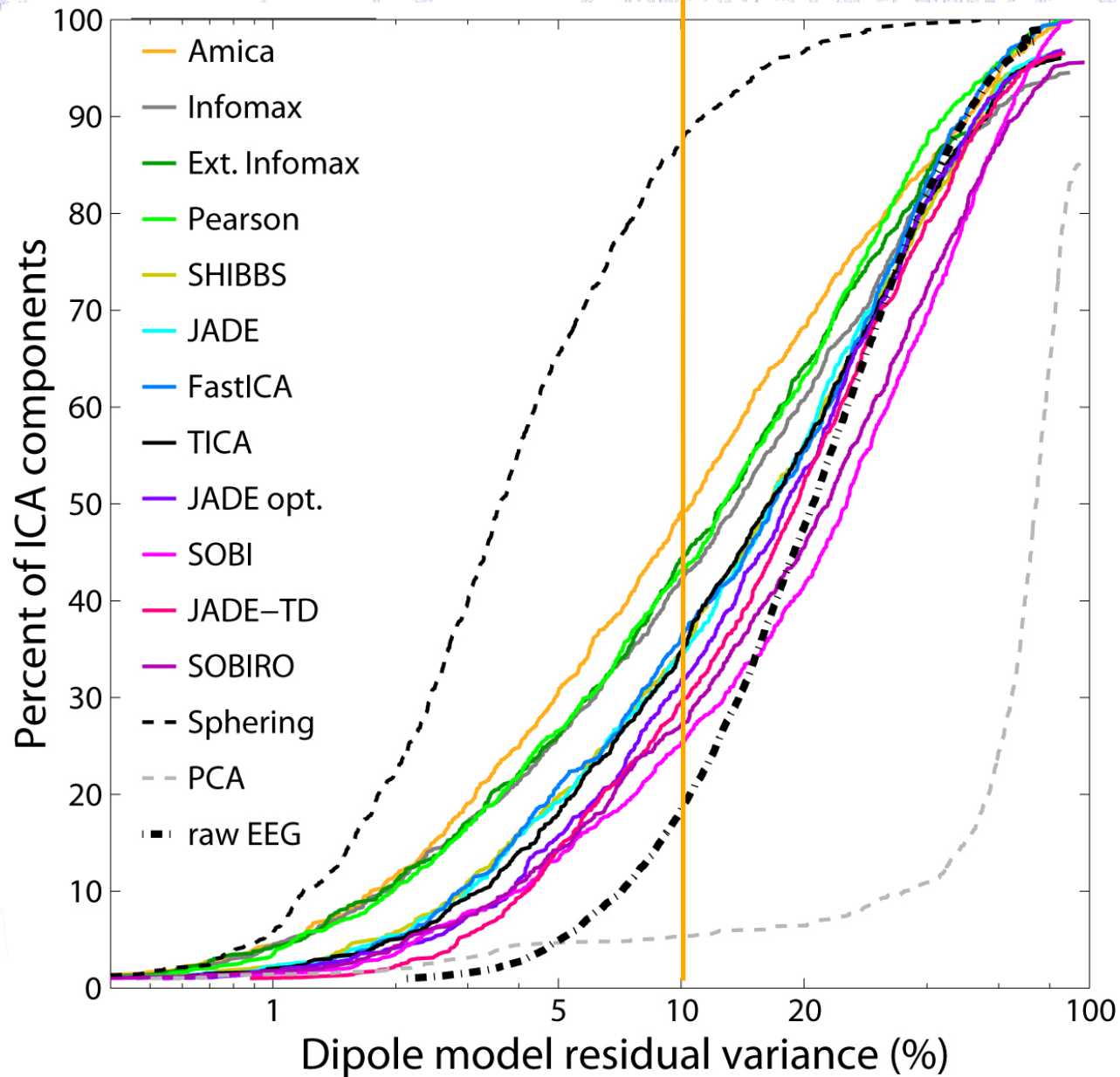
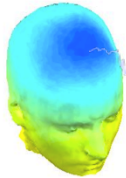




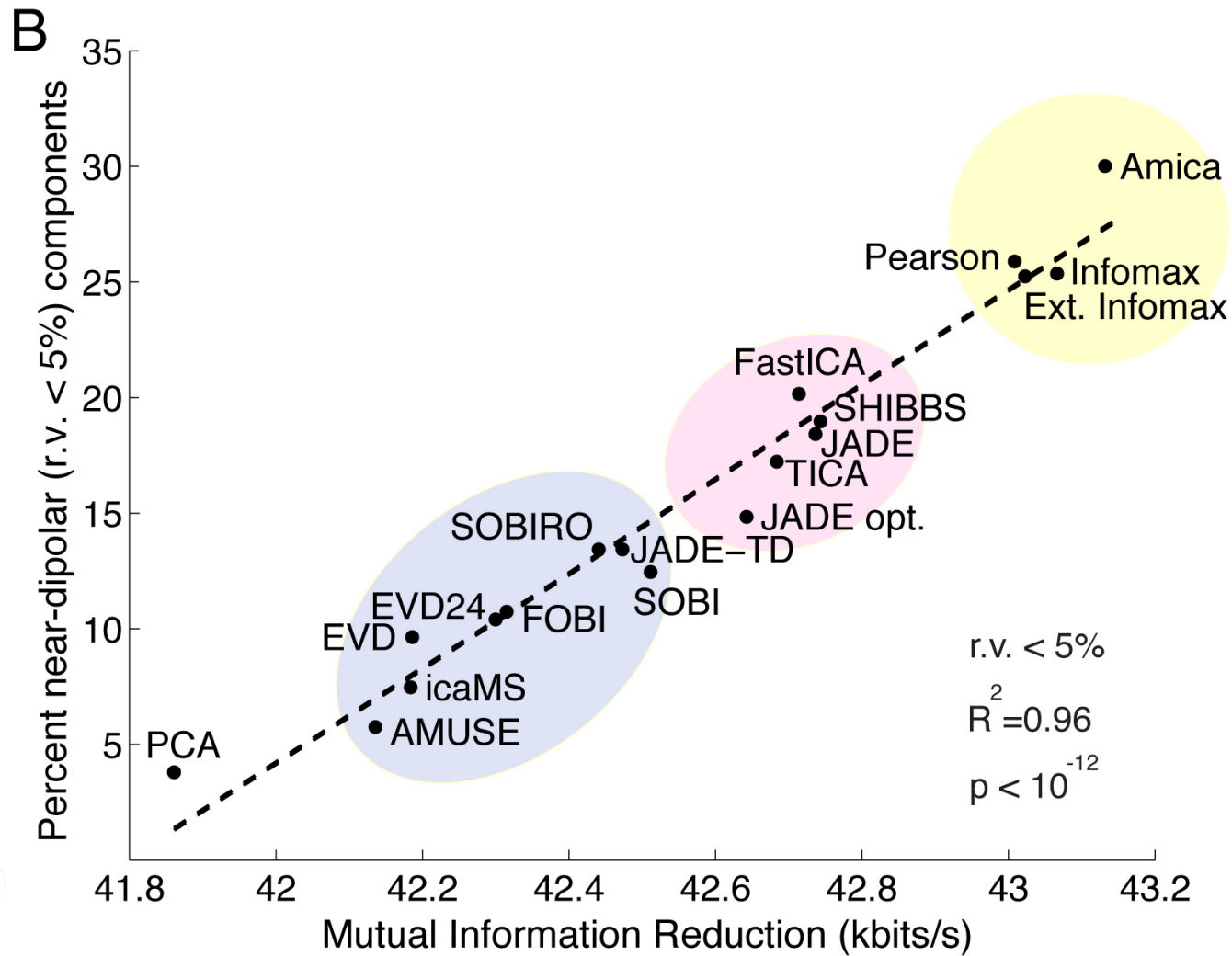
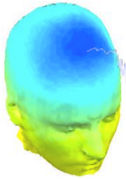
# Component examples

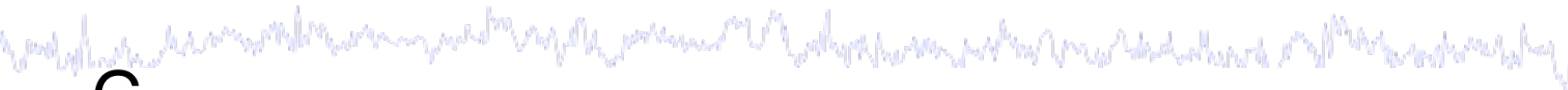
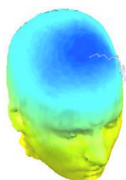


# Cumulative number of component below residual variance threshold

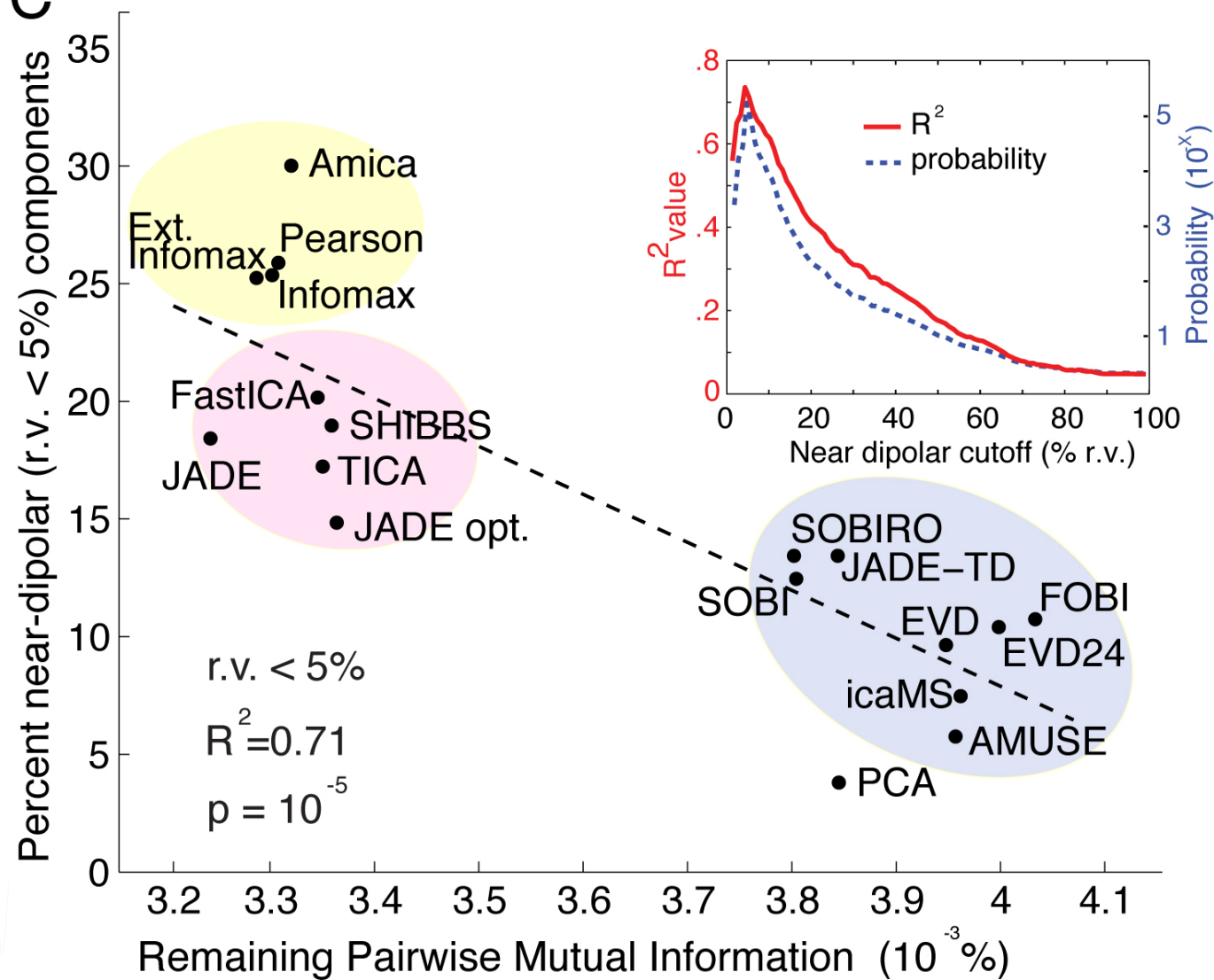


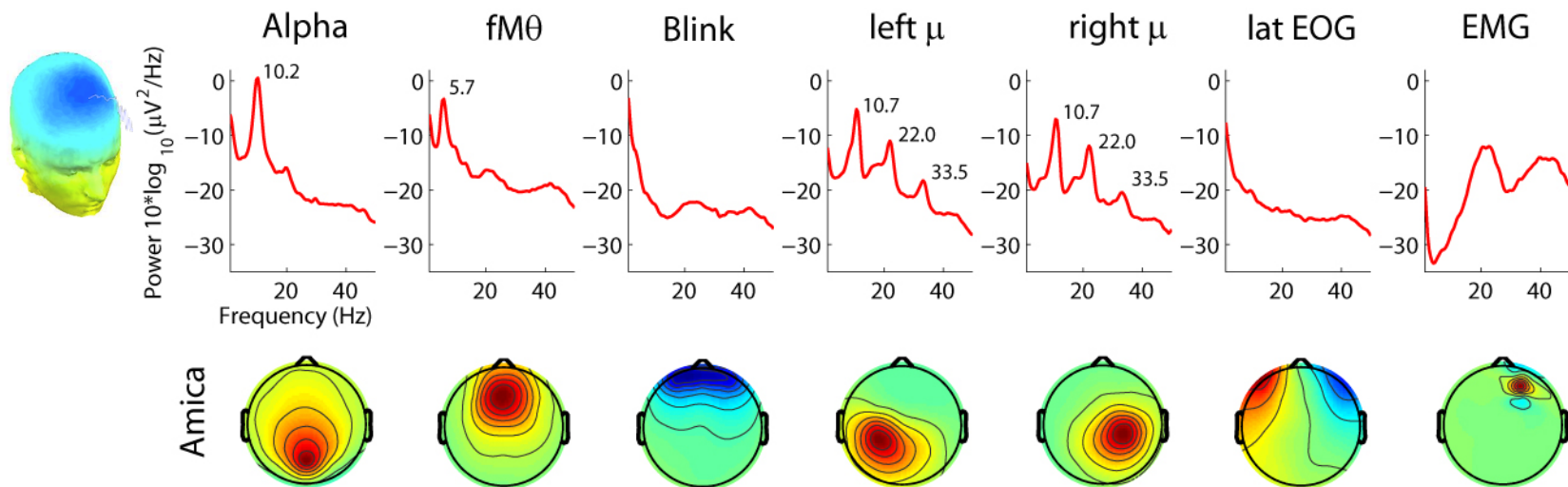
# More independence -> more biological components



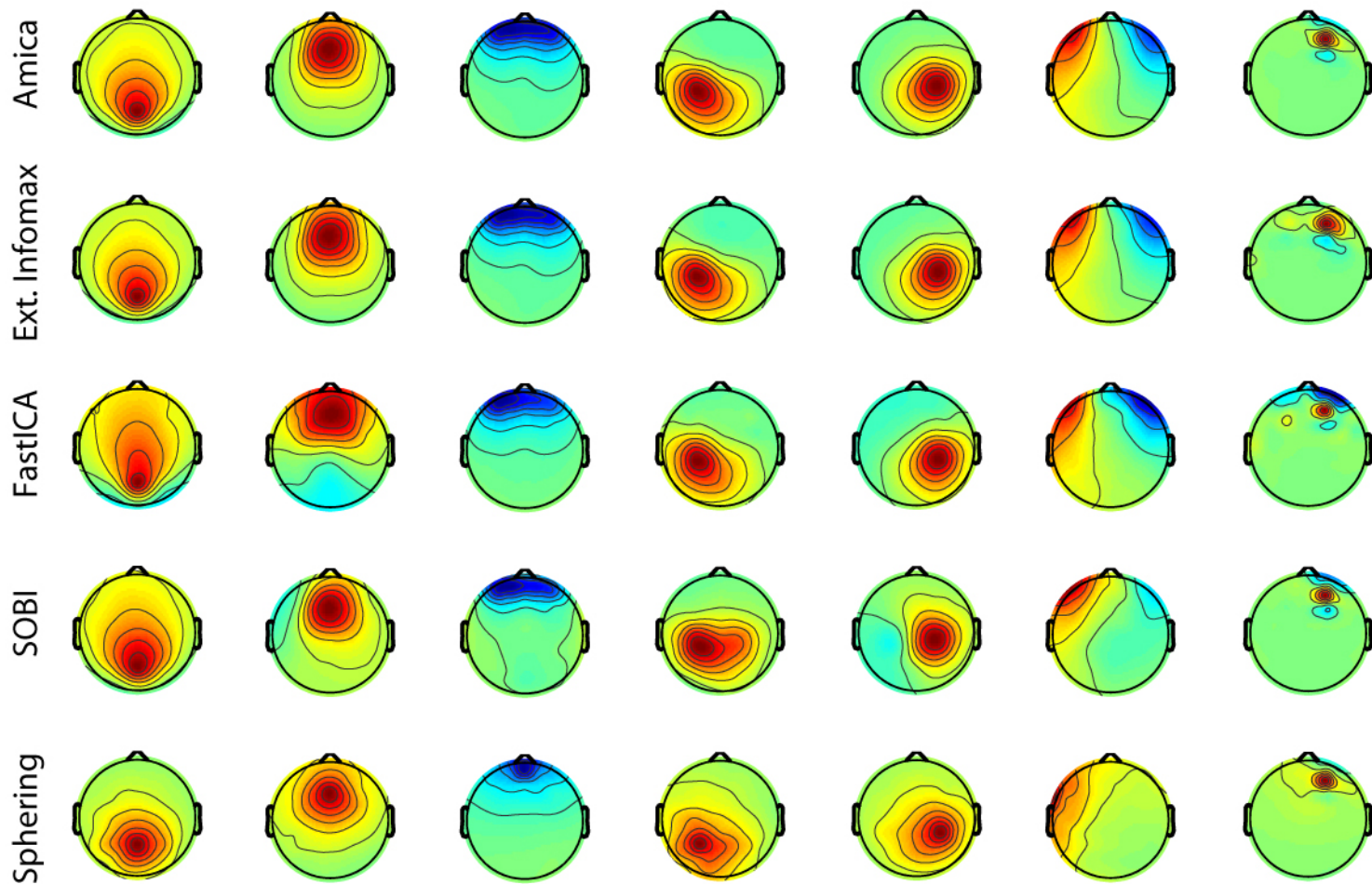
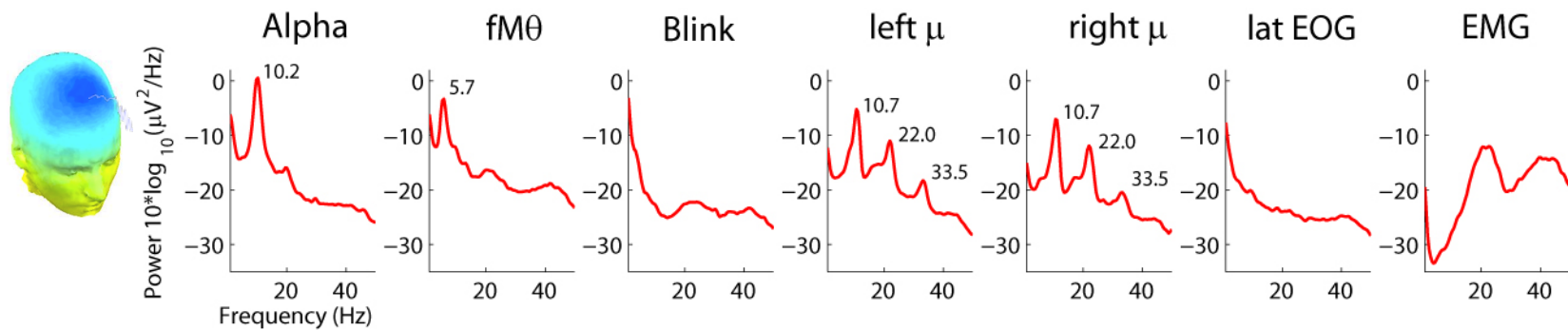


C

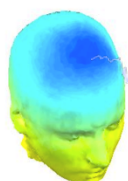






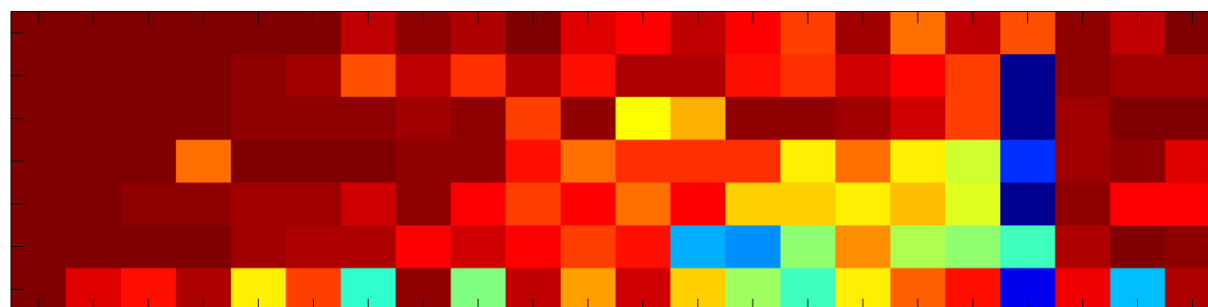


# Correlations between decompositions



Scalp map corr.

Alpha  
fM0  
Blink  
left  $\mu$   
right  $\mu$   
lat. EOG  
EMG

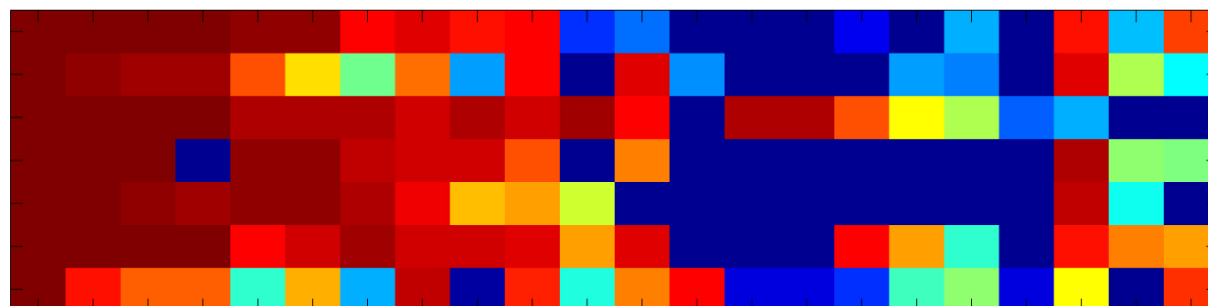


(r)  
1  
0.9  
0.8  
0.7  
0.6  
0.5

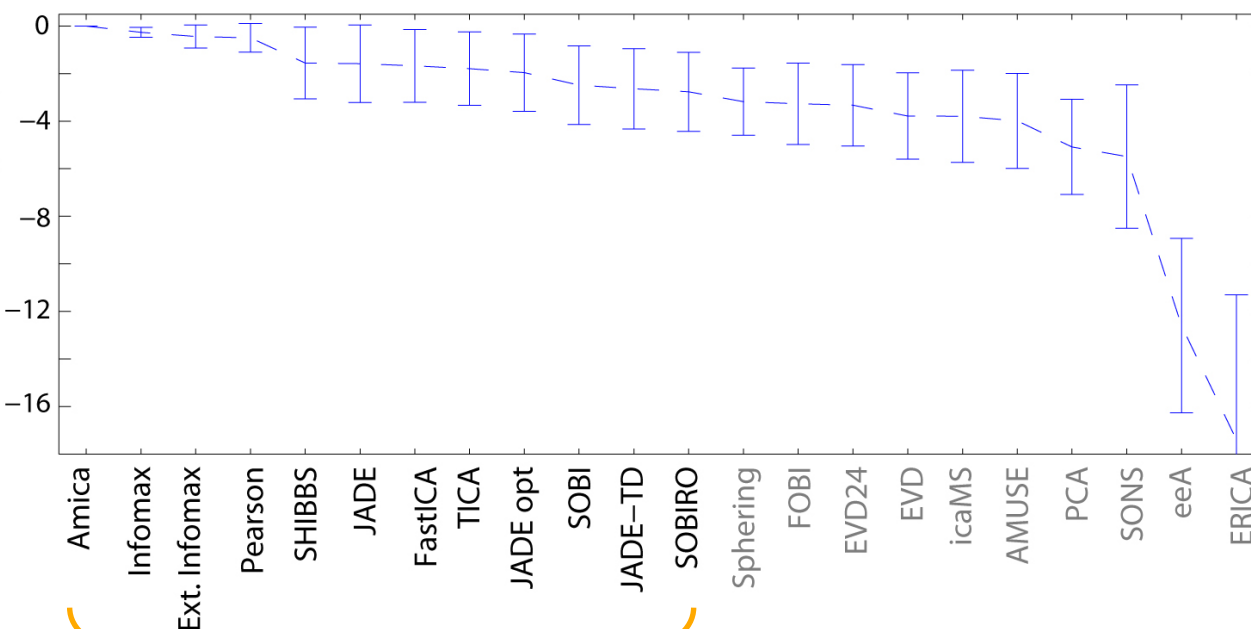


Raw data corr.

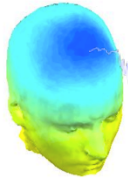
Alpha  
fM0  
Blink  
left  $\mu$   
right  $\mu$   
lat. EOG  
EMG



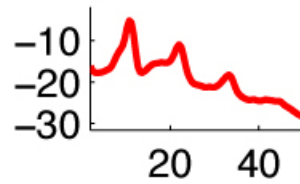
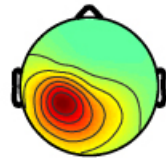
Information reduction  
difference (bits)



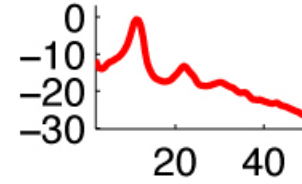
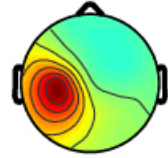
# Left $\mu$ cluster



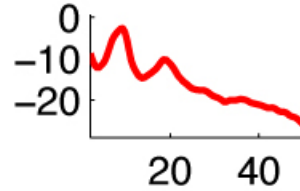
S2 IC47



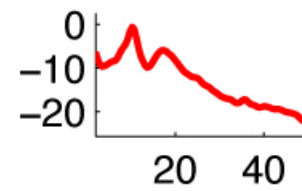
S3 IC47



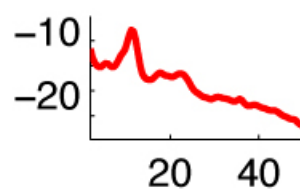
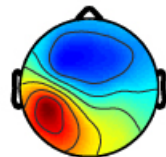
S4 IC37



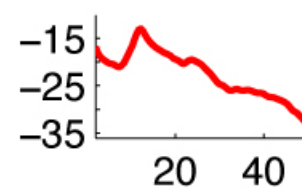
S5 IC48



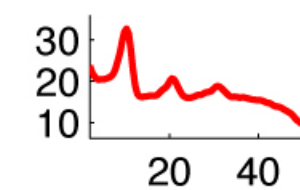
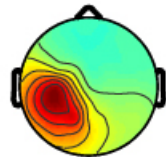
S6 IC46



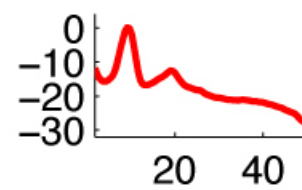
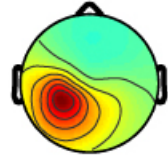
S7 IC35



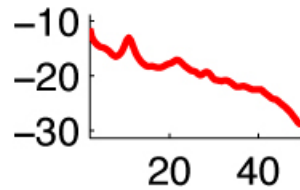
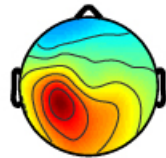
S9 IC7



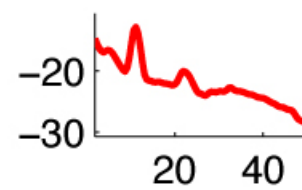
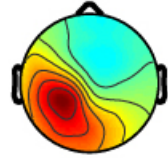
S11 IC45



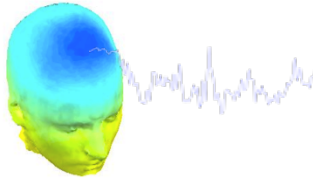
S12 IC45



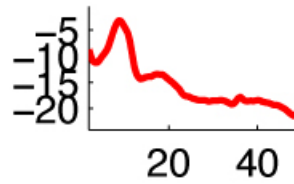
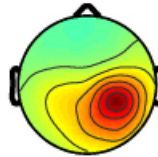
S14 IC45



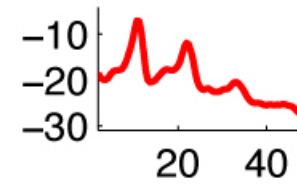
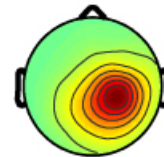
# Right $\mu$ cluster



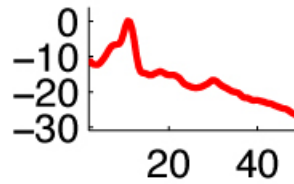
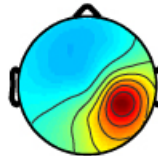
S1 IC51



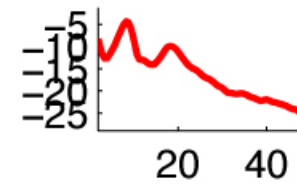
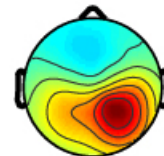
S2 IC41



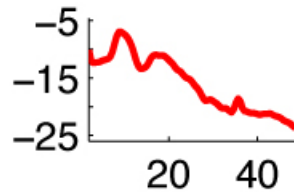
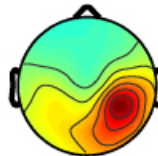
S3 IC41



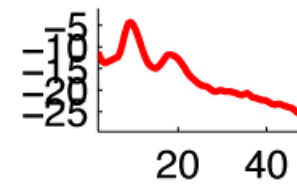
S4 IC50



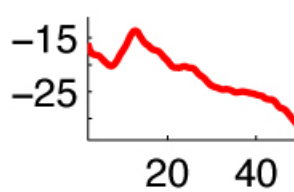
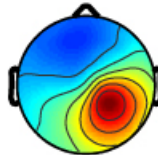
S5 IC51



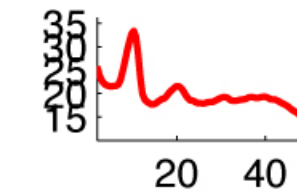
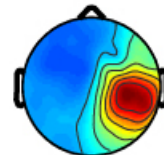
S6 IC6<sup>0</sup>



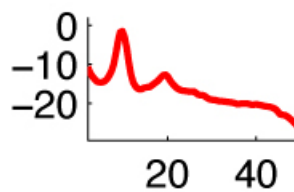
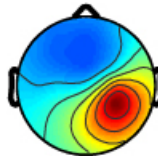
S7 IC48



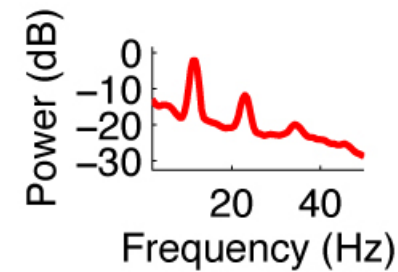
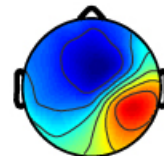
S9 IC39



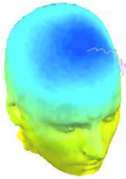
S11 IC49



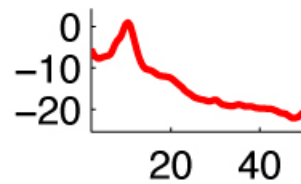
S14 IC49



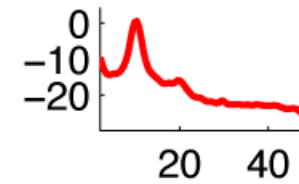
# Occipital $\alpha$ cluster



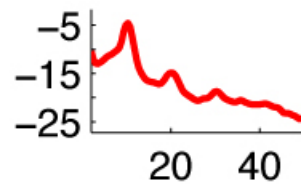
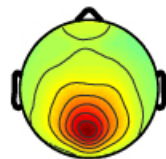
S1 IC67



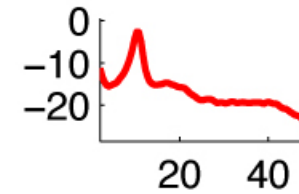
S2 IC67



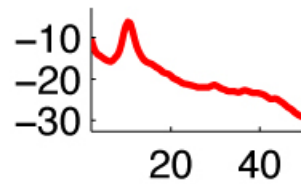
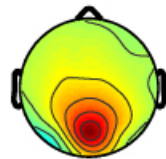
S3 IC51



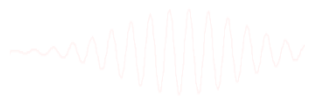
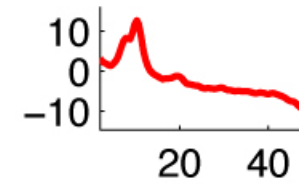
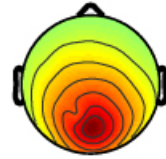
S11 IC65



S12 IC3<sup>8</sup>

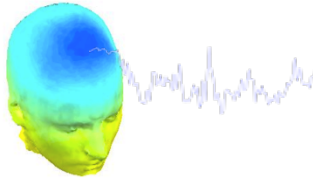


S13 IC65

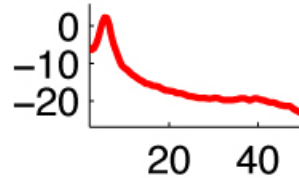




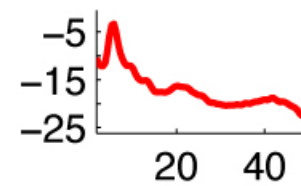
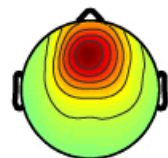
# Frontal Midline $\theta$ cluster



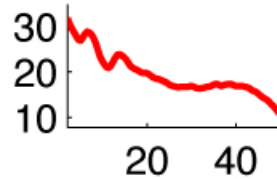
S1 IC63



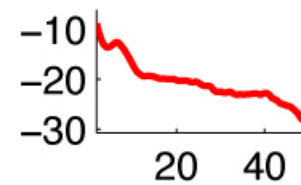
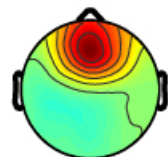
S2 IC18



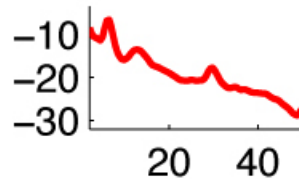
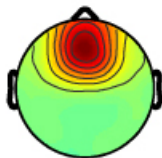
S9 IC16



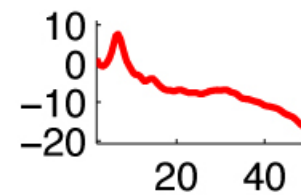
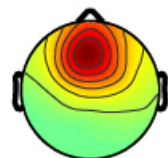
S11 IC16



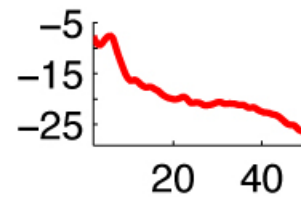
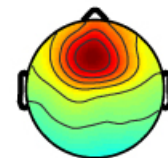
S12 IC15

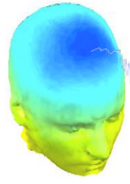


S13 IC15



S14 IC16





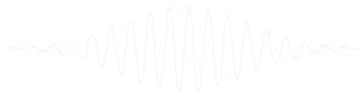
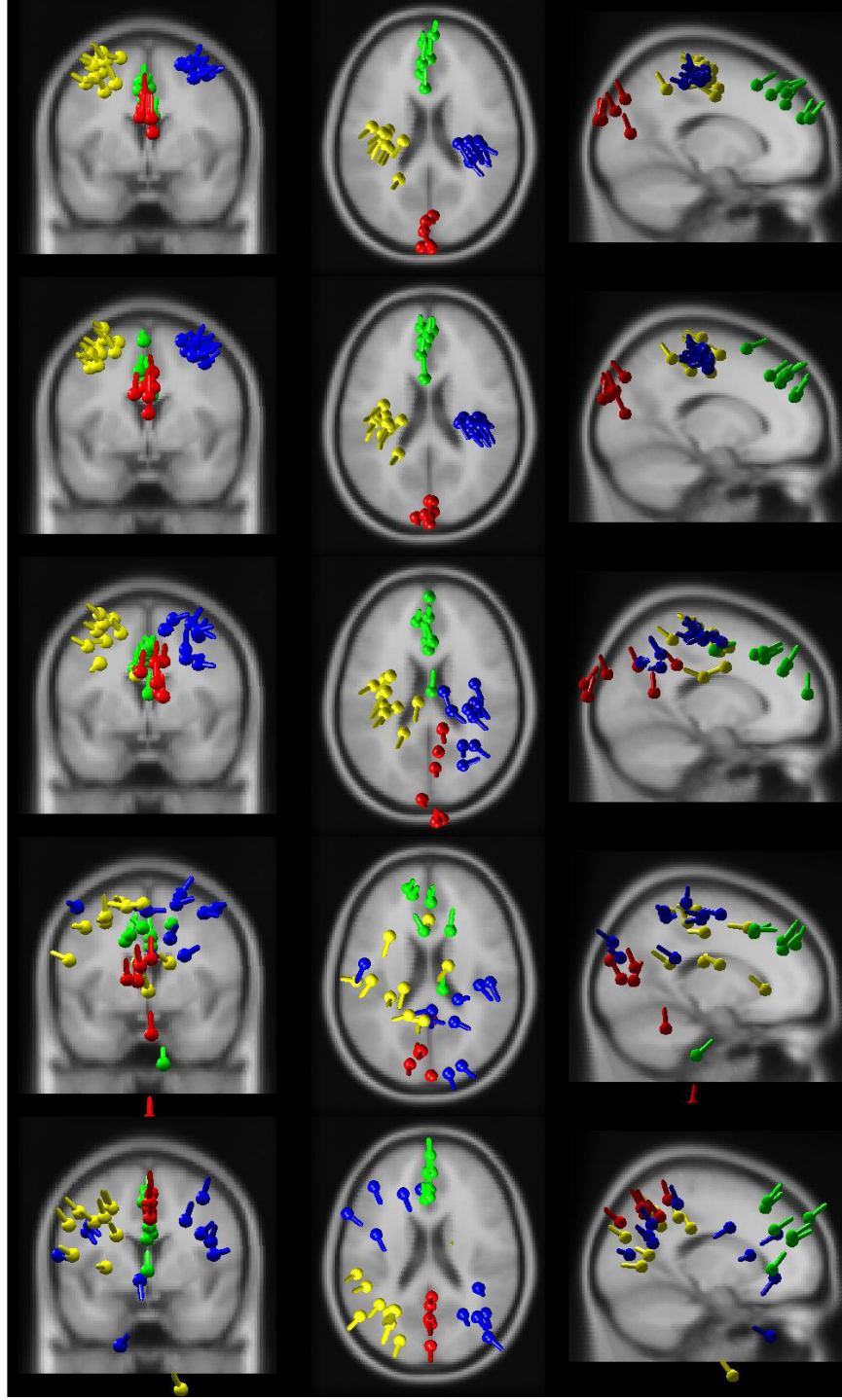
AMICA

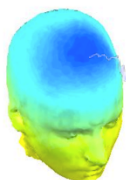
Ext. Infomax

FASTICA

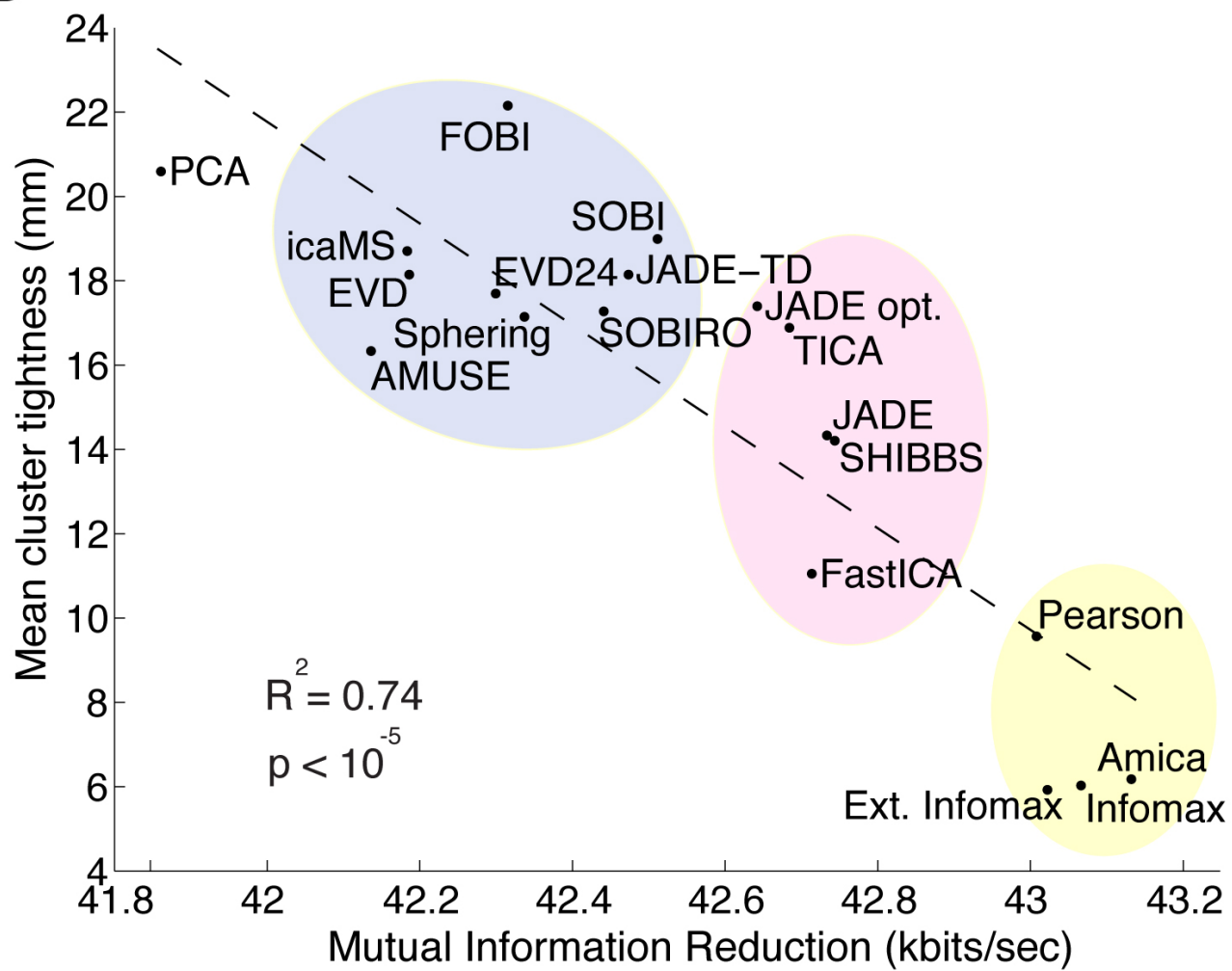
SOBI

Sphering

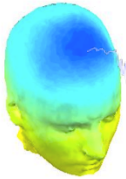




D

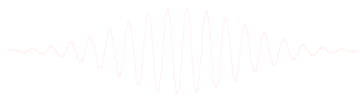




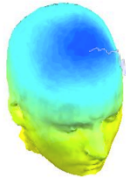


## Outline

- ICA clusters and reliability within subjects
- ICA clusters and reliability across subjects
- **Clustering in EEGLAB theory & Practice**



# Edit dataset info



pop\_study(): Pre-select components

Enter maximum residual (topo map - dipole proj.) var. (in %)  
NOTE: This will delete any existing component clusters!

15

☒ Keep only in-brain dipoles.

Cancel Help Ok

Create a new STUDY set -- pop\_study()

**Edit STUDY set information - remember to save changes**

STUDY set name: Sternberg

STUDY set task name: Sternberg

STUDY set notes:

	dataset filename	browse	subject	session	condition	group	Select by r.v.	
1	C:\Users\julie\Documents\Wor	...	S01		memorize		Comp.: 3 5 ...	Clear
2	C:\Users\julie\Documents\Wor	...	S01		ignore		Comp.: 3 5 ...	Clear
3	C:\Users\julie\Documents\Wor	...	S01		probe		Comp.: 3 5 ...	Clear
4	C:\Users\julie\Documents\Wor	...	S02		memorize		Comp.: 5 6 ...	Clear
5	C:\Users\julie\Documents\Wor	...	S02		ignore		Comp.: 5 6 ...	Clear
6	C:\Users\julie\Documents\Wor	...	S02		probe		Comp.: 5 6 ...	Clear
7	C:\Users\julie\Documents\Wor	...	S03		memorize		Comp.: 6 7 ...	Clear
8	C:\Users\julie\Documents\Wor	...	S03		ignore		Comp.: 6 7 ...	Clear
9	C:\Users\julie\Documents\Wor	...	S03		probe		Comp.: 6 7 ...	Clear
10	C:\Users\julie\Documents\Wor	...	S04		memorize		Comp.: 1 2 ...	Clear

Important note: Removed datasets will not be saved before being deleted from EEGLAB memory

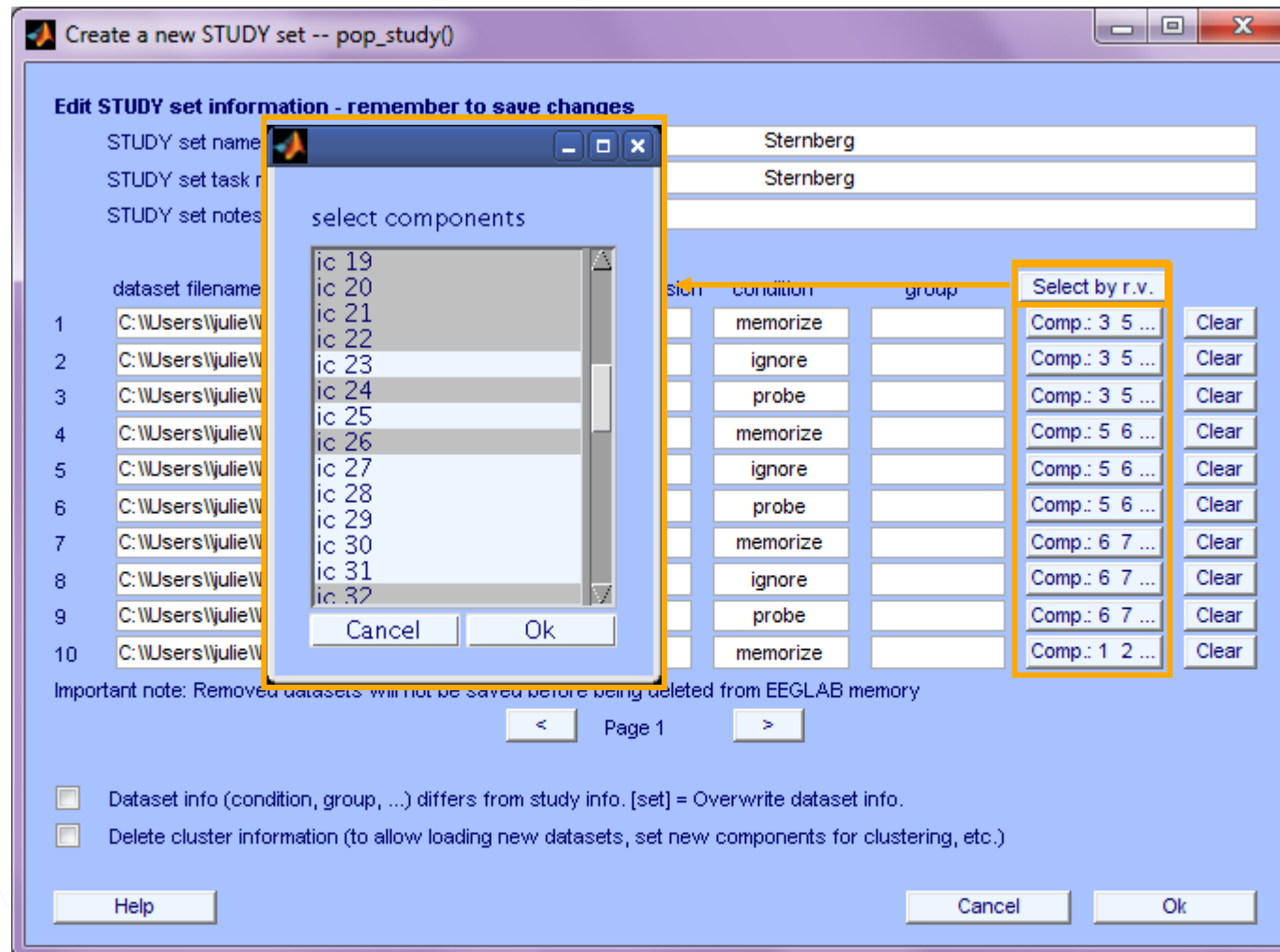
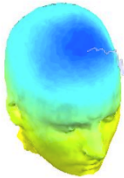
< Page 1 >

☐ Dataset info (condition, group, ...) differs from study info. [set] = Overwrite dataset info.

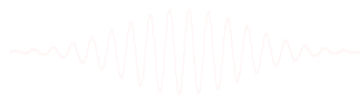
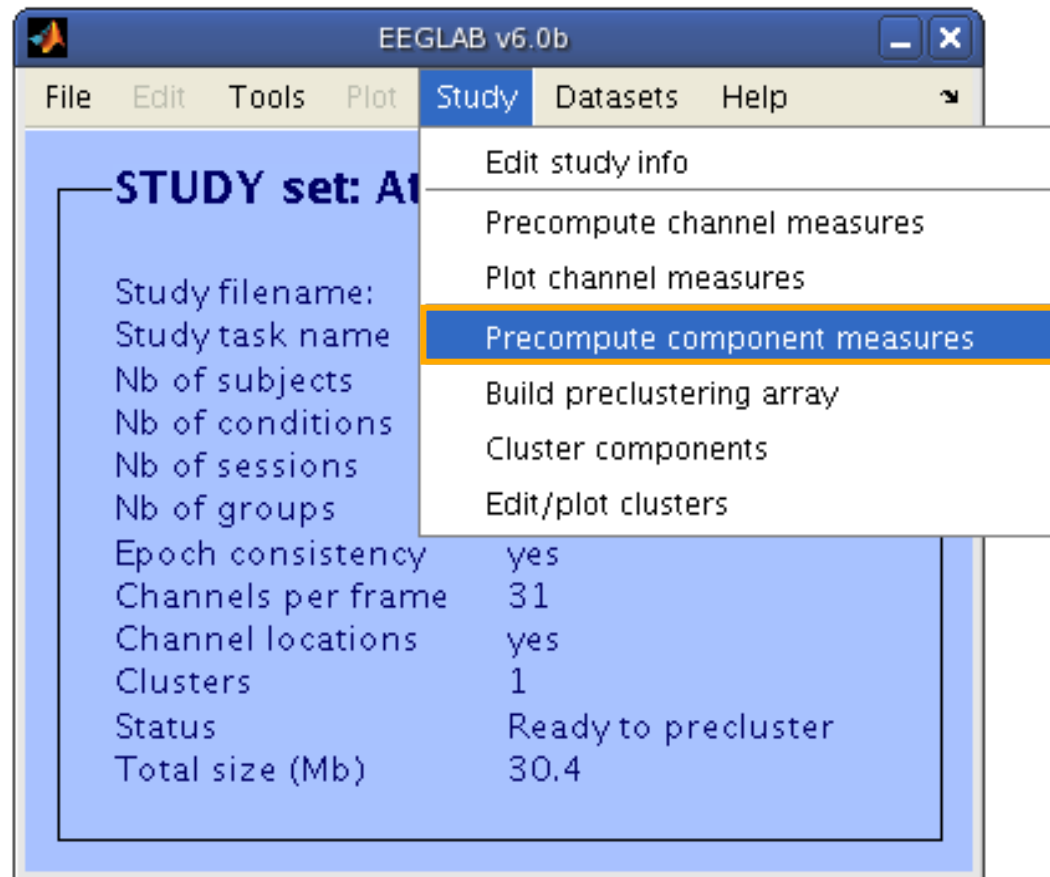
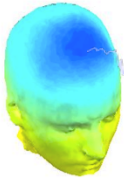
☐ Delete cluster information (to allow loading new datasets, set new components for clustering, etc.)

Help Cancel Ok

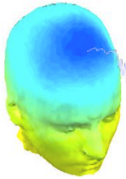
# ICs to cluster



# Precompute data measures



# Precompute data measures



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

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

**Pre-compute component measures for STUDY 'Sternberg'**

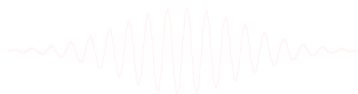
☒ Compute ERP/spectrum/ERSP only for components selected by RV (set) or for all components (unset)

**List of measures to precompute**

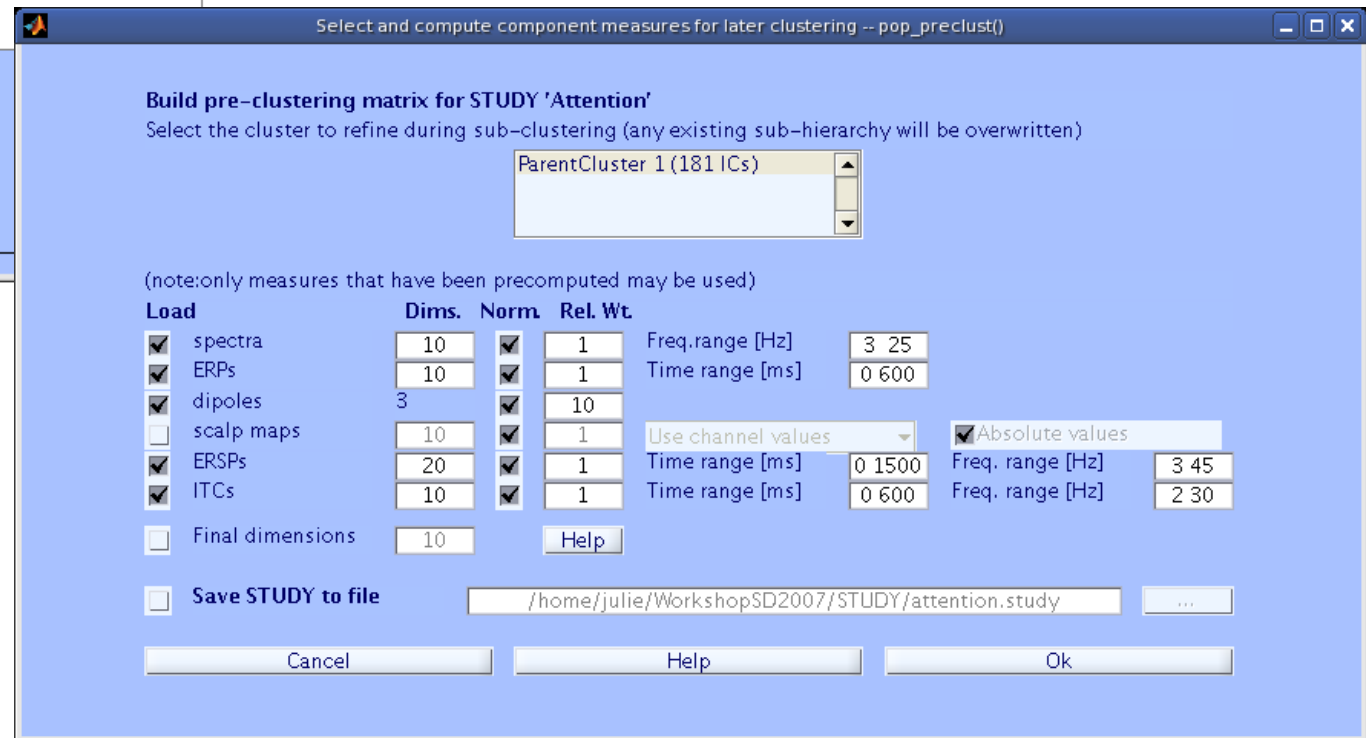
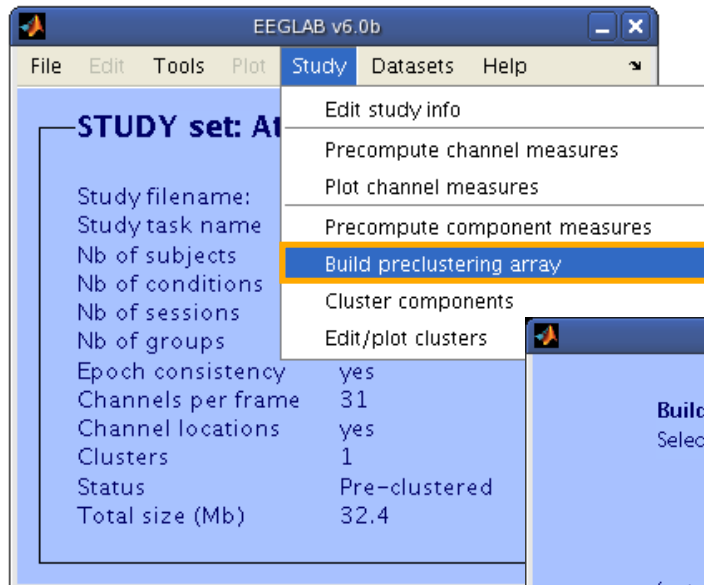
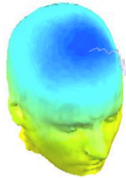
<input checked="" type="checkbox"/> ERPs	Baseline ([min max] in ms)	<input type="text" value="[-200 0]"/>	
<input checked="" type="checkbox"/> Power spectrum	Spectopo parameters	<input type="text" value=""/>	<input type="button" value="Test"/>
<input checked="" type="checkbox"/> ERSPs	Time/freq. parameters	<input type="text" value="'cycles', [3 0.5], 'nfreqs', 100"/>	<input type="button" value="Test"/>
<input checked="" type="checkbox"/> ITCs			
<input checked="" type="checkbox"/> Scalp maps			

☐ Recompute even if present on disk

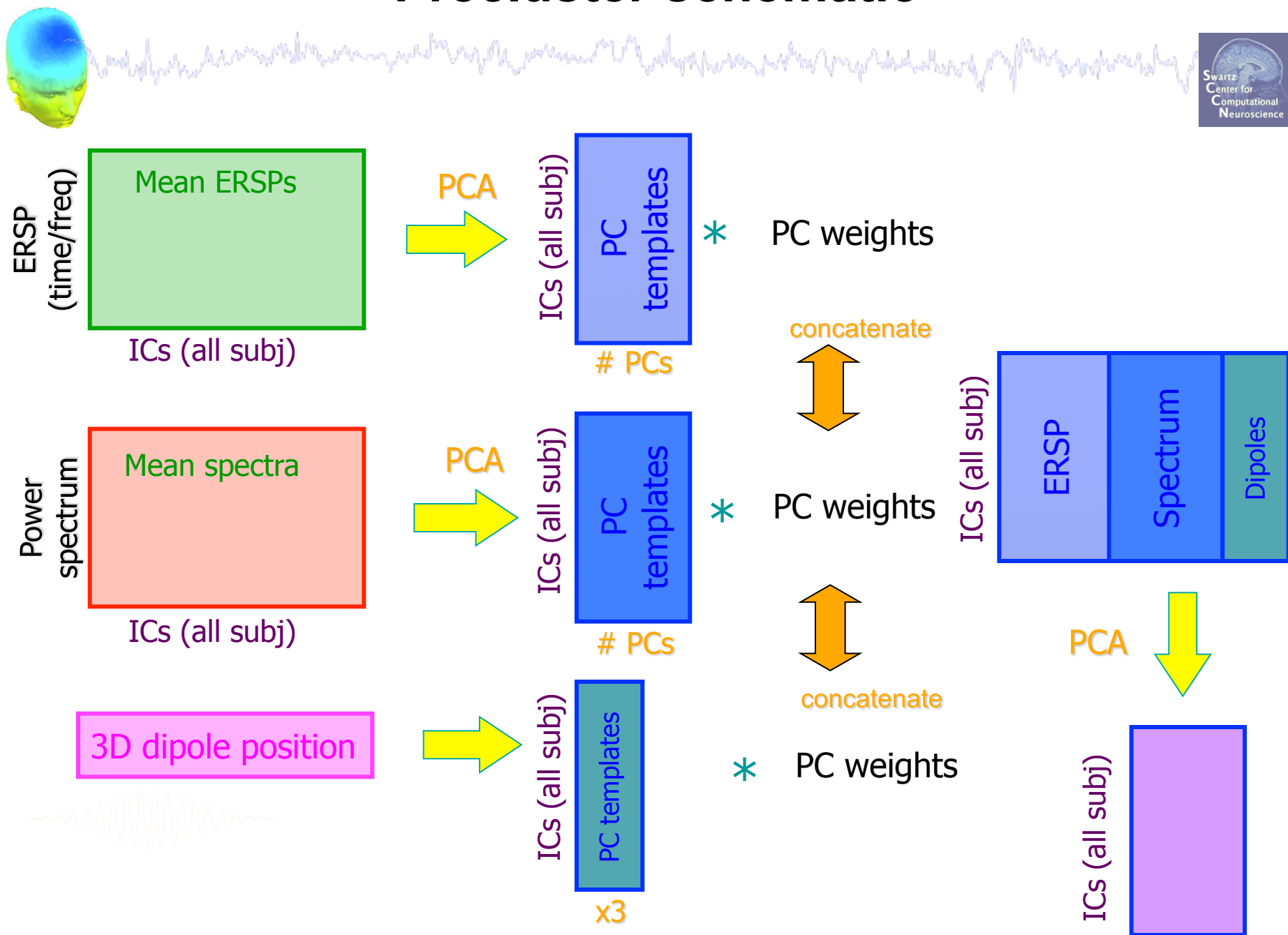
**Time-frequency options**



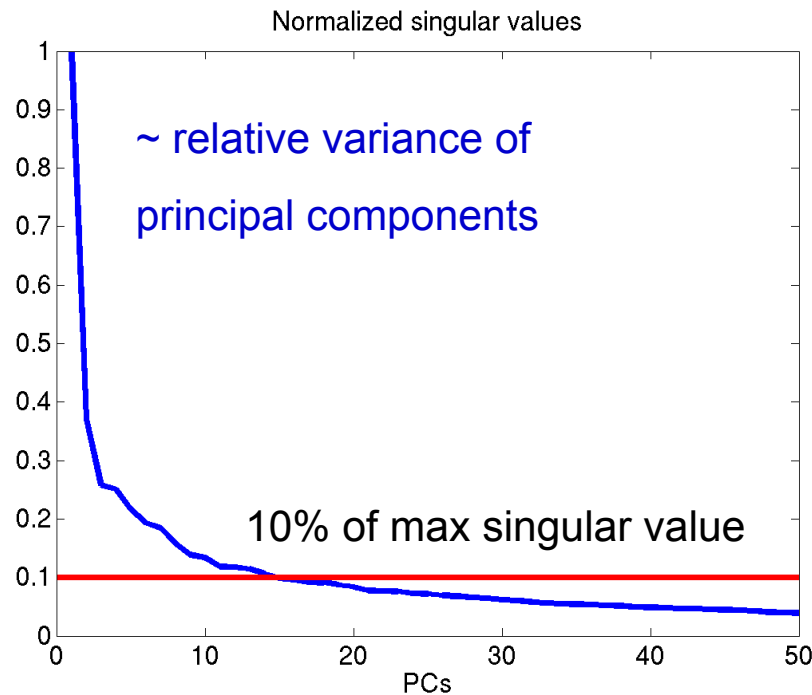
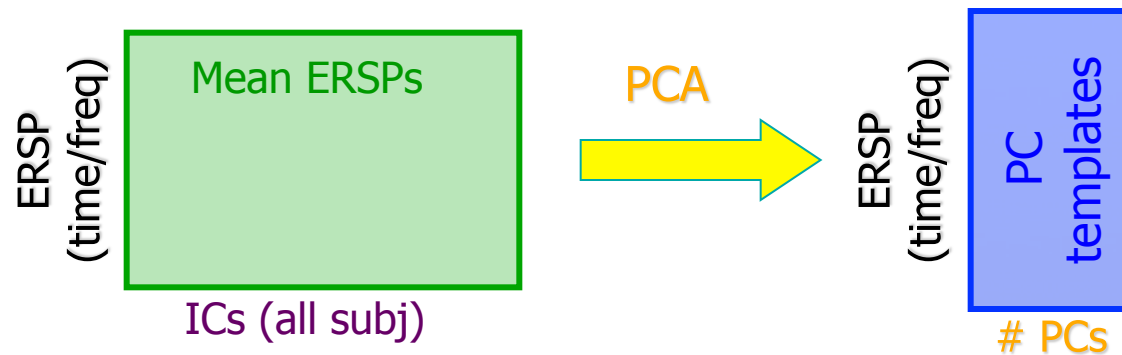
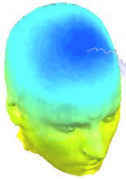
# 3. Cluster components



# Precluster schematic



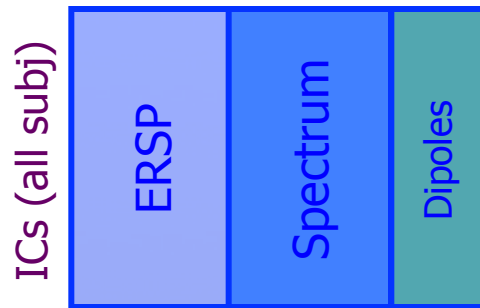
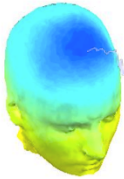
# Precluster: Use singular values from PCA



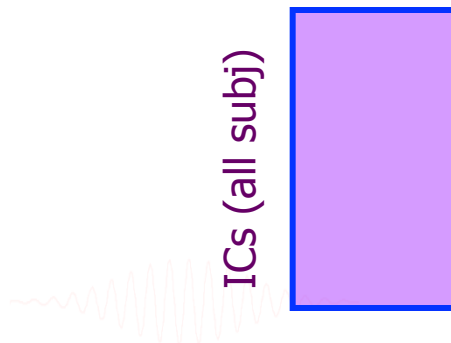
Credit: Julie Onton



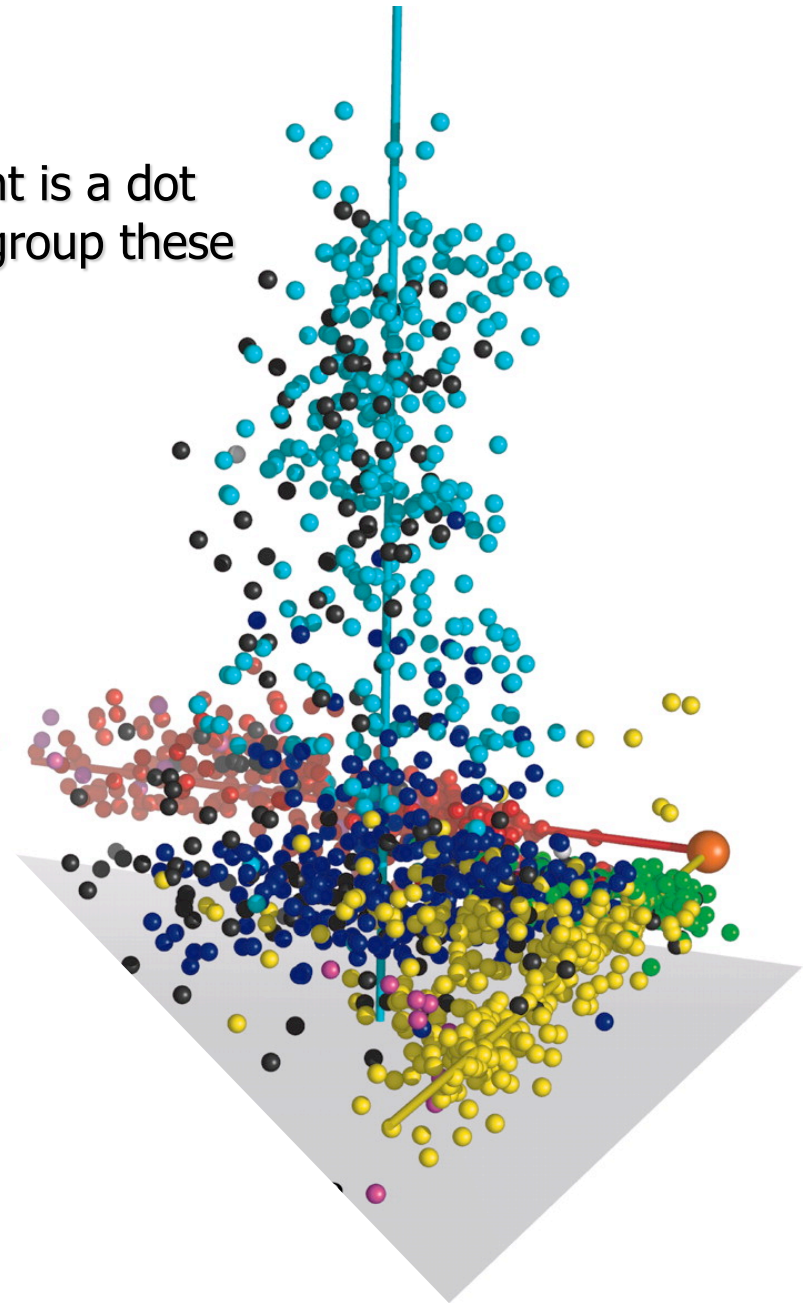
# Precluster schematic



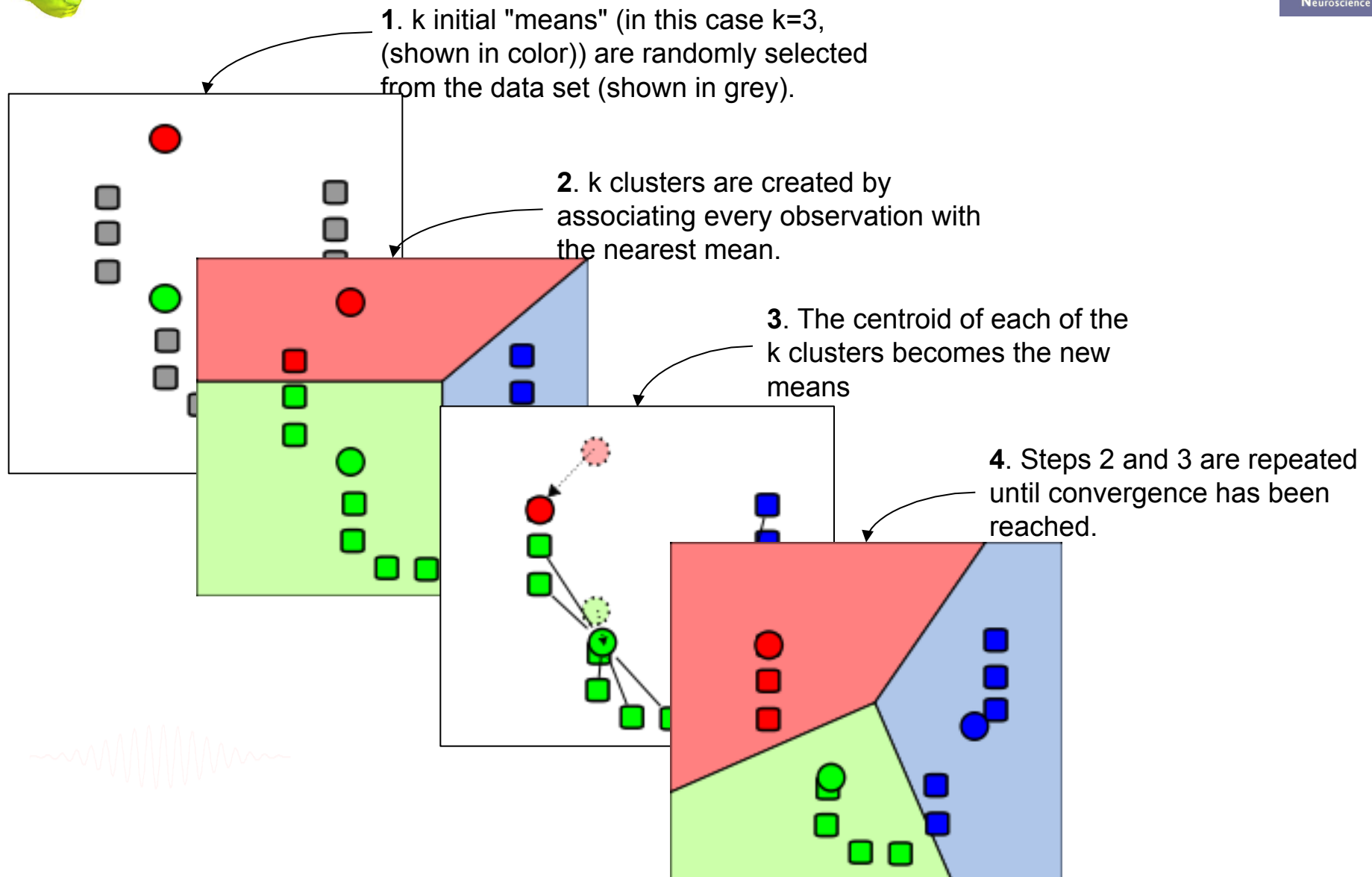
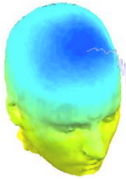
OR

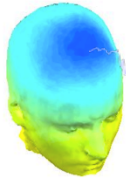


Each component is a dot  
Clustering will group these dots



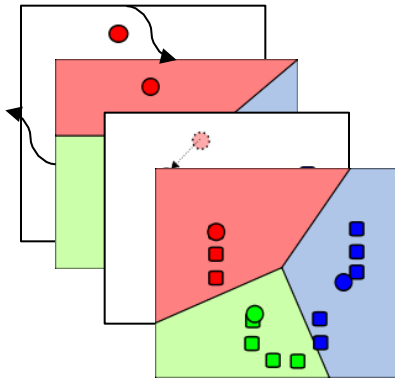
# Classical KMean





# Customized KMean

(no more than 1 session per cluster)



1. A first KMean solution is computed for N clusters

2. Select the cluster with minimum residual distance to centroid

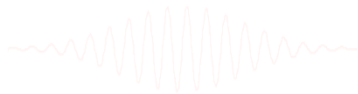
3. Keeps at most one component per session  
(min dist. to centroid)

4. Store the resulting cluster

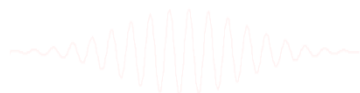
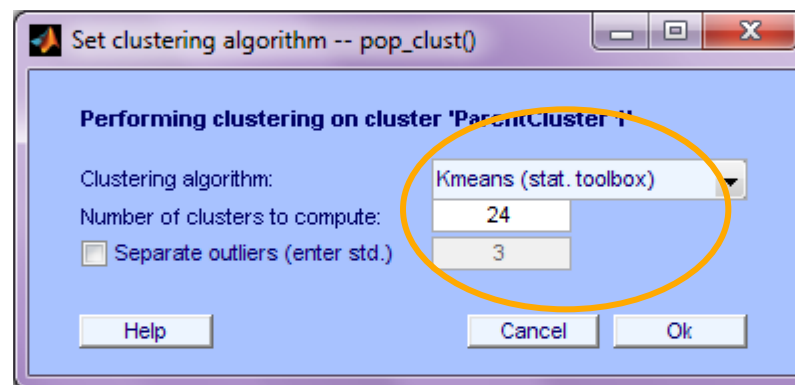
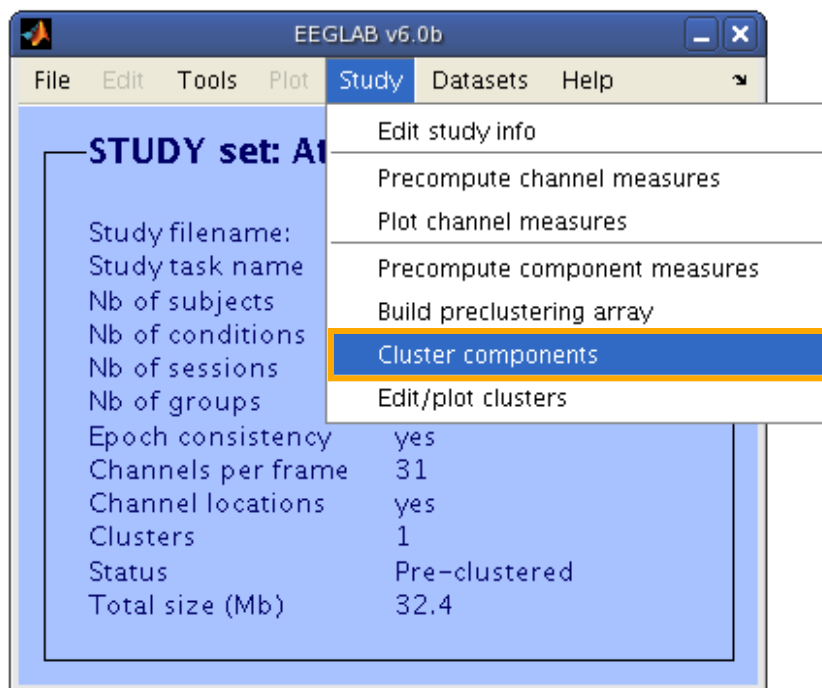
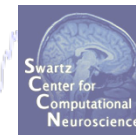
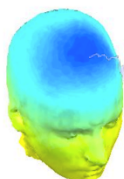
5. Remove the cluster's ICs from the pool of all ICs

6. Compute a new KMean solution for N-1 clusters on the new pool

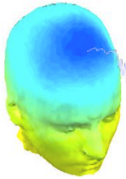
7. Loop until the desired number of selected clusters is reached



# Cluster components



# Choosing data measures



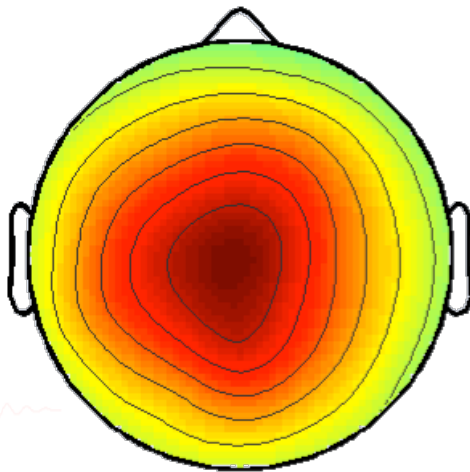
What measure(s) should you use?

- It depends on your final cluster criteria...
  - If for example, your priority is dipole location, then cluster only based on dipole location...

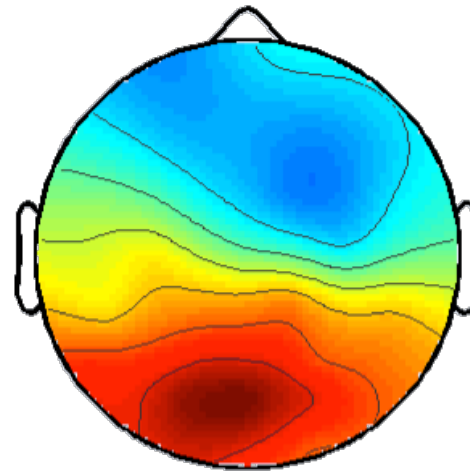
But consider:

- What is the difference between these two components?

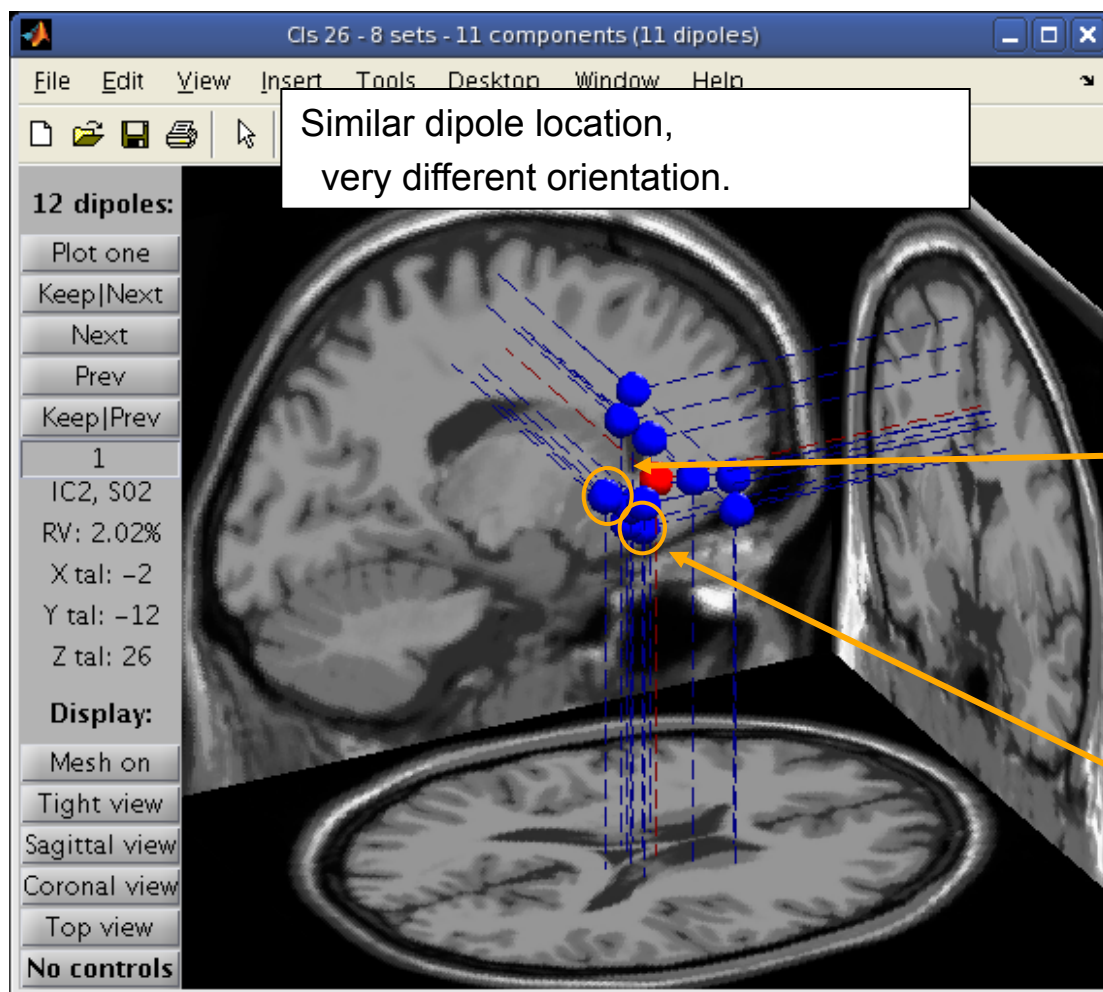
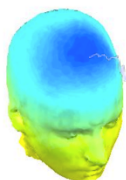
IC2 / S02, Cls 26



IC5 / S05, Cls 26

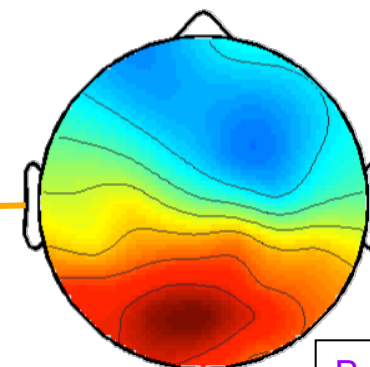


# Choosing data measures

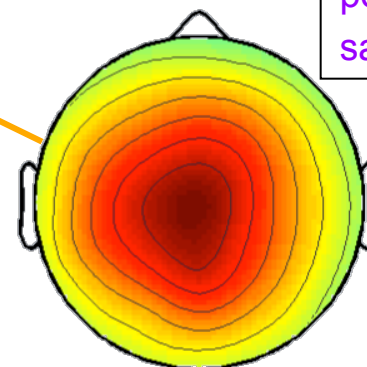


Obvious dramatic effect on  
scalp map topography:

IC5 / S05, Cls 26

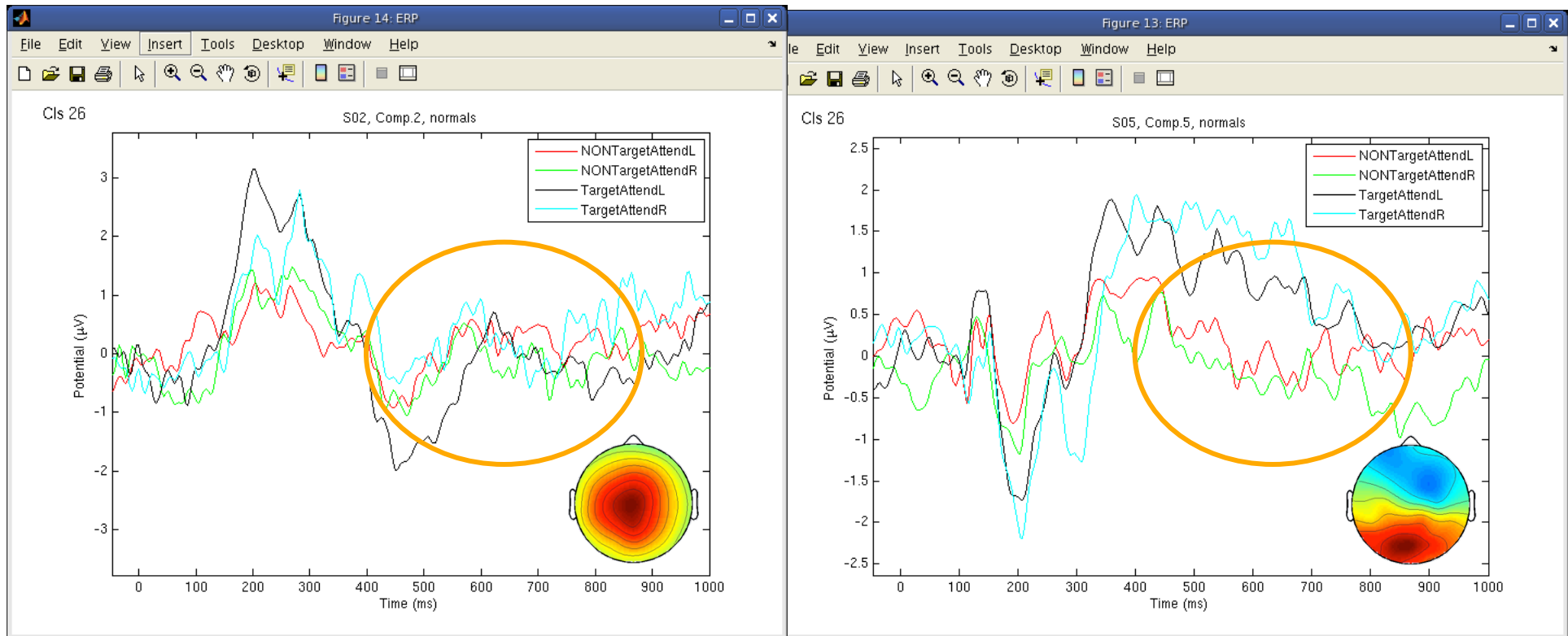
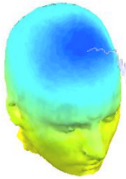


IC2 / S02, Cls 26

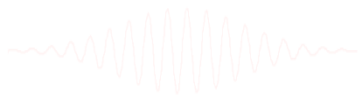


But, do they  
perform the  
same functions?

# Choosing data measures

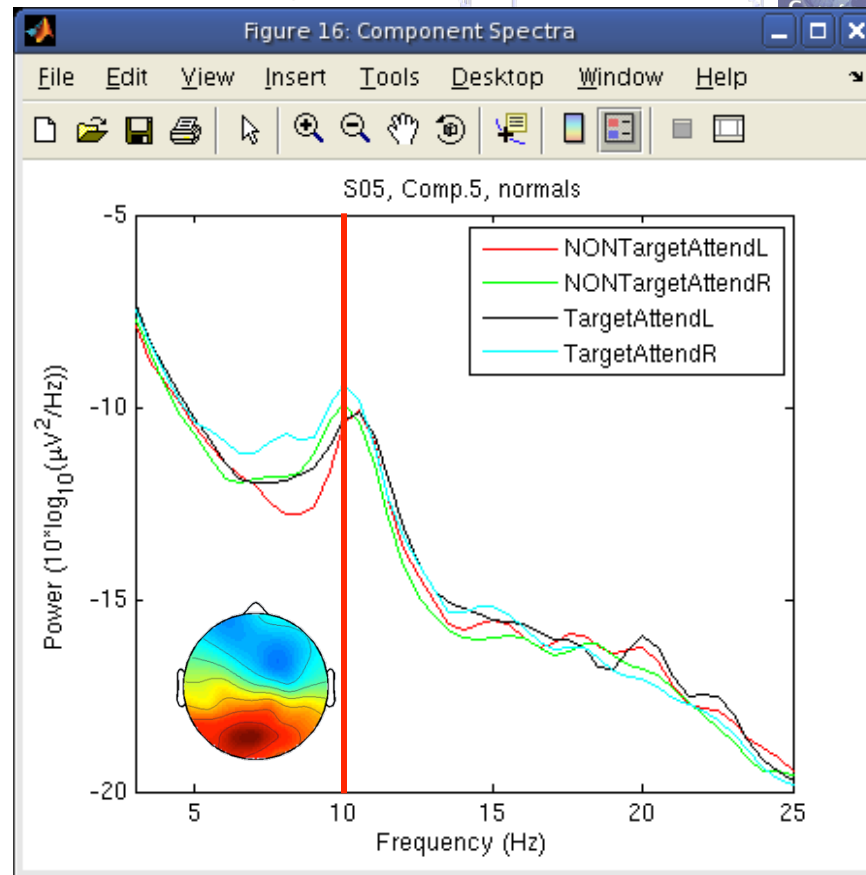
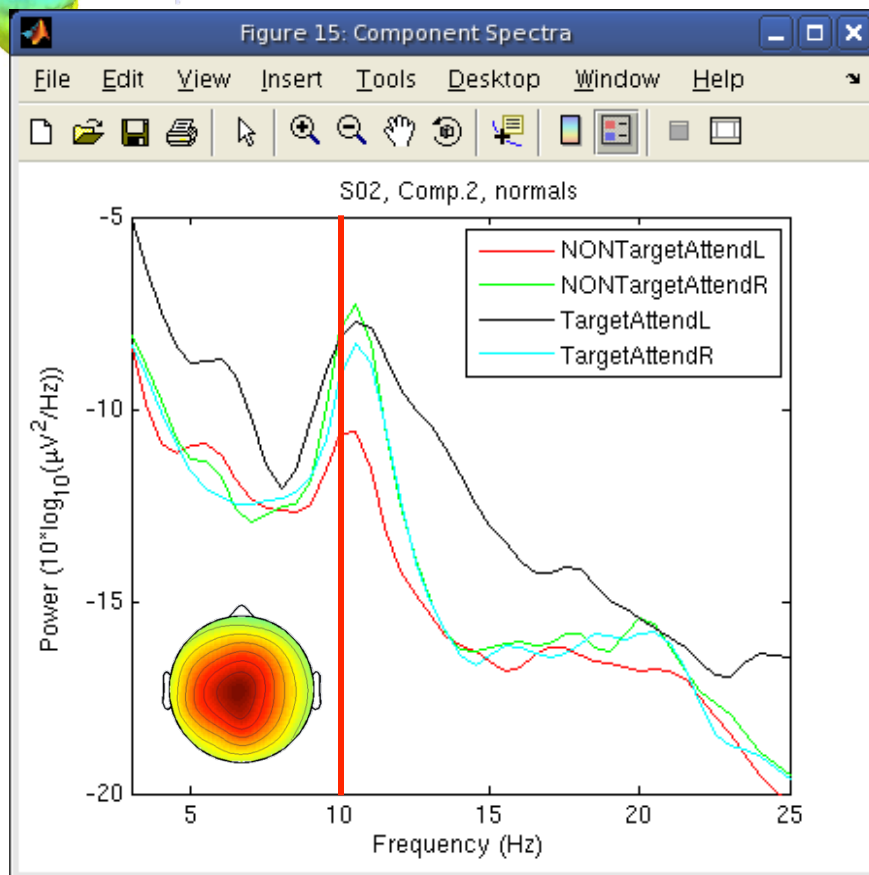


ERPs seem different...



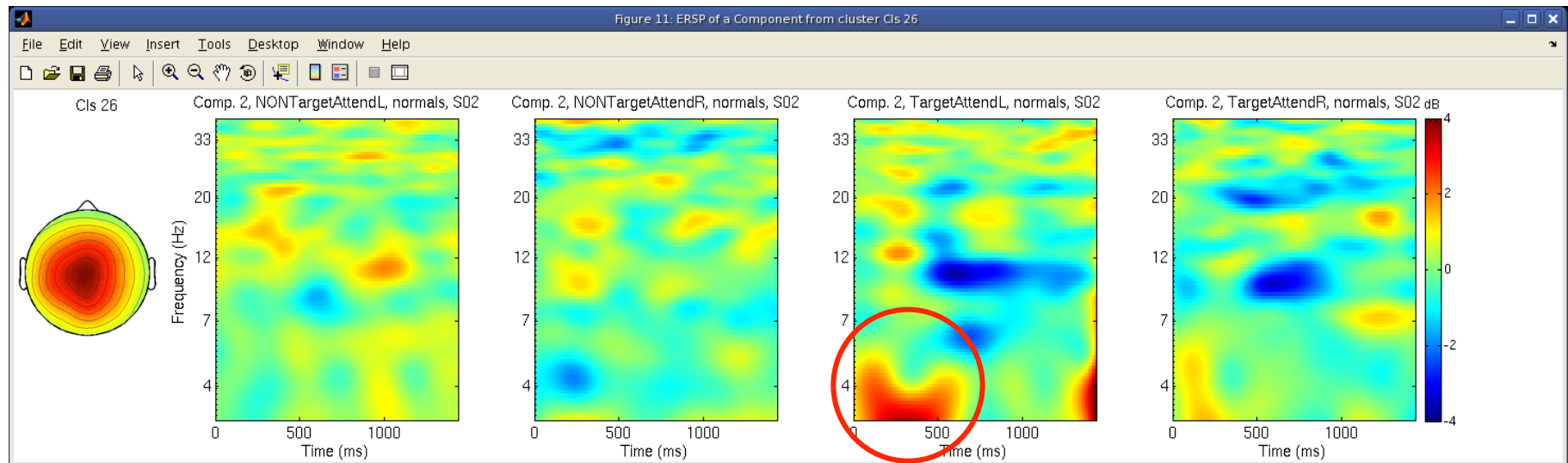
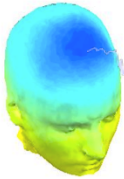


# Choosing data measures

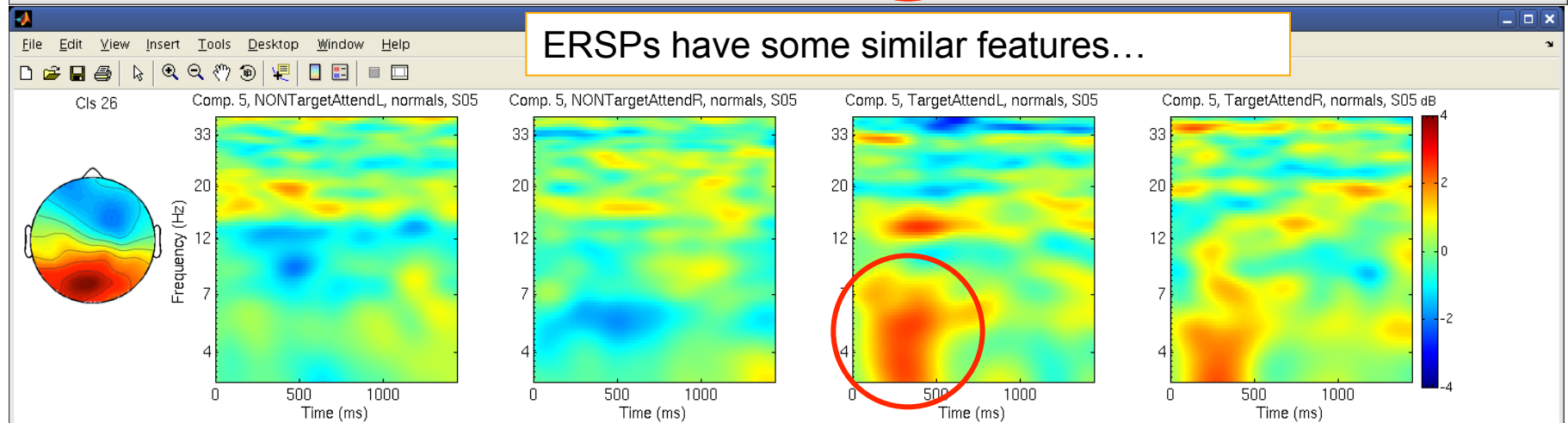


Spectra are similar, but they have  
variable responses to different conditions...

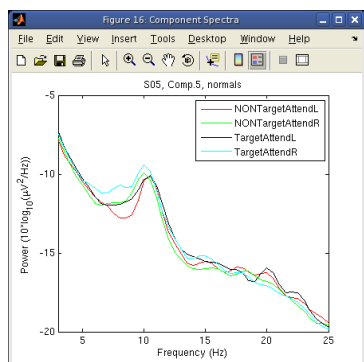
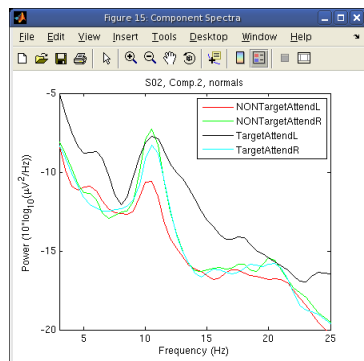
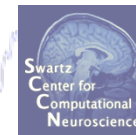
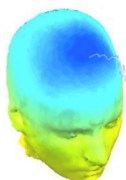
# Choosing data measures



ERSPs have some similar features...



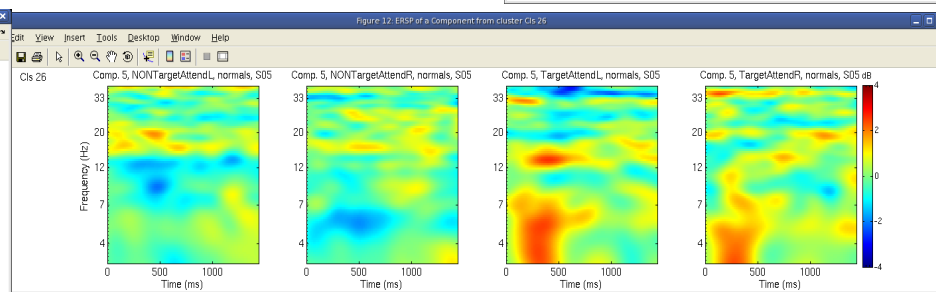
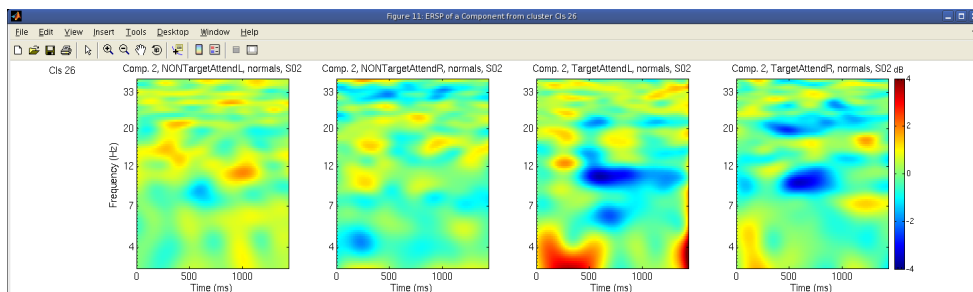
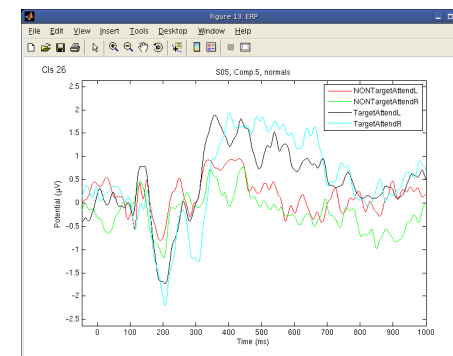
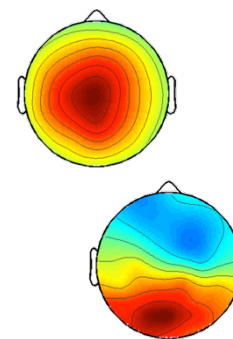
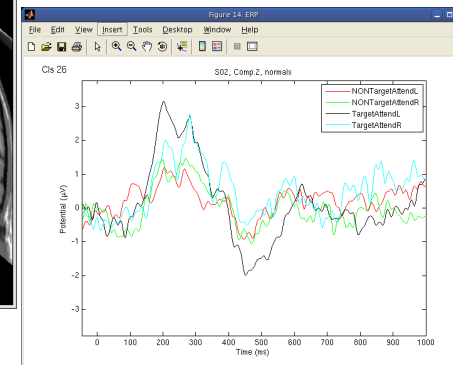
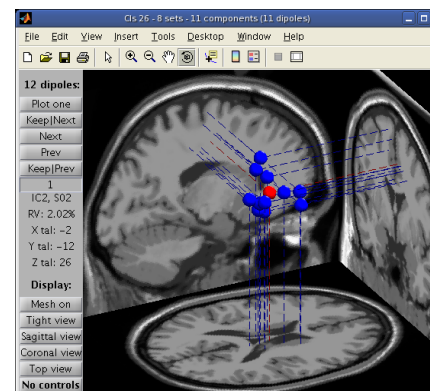
# Choosing data measures



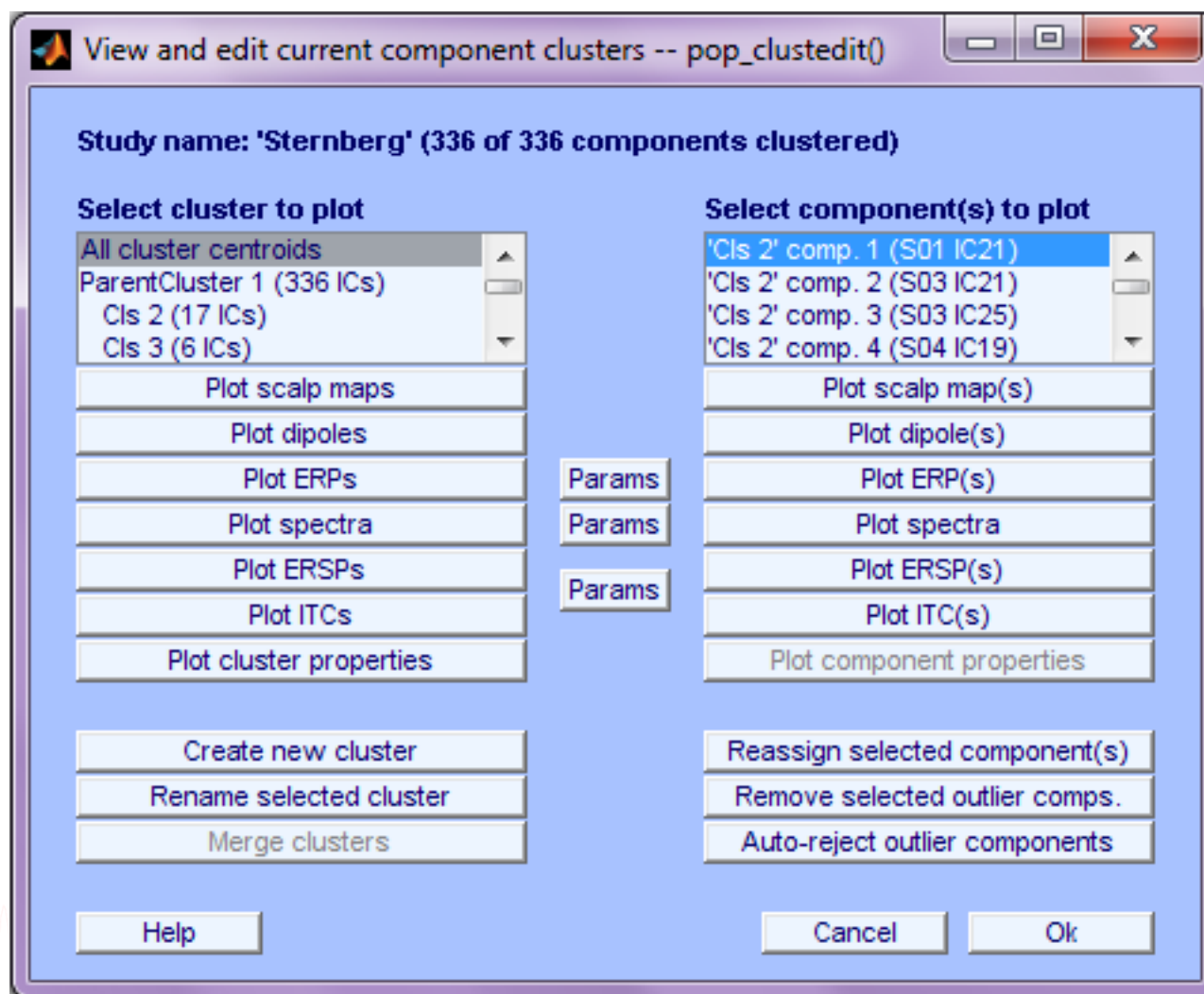
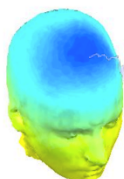
What data measures  
should you use?

It depends...

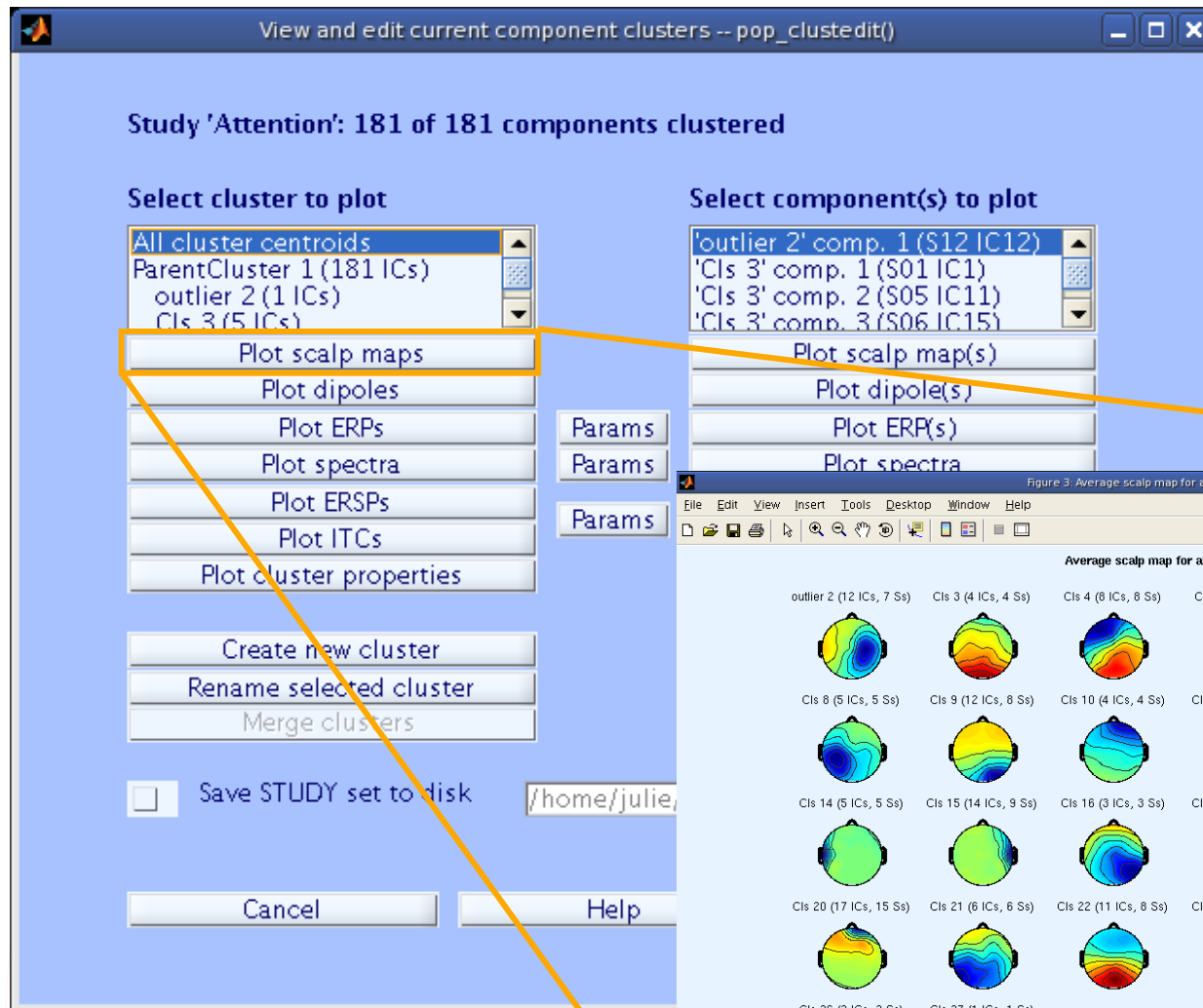
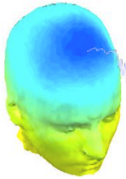
- broadly-matched ICs: use many/all of the measures.
- specifically-matched ICs: use one/few of the measures.



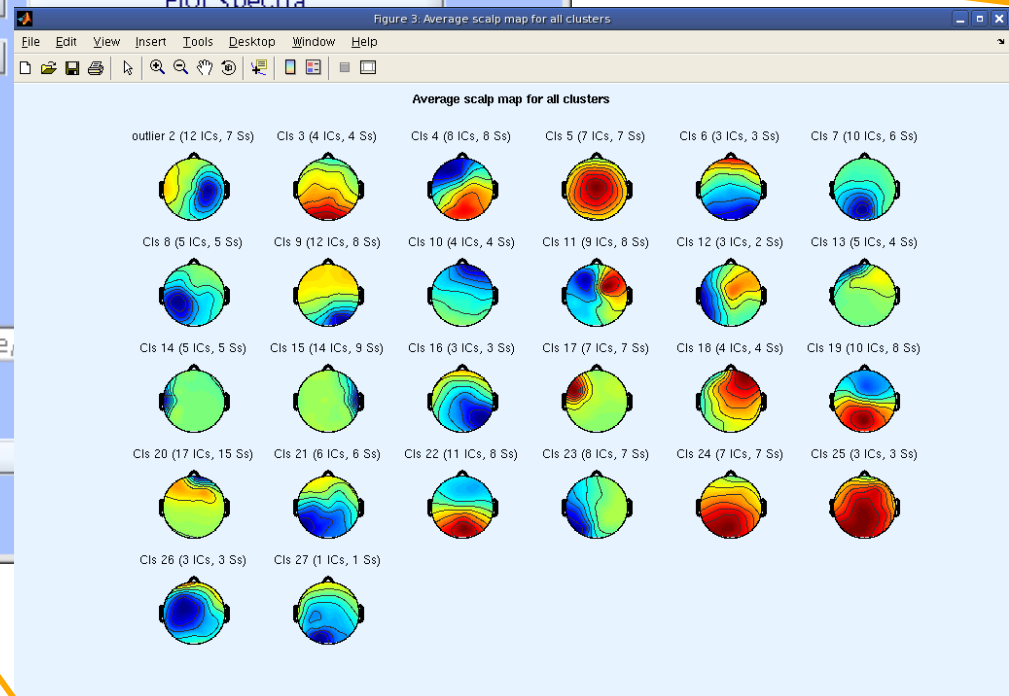
# Plot/edit clusters



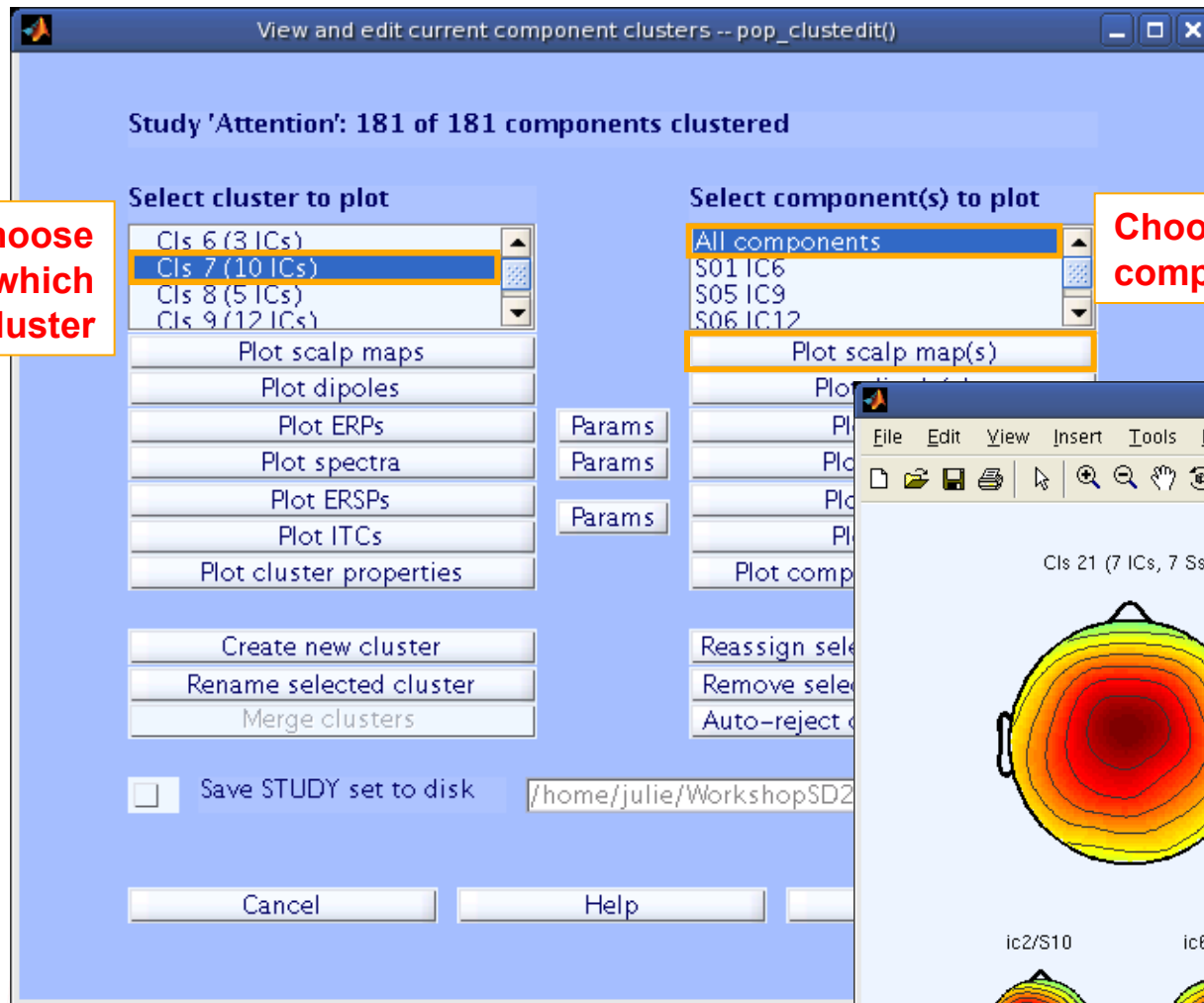
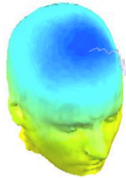
# Plot cluster data



Plot mean scalp maps for easy reference

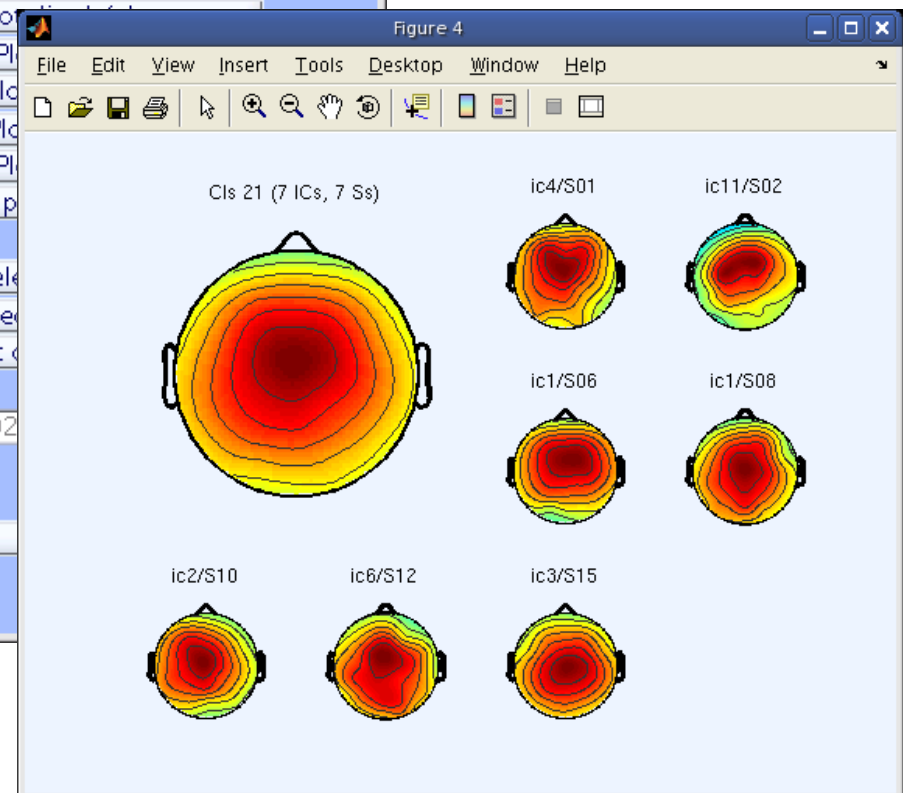


# Plot cluster data

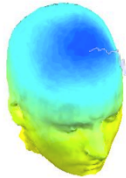


Choose which cluster

Choose which components



# Plot cluster data



Clis 19 - 5 sets - 14 components (14 dipoles)

File Edit View Insert Tools Desktop Window Help

15 dipoles:

Plot one  
Keep|Next  
Next  
Prev  
Keep|Prev

1  
IC3, S02  
RV: 2.62%  
X tal: -6  
Y tal: -13  
Z tal: 21

Display:

Mesh on  
Tight view  
Sagittal view  
Coronal view  
Top view  
No controls

Plot scalp maps  
Plot dipoles  
Plot ERPs  
Plot spectra  
Plot ERSPs  
Plot ITCs  
Plot cluster properties

Create new cluster  
Rename selected cluster  
Merge clusters

☐ Save STUDY set to disk

/home/julie/workshop06/5subjects/WSstudy.study

Cancel Help Ok

component clusters -- pop\_clustedit()

Select component(s) to plot

All components  
S02 IC3  
S02 IC11  
S02 IC12  
S02 IC17

Plot scalp map(s)  
Plot dipole(s)  
Plot ERP(s)  
Plot spectra  
Plot ERSP(s)  
Plot ITC(s)  
Plot component properties

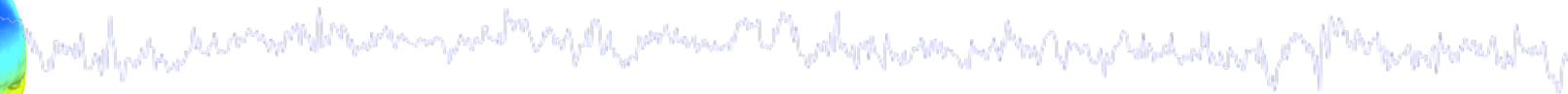
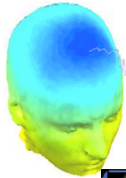
Params  
Params  
Params

Reassign selected component(s)  
Remove selected outlier comps.  
Auto-reject outlier components





# Plot cluster data



View and edit current component clusters -- pop\_clustedit()

Study #: 151 of 151 components clustered

Select cluster to plot

- Cls 15 (8 ICs)
- Cls 16 (6 ICs)
- Cls 17 (4 ICs)
- Cls 18 (14 ICs)
- Cls 19 (14 ICs)

Plot scalp maps

Plot dipoles

Plot ERPs

Plot spectra

Params

Params

Select component(s) to plot

- All components
- S02 IC3
- S02 IC11
- S02 IC12
- S02 IC17

Plot scalp map(s)

Plot dipole(s)

Plot ERP(s)

Plot spectra

Set ERP plotting parameters -- pop\_erpparams()

Time range in ms [low high]

Plot scalp map at latency [ms]

☐ Plot conditions on the same panel

☐ Plot groups on the same panel

Statistical method to use

☐ Compute condition statistics

☐ Compute group statistics

☐ Use single trials (when available)

☐ Use False Discovery Rate to correct for multiple comparisons

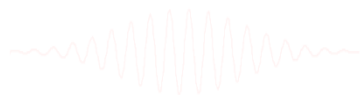
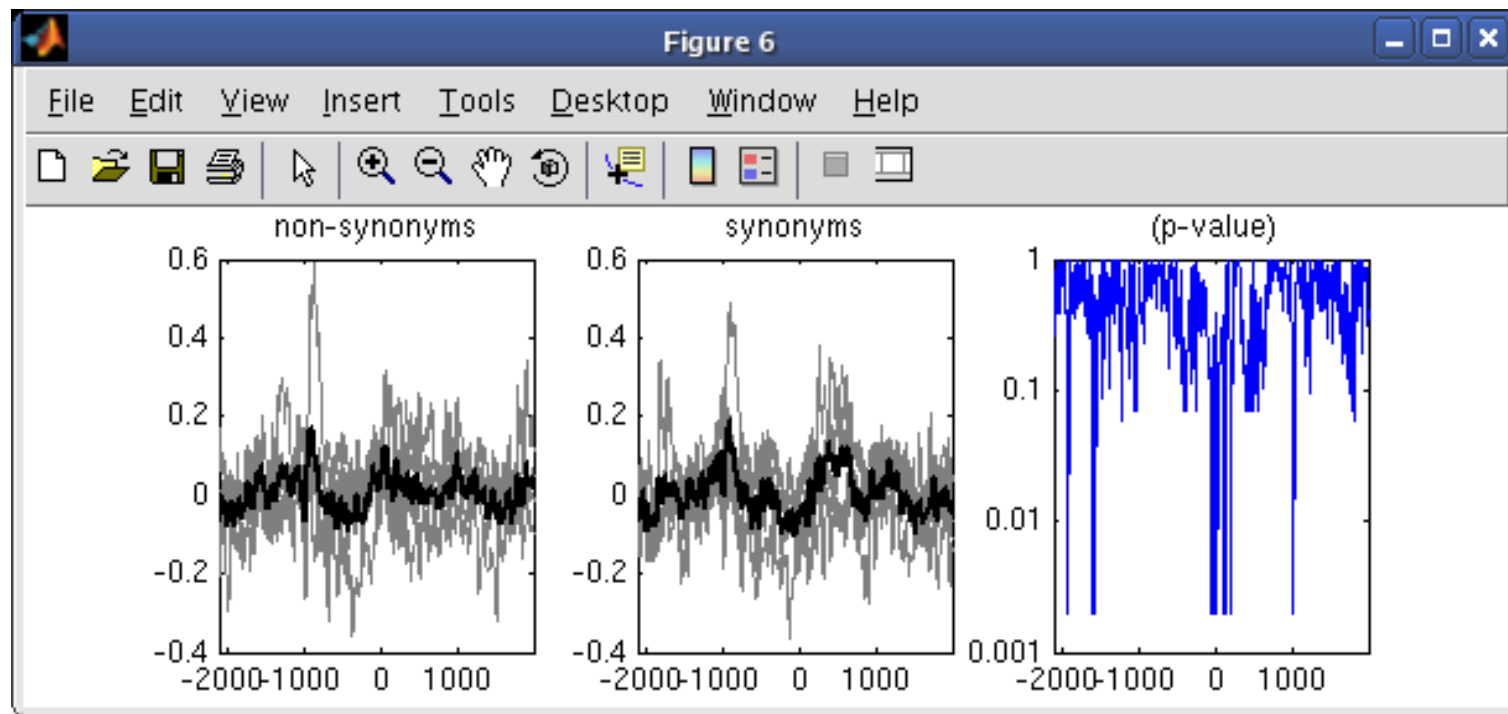
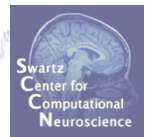
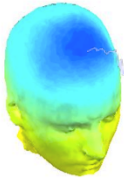
Plot limits in uV [low high]

Display filter in Hz [high]

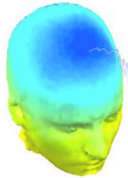
Statistical threshold (p<)

Help Cancel Ok

# Plot cluster ERP



# Other plotting options...



Set ERP plotting parameters -- pop\_erpparams()

Time range in ms [low high]

Plot limits in uV [low high]

Plot scale map at latency [ms]  NaN

Display filter in Hz [high]

☐ Plot conditions on the same panel

☐ Plot groups on the same panel

Statistical method to use  Parametric

Statistical threshold (p<)

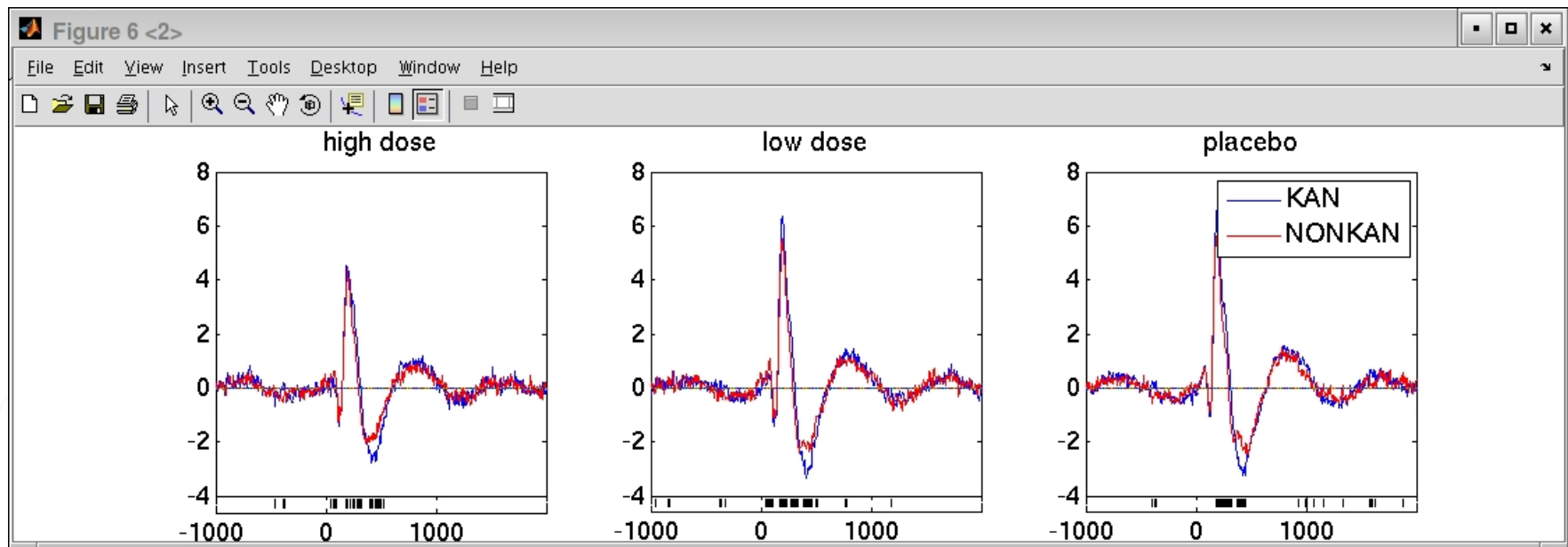
☐ Compute condition statistics

☐ Compute group statistics

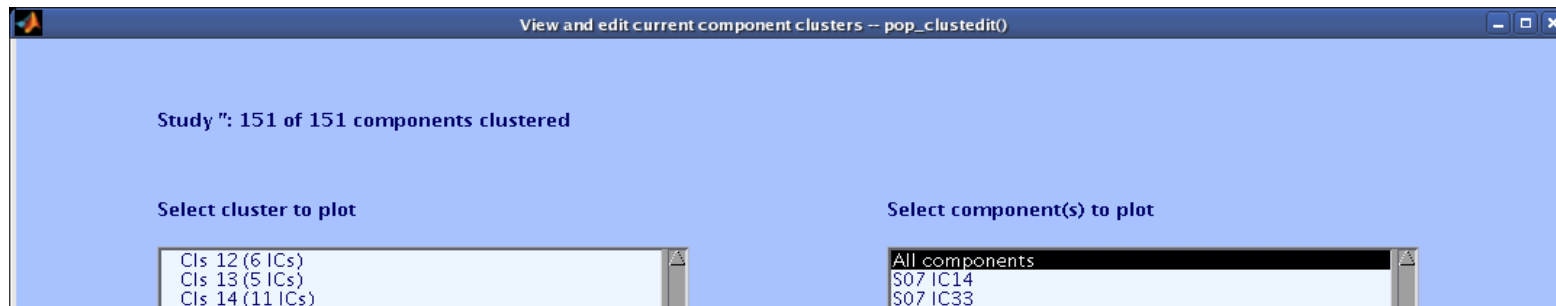
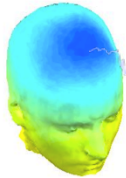
☐ Use single trials (when available)

☐ Use False Discovery Rate to correct for multiple comparisons

Help Cancel Ok



# Reassigning components



Select component(s) to plot

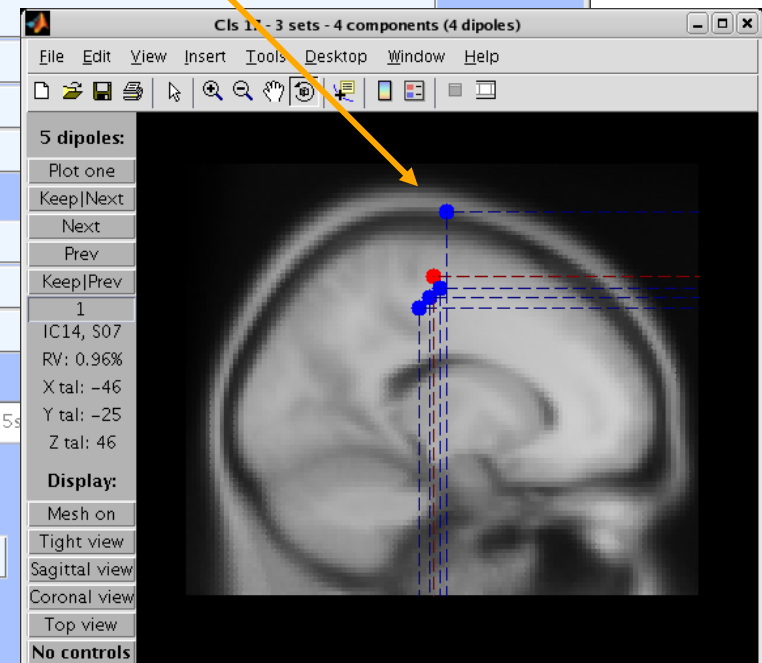
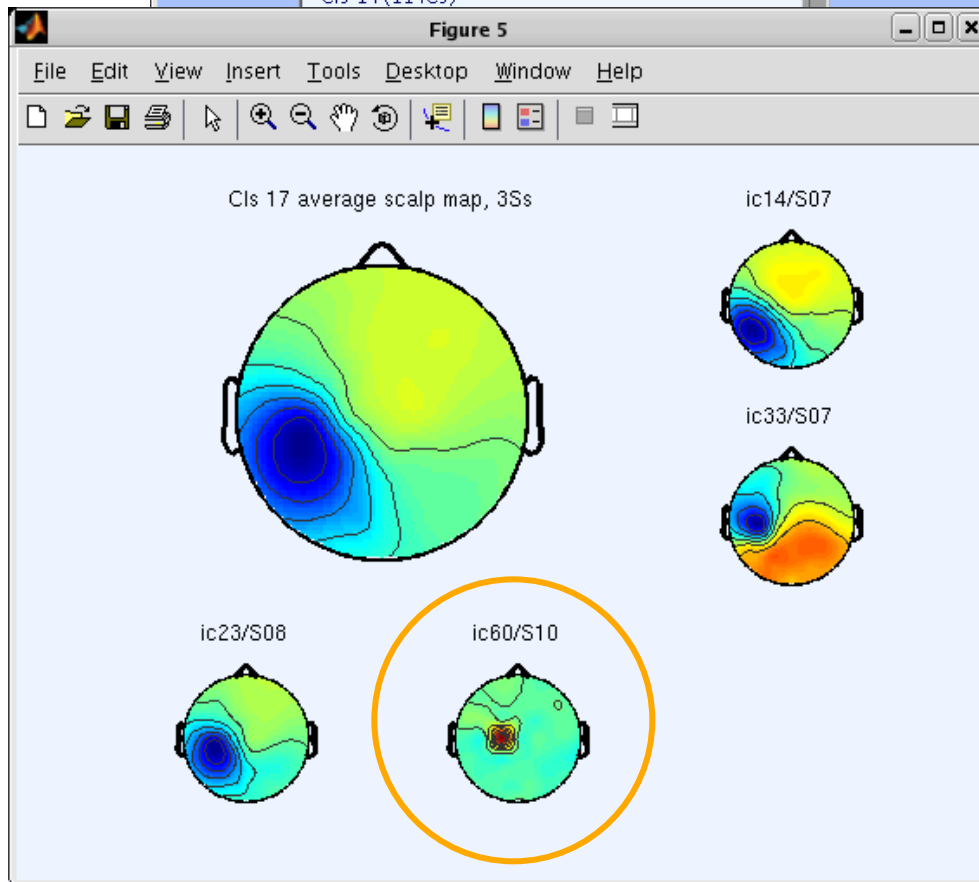
All components

- S07 IC14
- S07 IC33
- S08 IC23
- S10 IC60

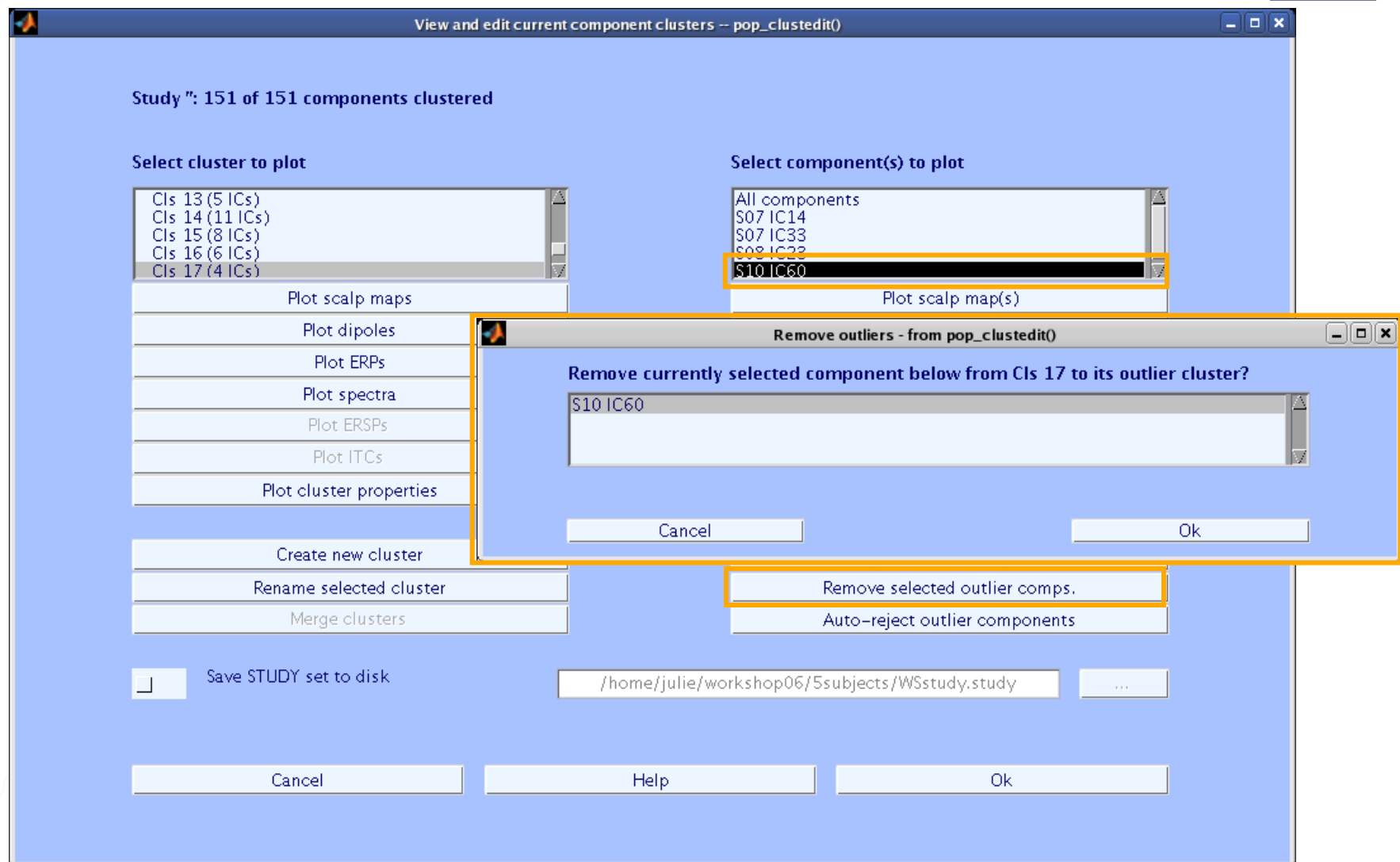
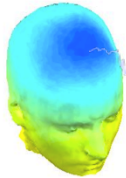
Plot scalp map(s)

Plot dipole(s)

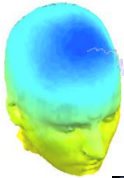
Plot ERP(s)



# Reassigning components



# Outlier cluster reassignment



View and edit current component clusters -- pop\_clustedit()

Study ": 151 of 151 components clustered

Select cluster to plot

- Cls 16 (6 ICs)
- Cls 17 (3 ICs)
- Cls 18 (14 ICs)
- Cls 19 (14 ICs)
- Outliers Cls 17 20 (1 ICs)

Plot scalp maps

Plot dipoles

Plot ERPs

Plot spectra

Plot ERSPs

Plot ITCs

Plot cluster properties

Create new cluster

Rename selected cluster

Merge clusters

Save STUDY set to disk

/home/julie/workshop06/5subjects/WSstudy.study

Cancel Help Ok

Select component(s) to plot

- All components
- S10 IC60

Plot scalp map(s)

Plot dipole(s)

Plot ERP(s)

Plot spectra

Plot ERSP(s)

Plot ITC(s)

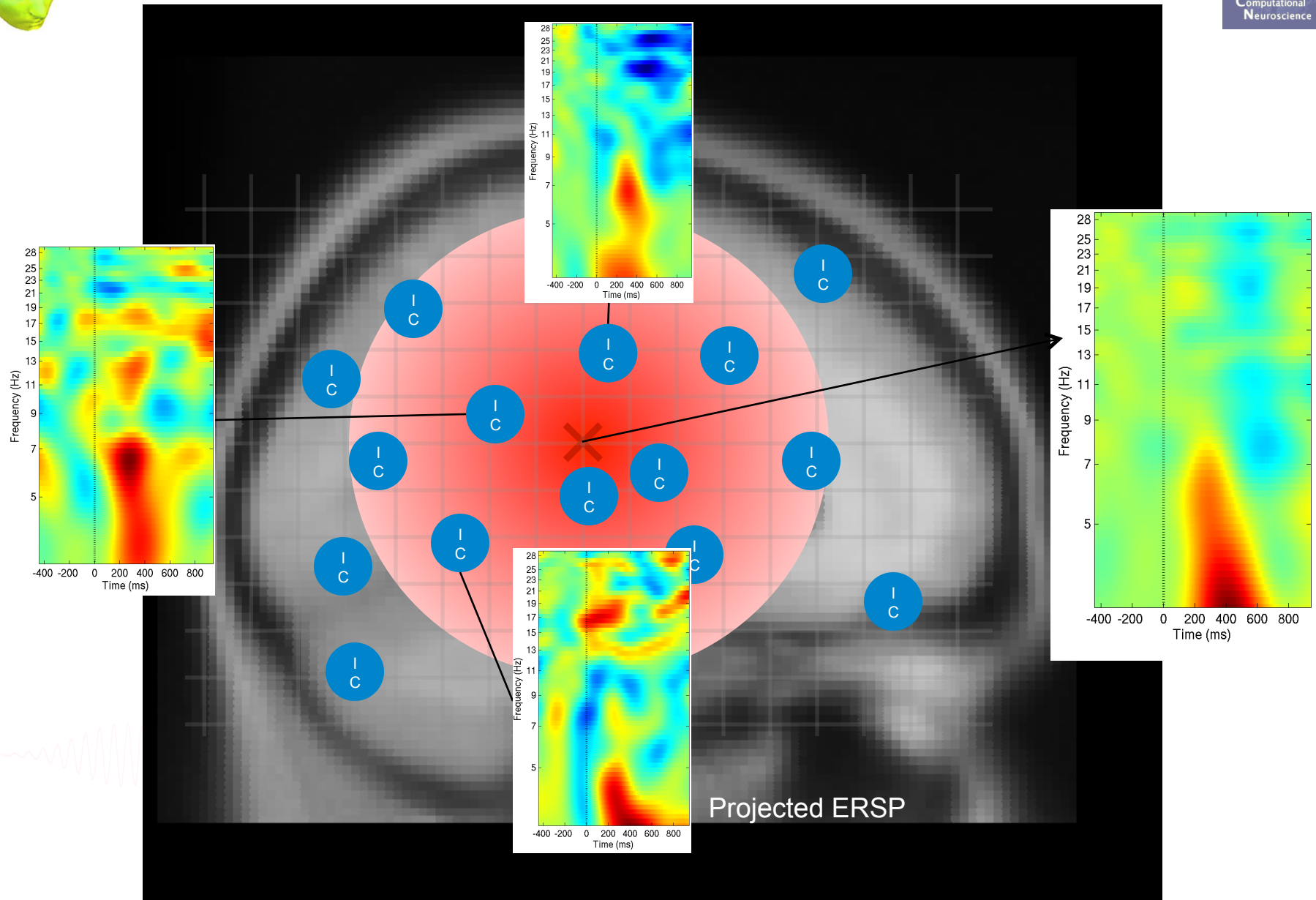
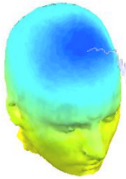
Plot component properties

Reassign selected component(s)

Remove selected outlier comps.

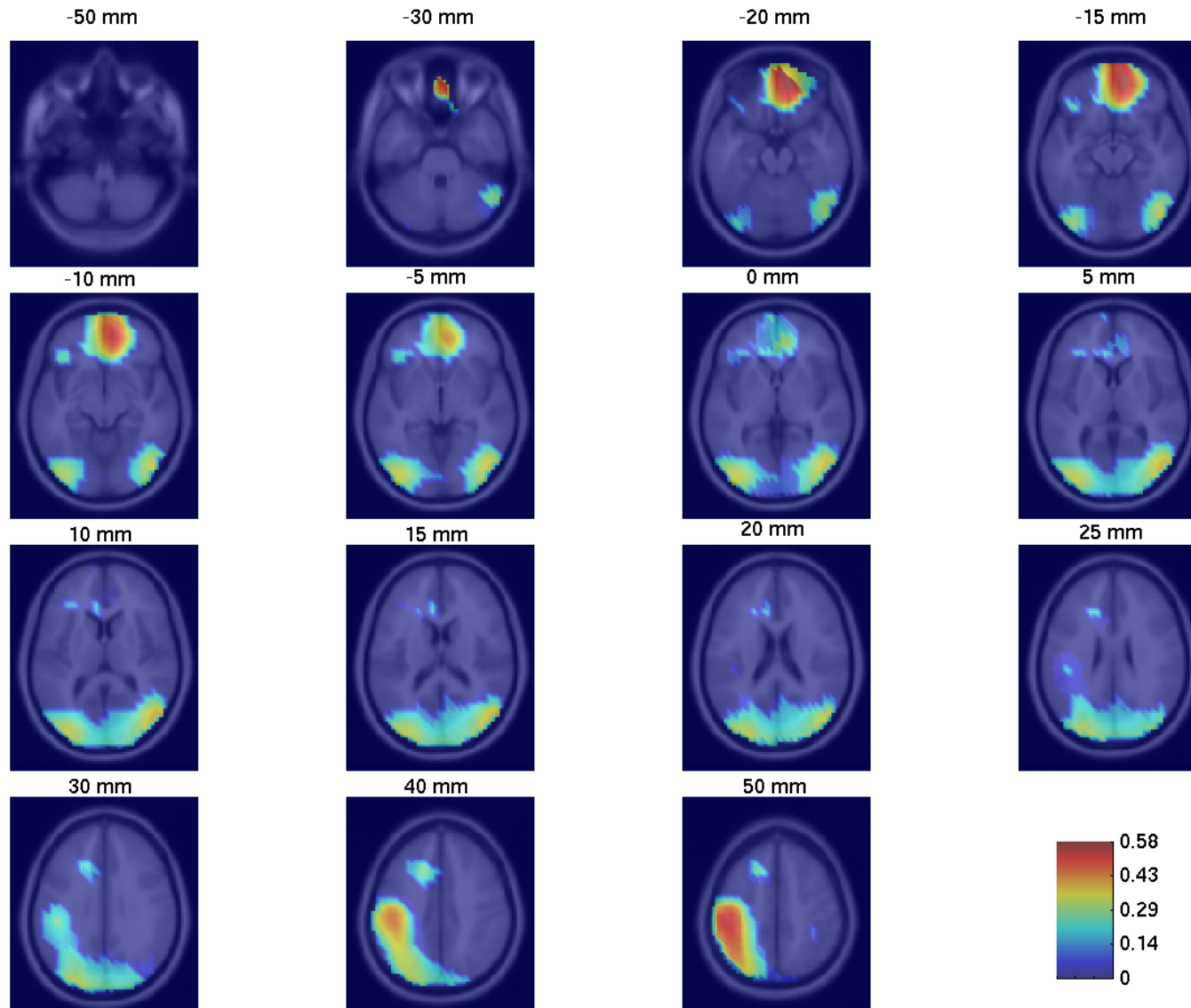
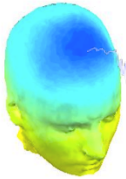
Auto-reject outlier components

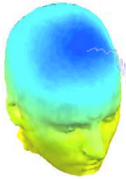
# Measure Projection Toolbox





# Measure Projection Toolbox





## Exercise

Precluster (pre-computation already done) and cluster components using measures of your choice. Experiment with different measures.

