

# Independent Component Clustering



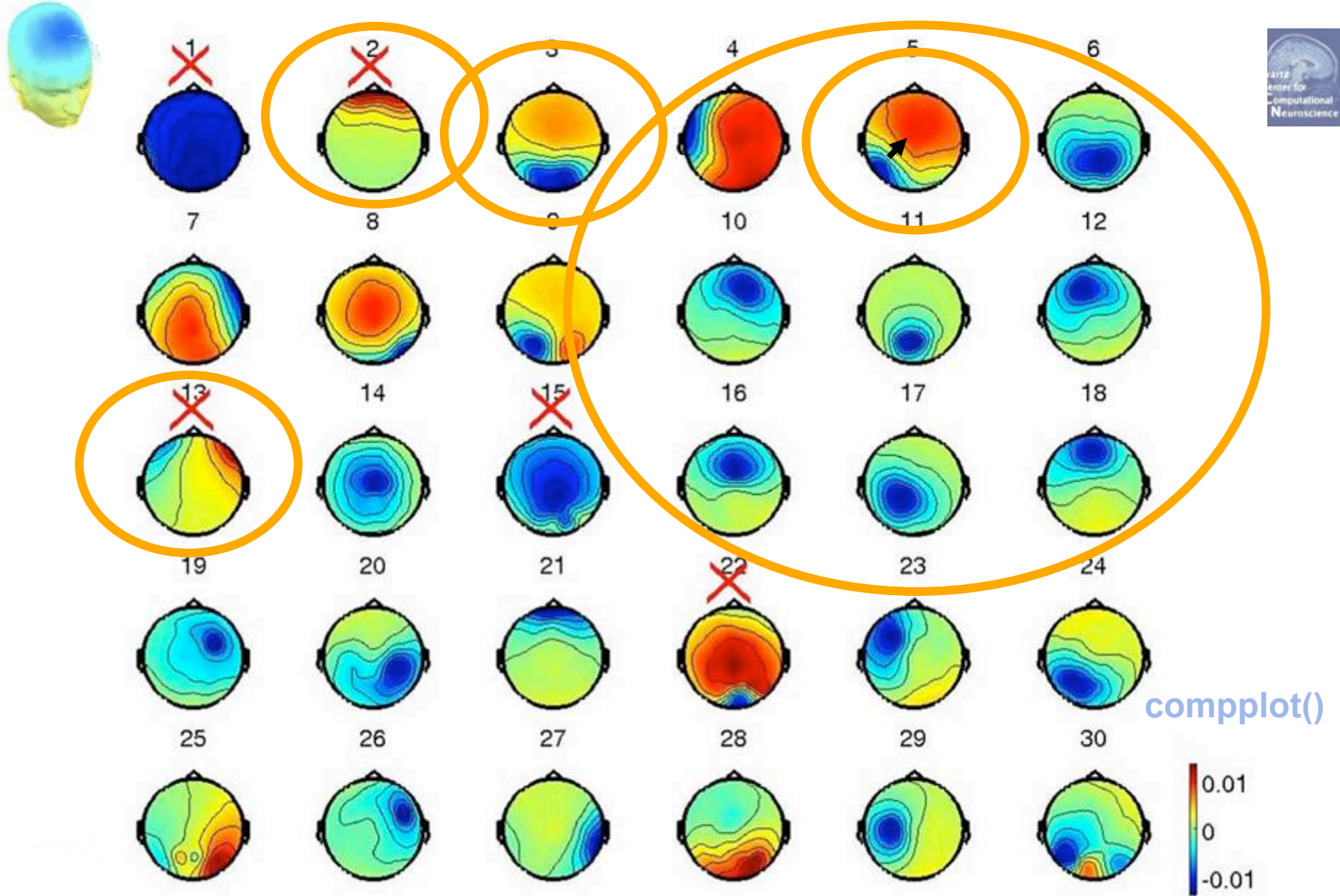
**Scott Makeig**

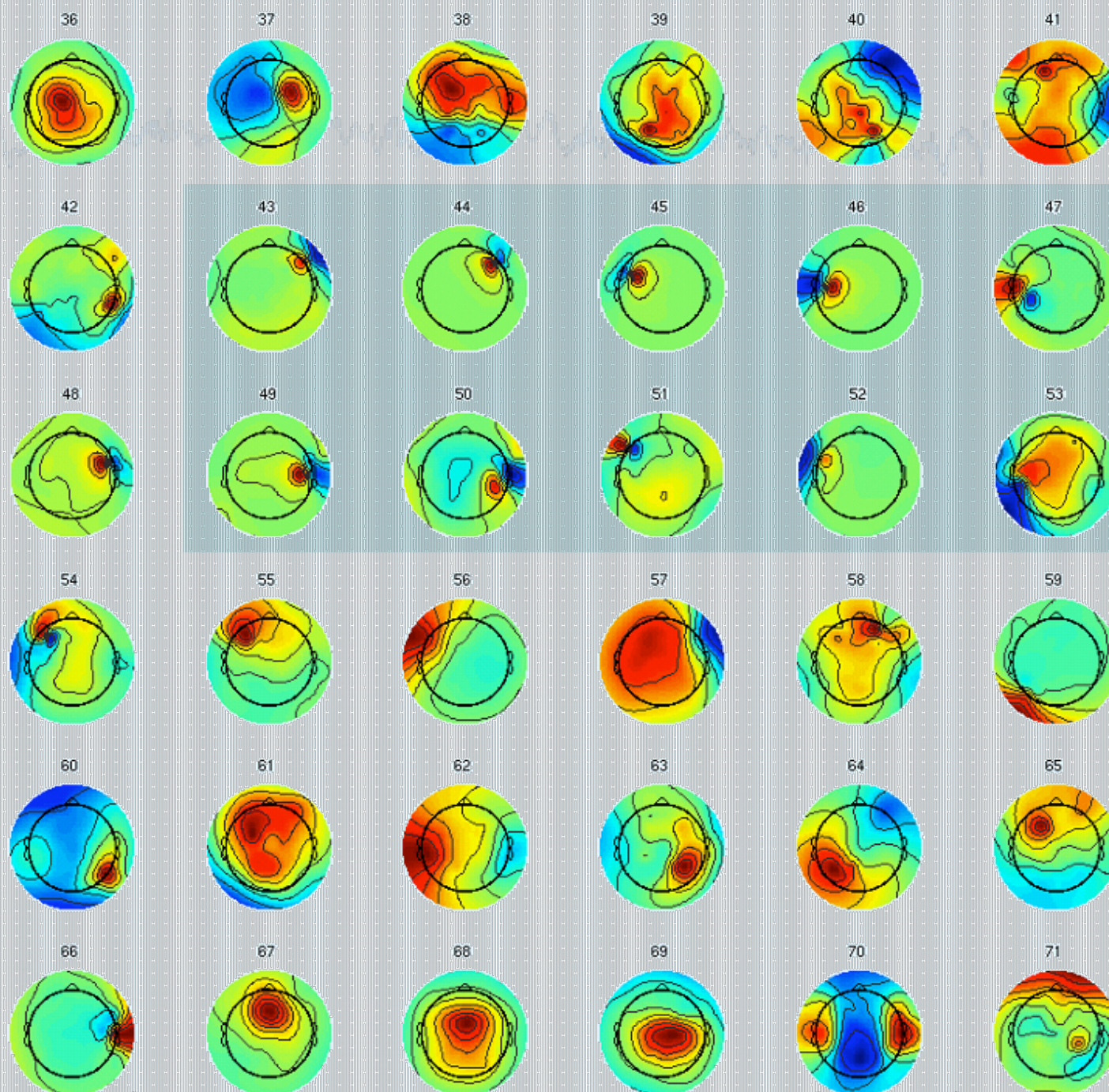
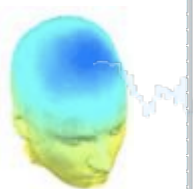
Institute for Neural Computation  
University of California San Diego

EEGLAB June, 2009

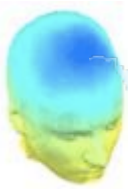
Aspet, France

# Largest 30 independent components (single subject)





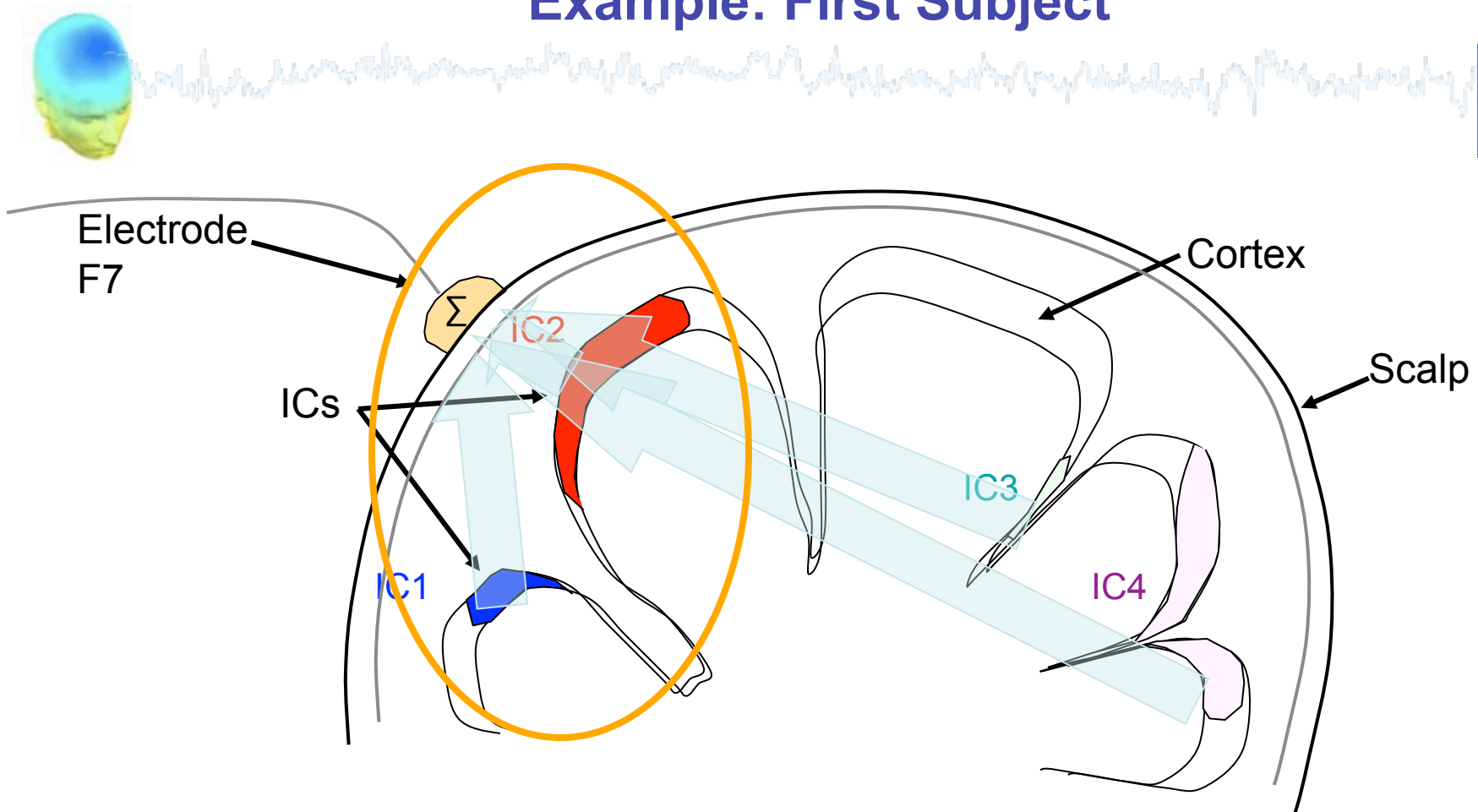
# Channel-based clustering



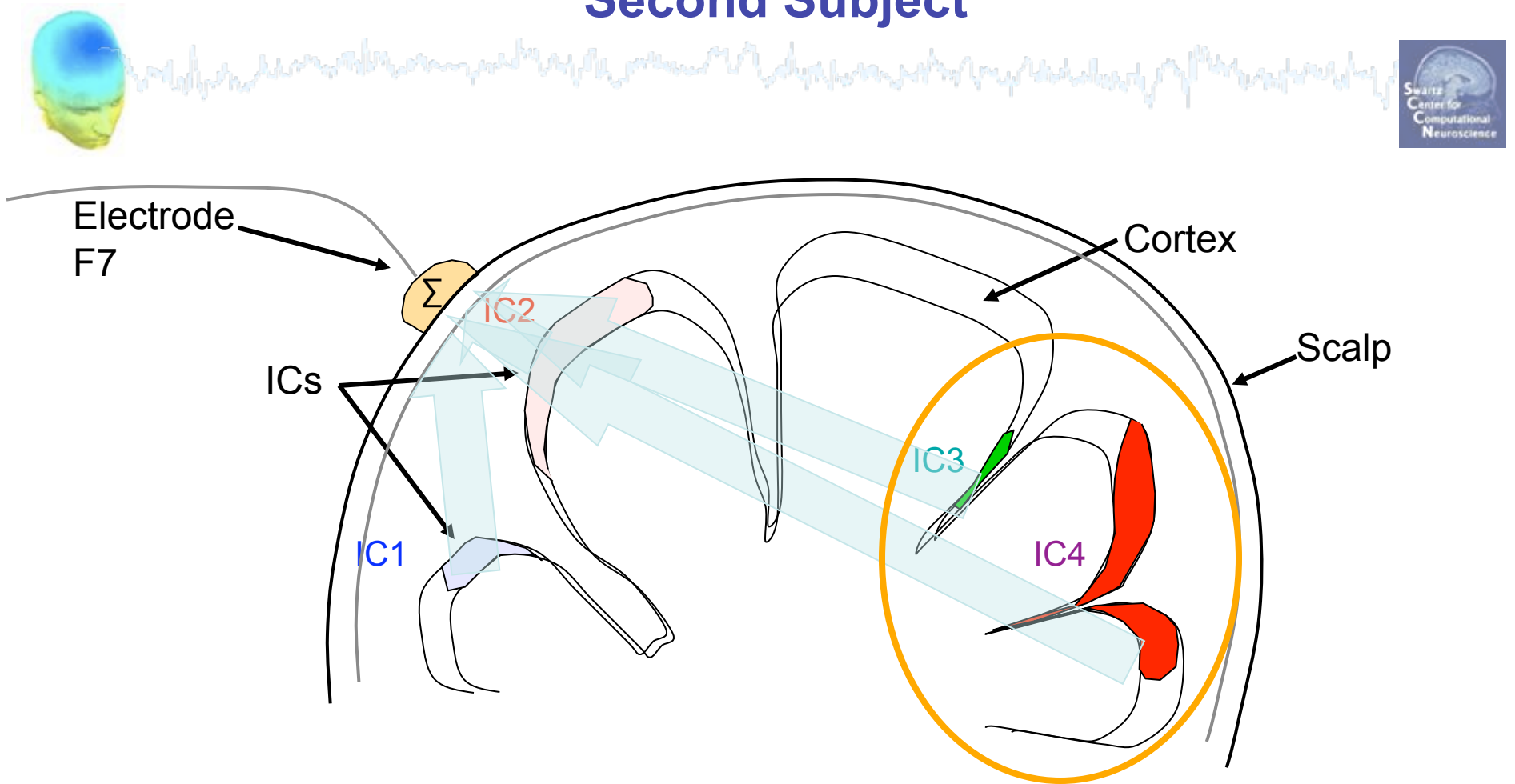
- ICA transforms the data from a ***channel basis*** (activity recorded at each scalp channel)
  - to a ***component basis*** (activity computed at each independent spatially-filtered (cortical or non-cortical) component process).
- Normally, EEG researchers assume that each electrode, say F7 is equivalent to the F7 of each subject – and then ‘cluster’ their data by identifying channels with equivalent scalp locations ...
- But this is only *roughly* correct!



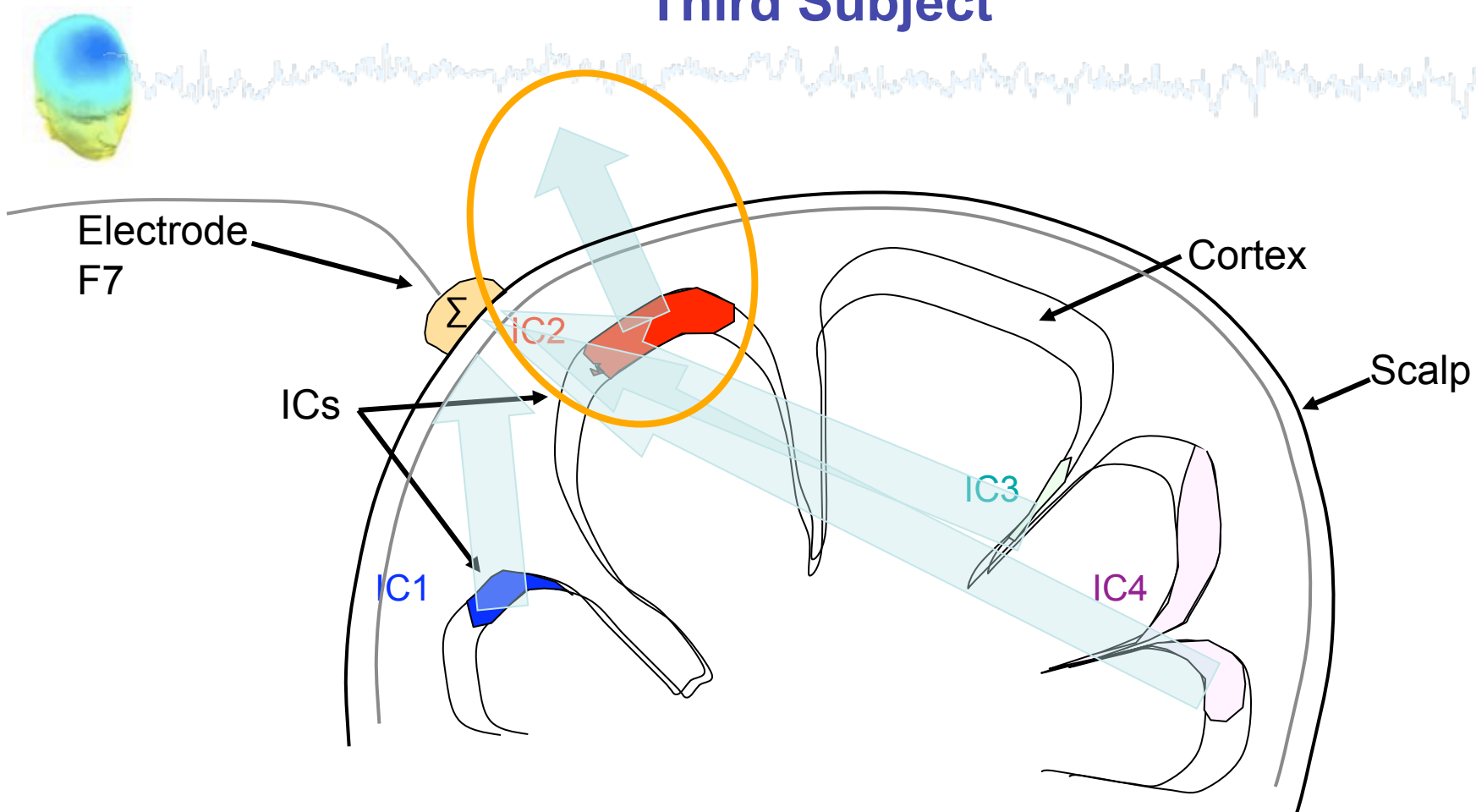
## Example: First Subject



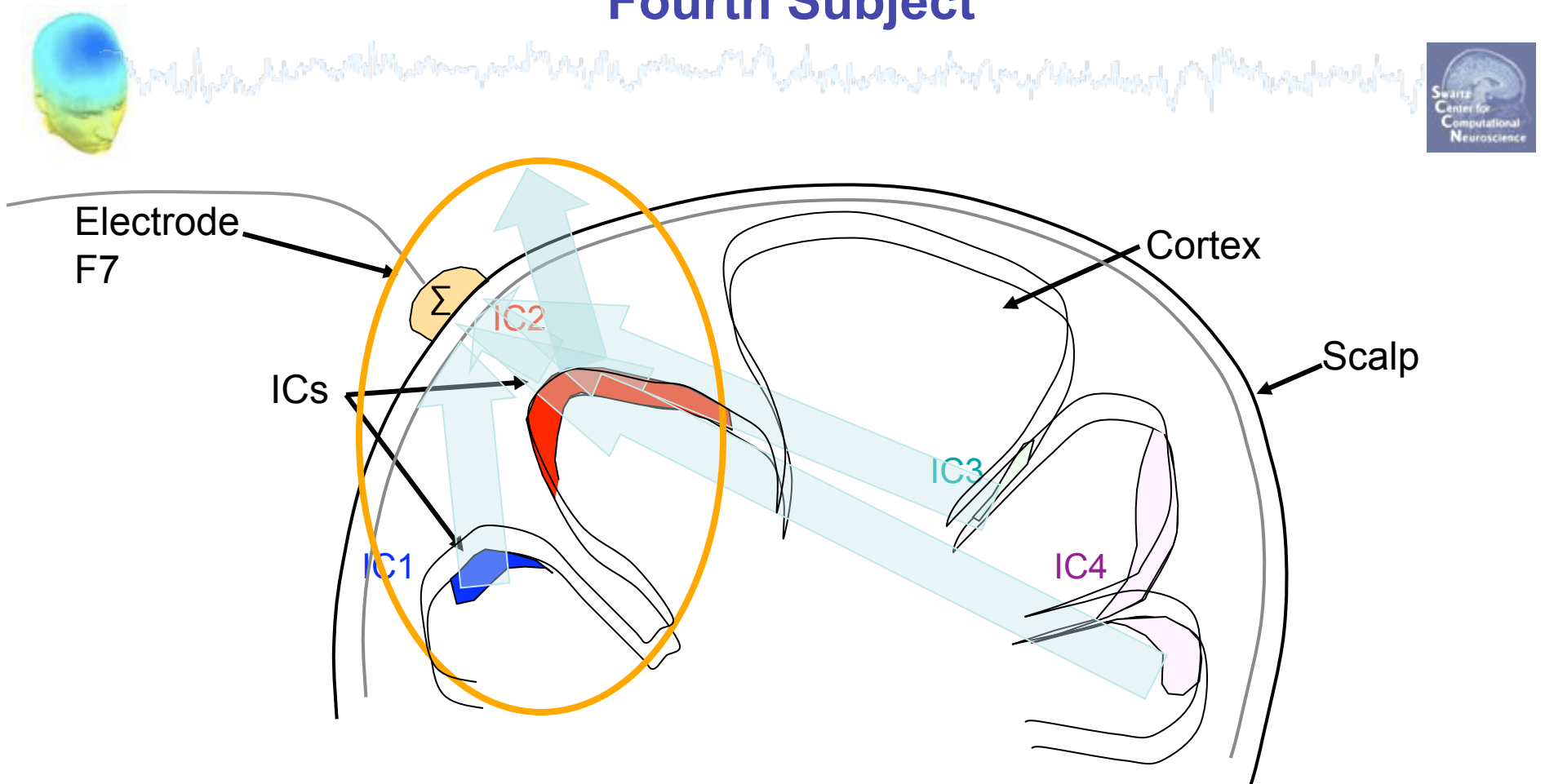
## Second Subject



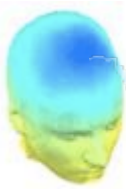
## Third Subject



## Fourth Subject



# How to cluster independent components?



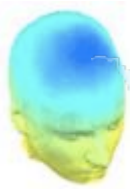
## The same problems hold for clustering independent components

Across Ss, components don't even have "the same" scalp maps!

→ Are "the same" components found across subjects??

- What should define "the same" (i.e., "component equivalence")?
  - Similar scalp maps?
  - Similar cortical or 3-D equivalent dipole locations?
  - Similar activity power spectra?
  - Similar ERPs?
  - Similar ERSPs?
  - Similar ITCs?
  - OR ..., Similar *combinations* of the above? ...





First, we should ask...



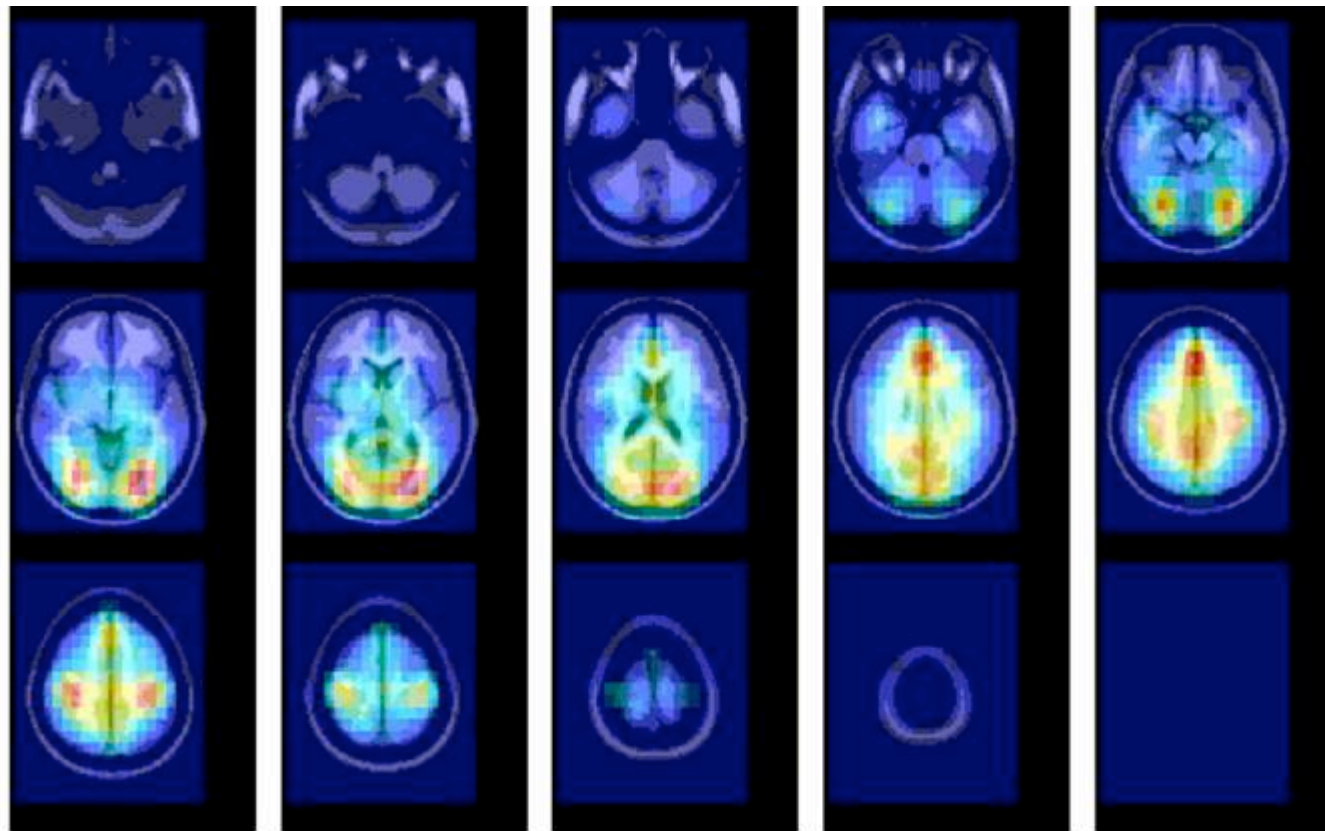
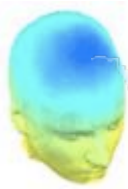
Does the spatial distribution of independent components depend on the task the subject performs?

i.e.

Do “the same” components (and component clusters) appear for every subject task?



# Equivalent dipole density

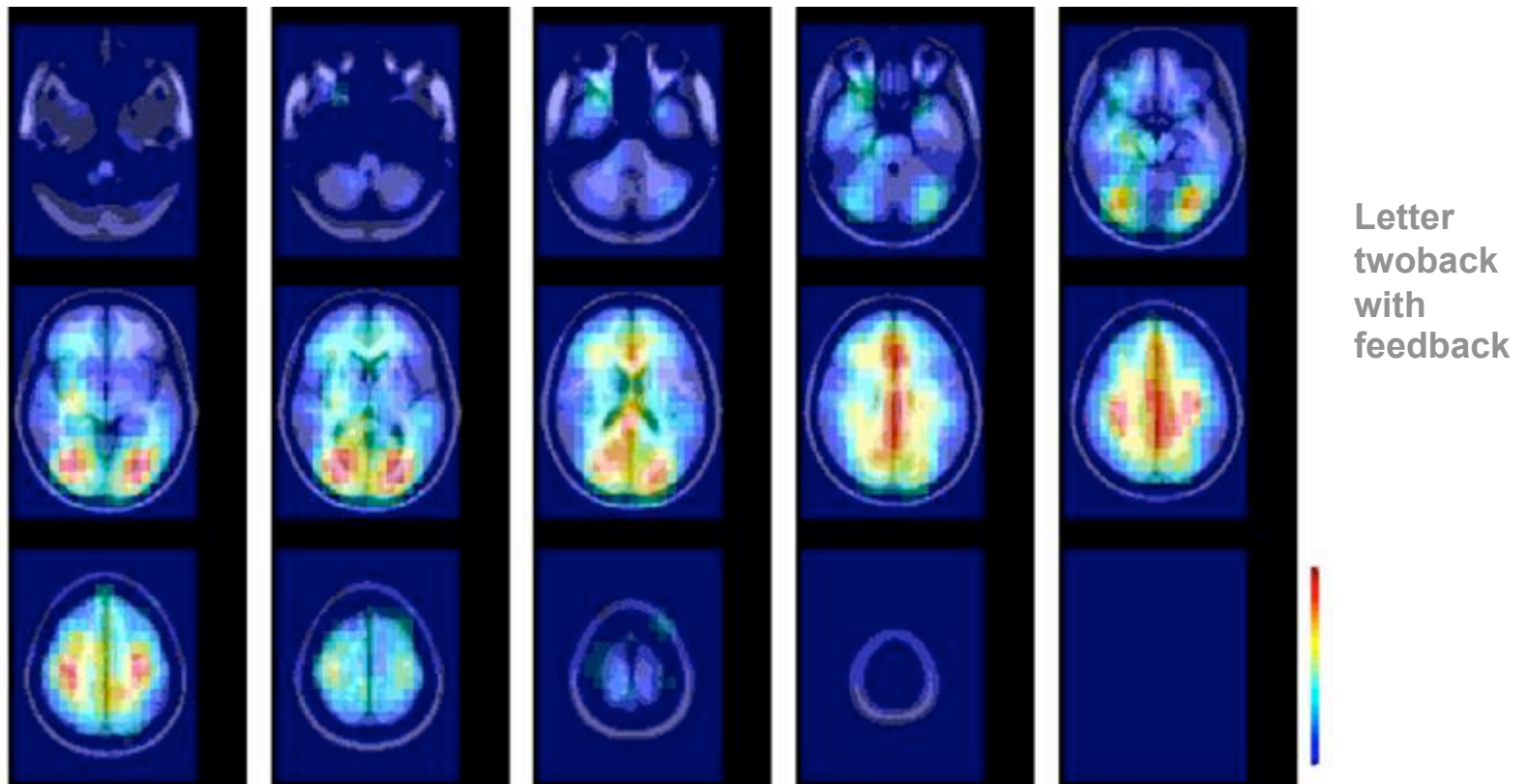
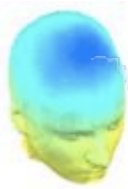


Sternberg  
letter  
memory  
task

>> dipoledensity()



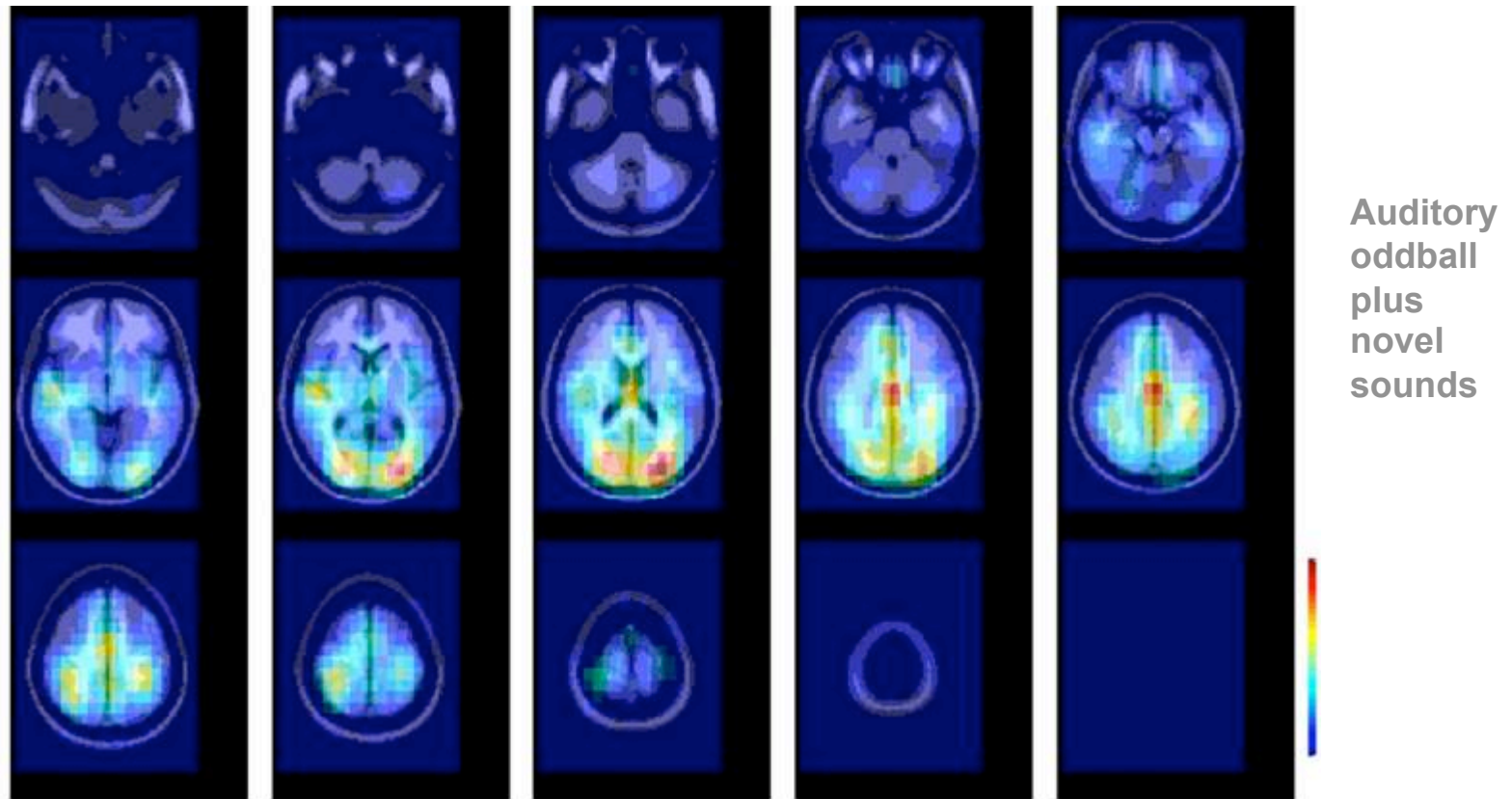
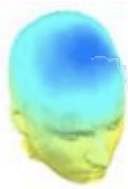
# Equivalent dipole density



>> dipoledensity()



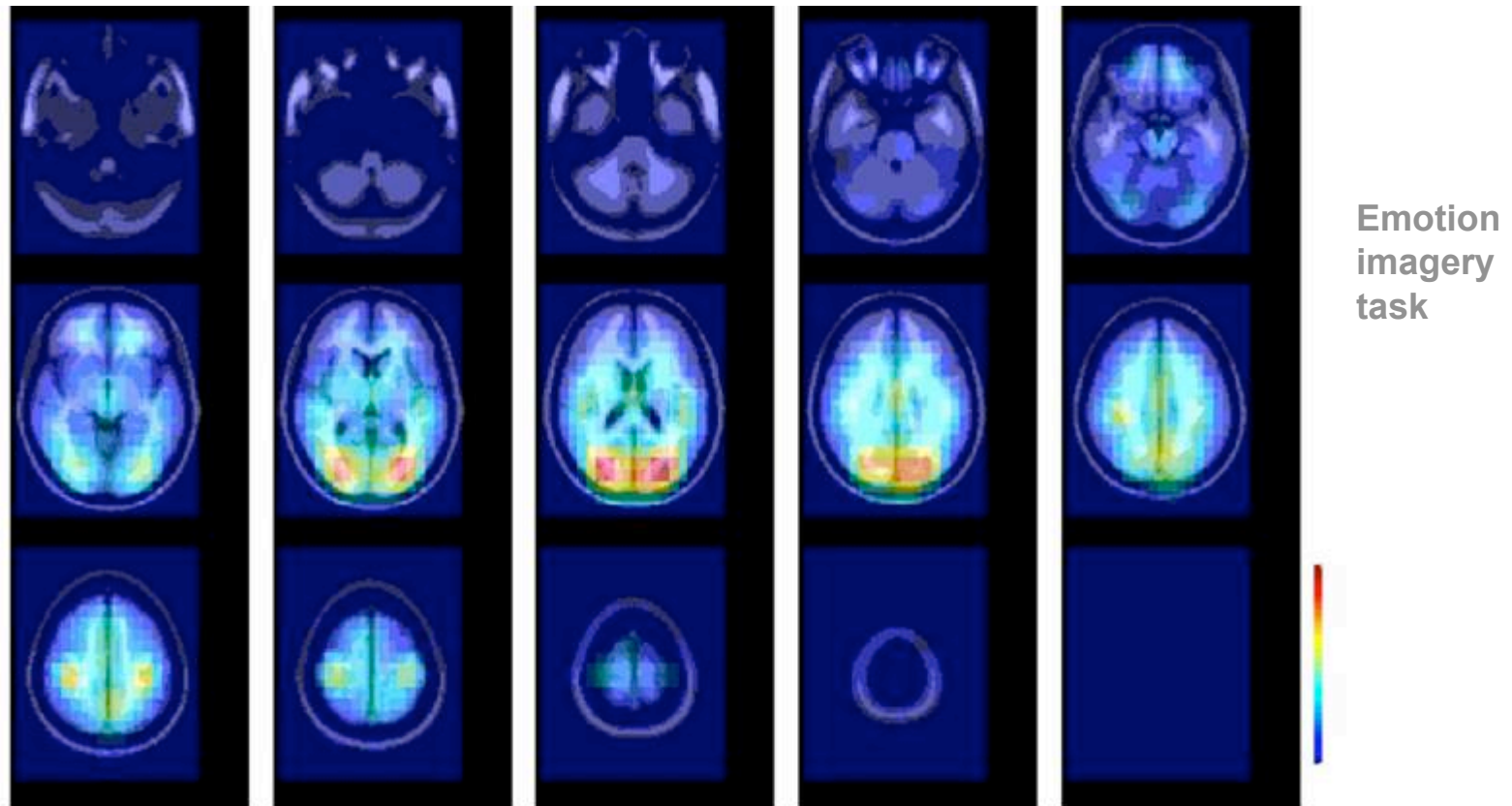
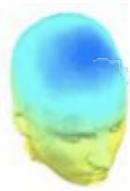
# Equivalent dipole density



>> dipoledensity()



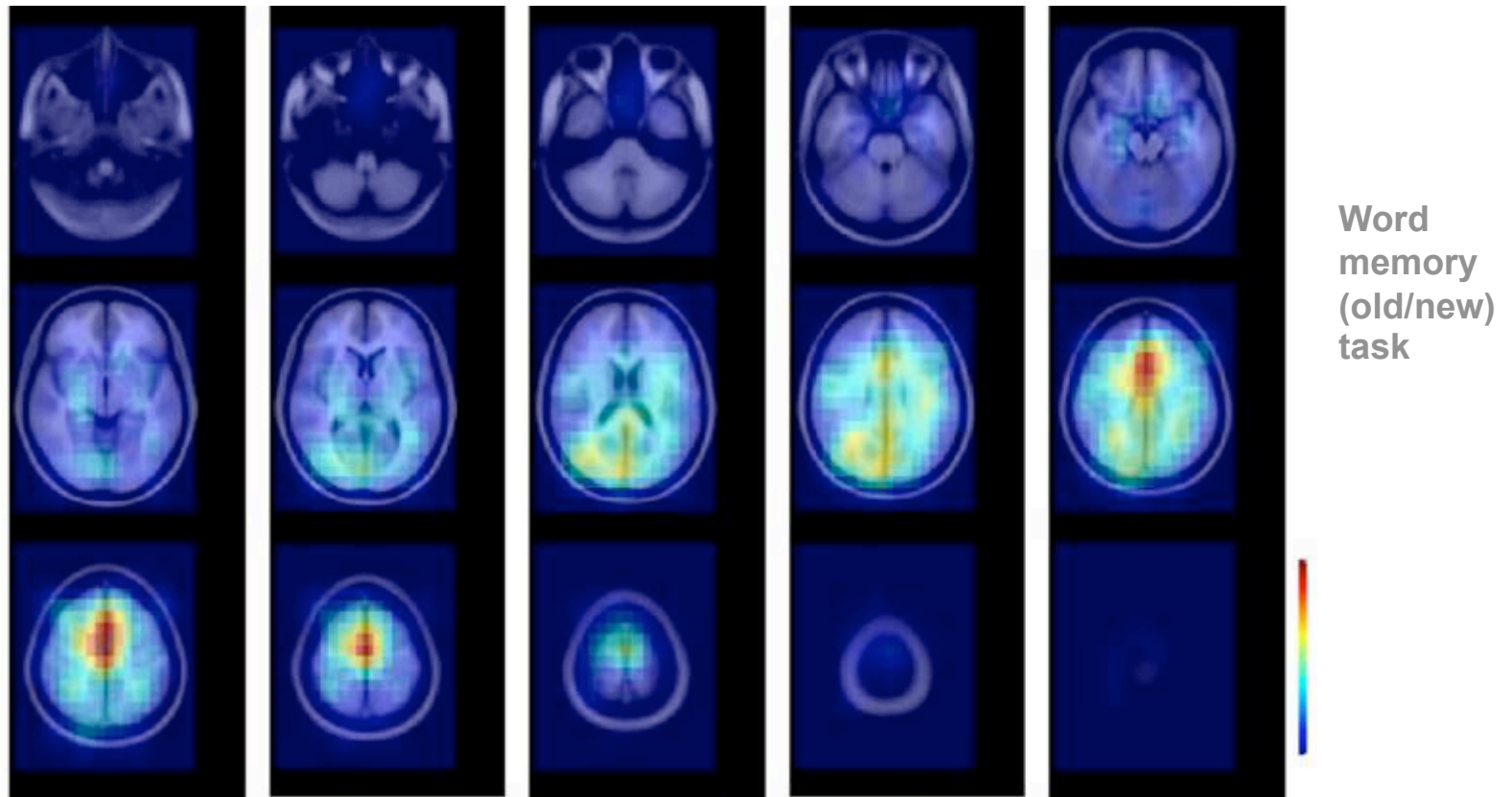
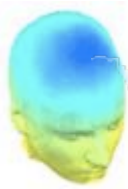
# Equivalent dipole density



>> dipoledensity()



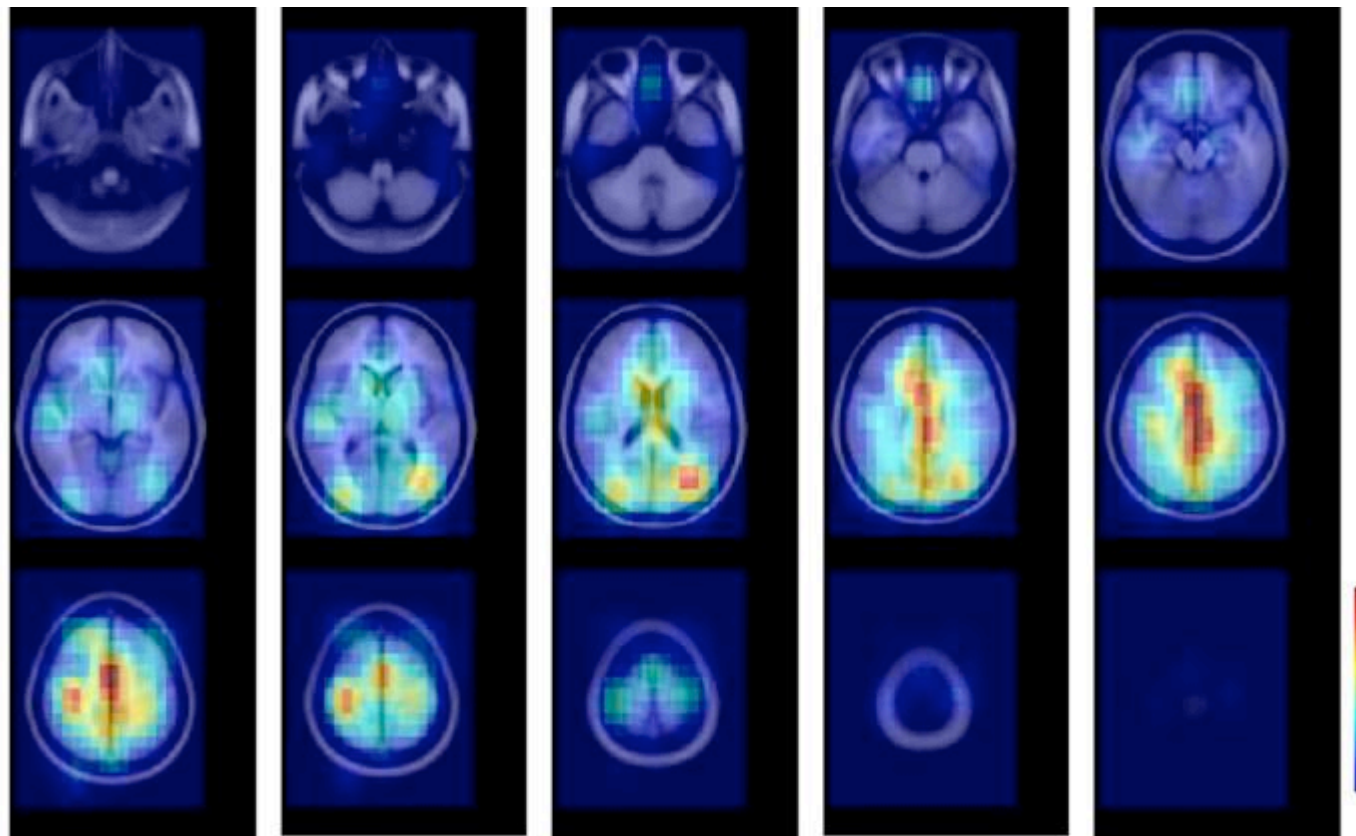
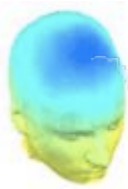
# Equivalent dipole density Exp I



>> dipoledensity()



# Equivalent dipole density Exp II



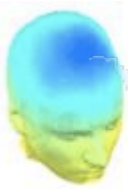
Visually  
cued  
button  
press  
task



>> `dipoledensity()`



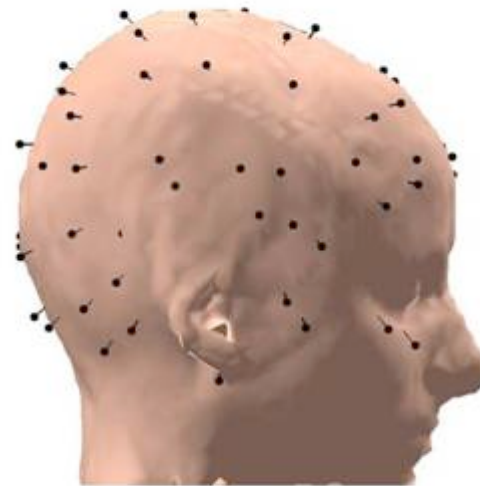
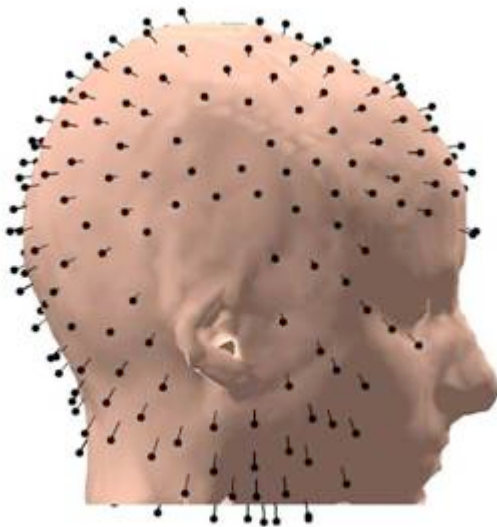
## ... Some caveats



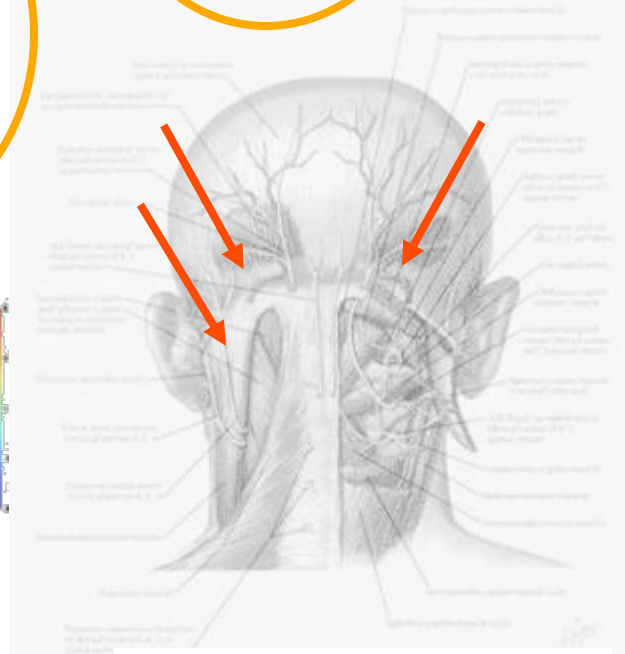
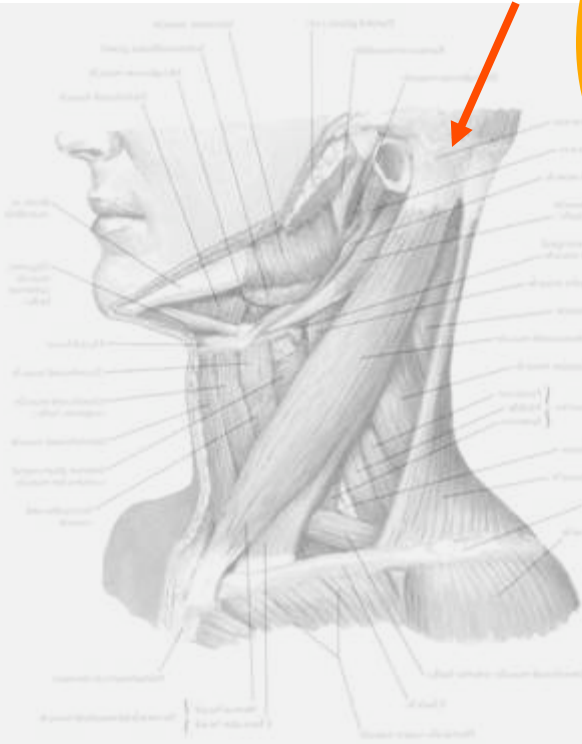
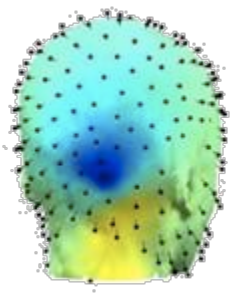
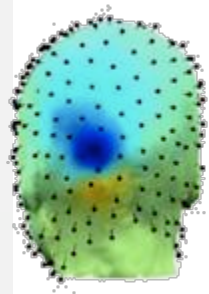
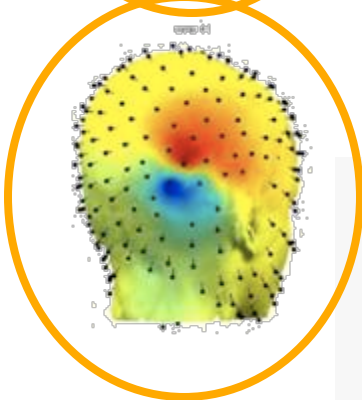
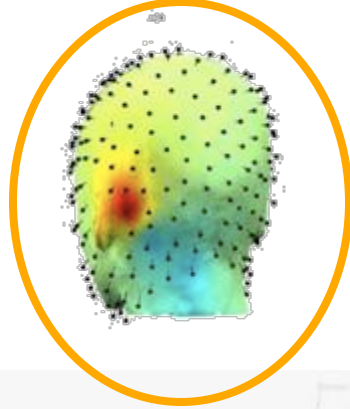
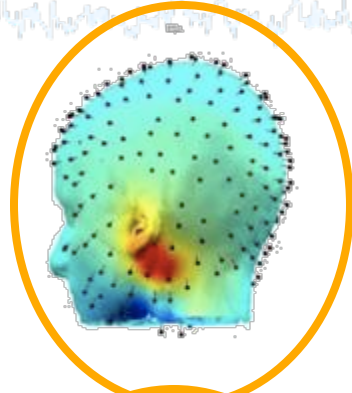
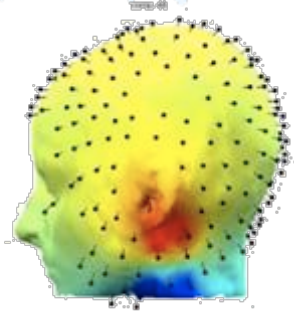
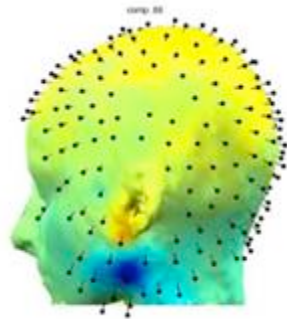
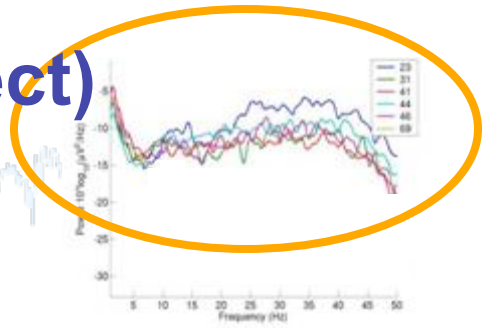
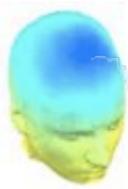
In this preliminary study ...

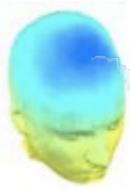


- The electrode locations were not individualized.
- MR images were not available → co-registration crude.
- Single versus dual-dipole model selection was subjective.
- Different electrode montages → possible location effects



# Clustering by spectra (1 subject)





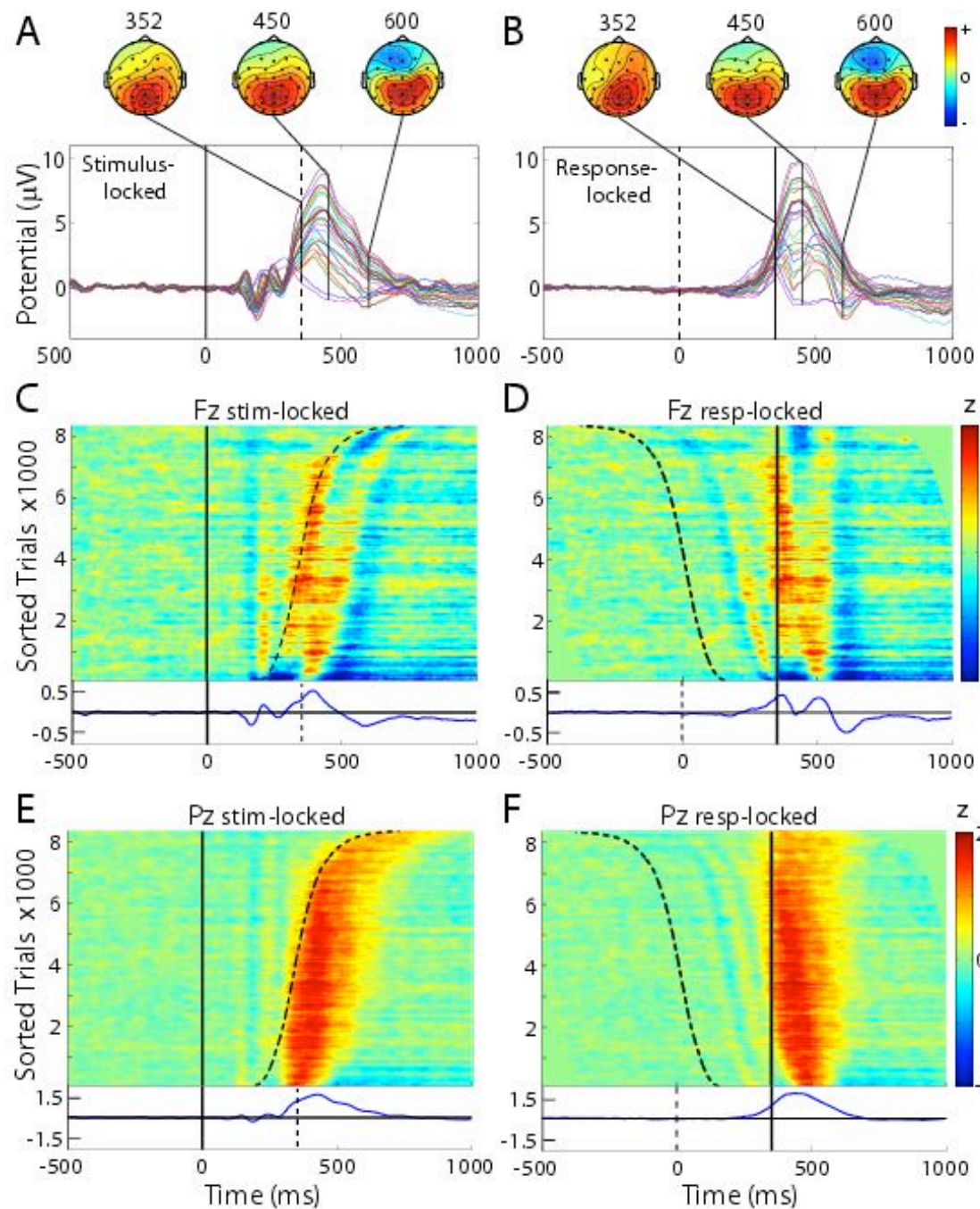
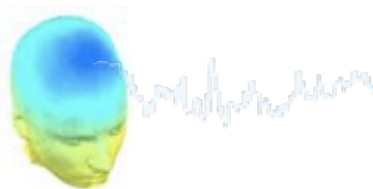
# Visual Selective Attention Task



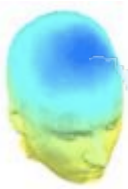
+

**15 subjects**

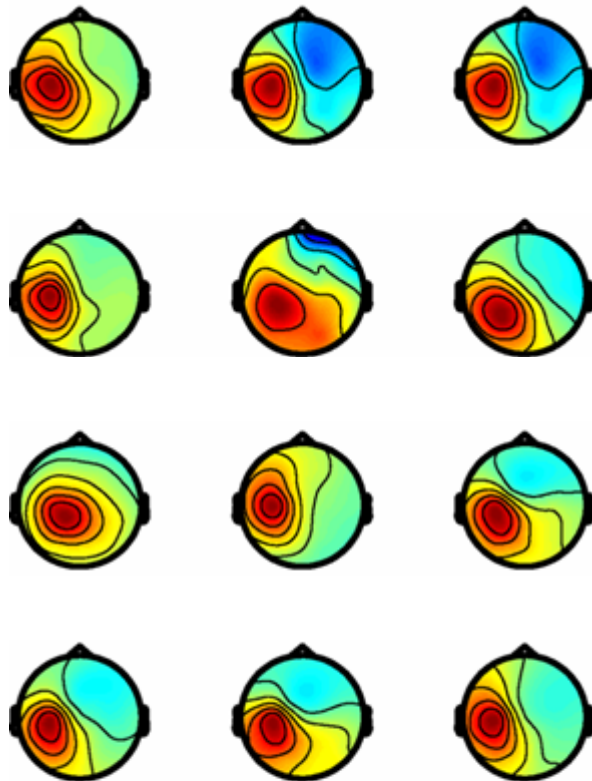




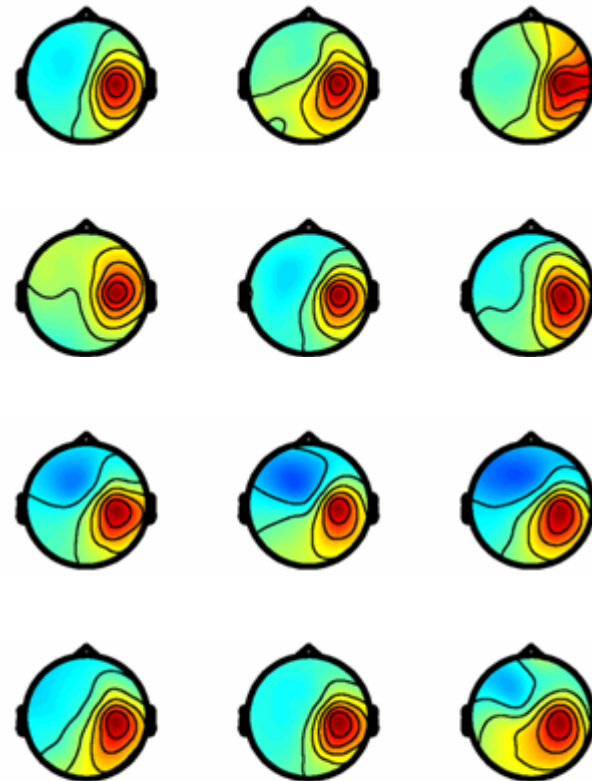
# Clustering ICA components by eye



Left mu



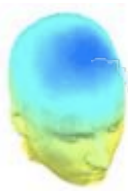
Right mu



EEGLAI

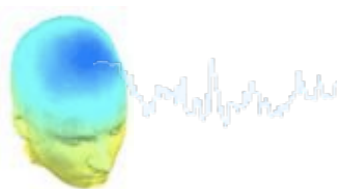
Makeig et al., *submitted*

# Semi-automated clustering

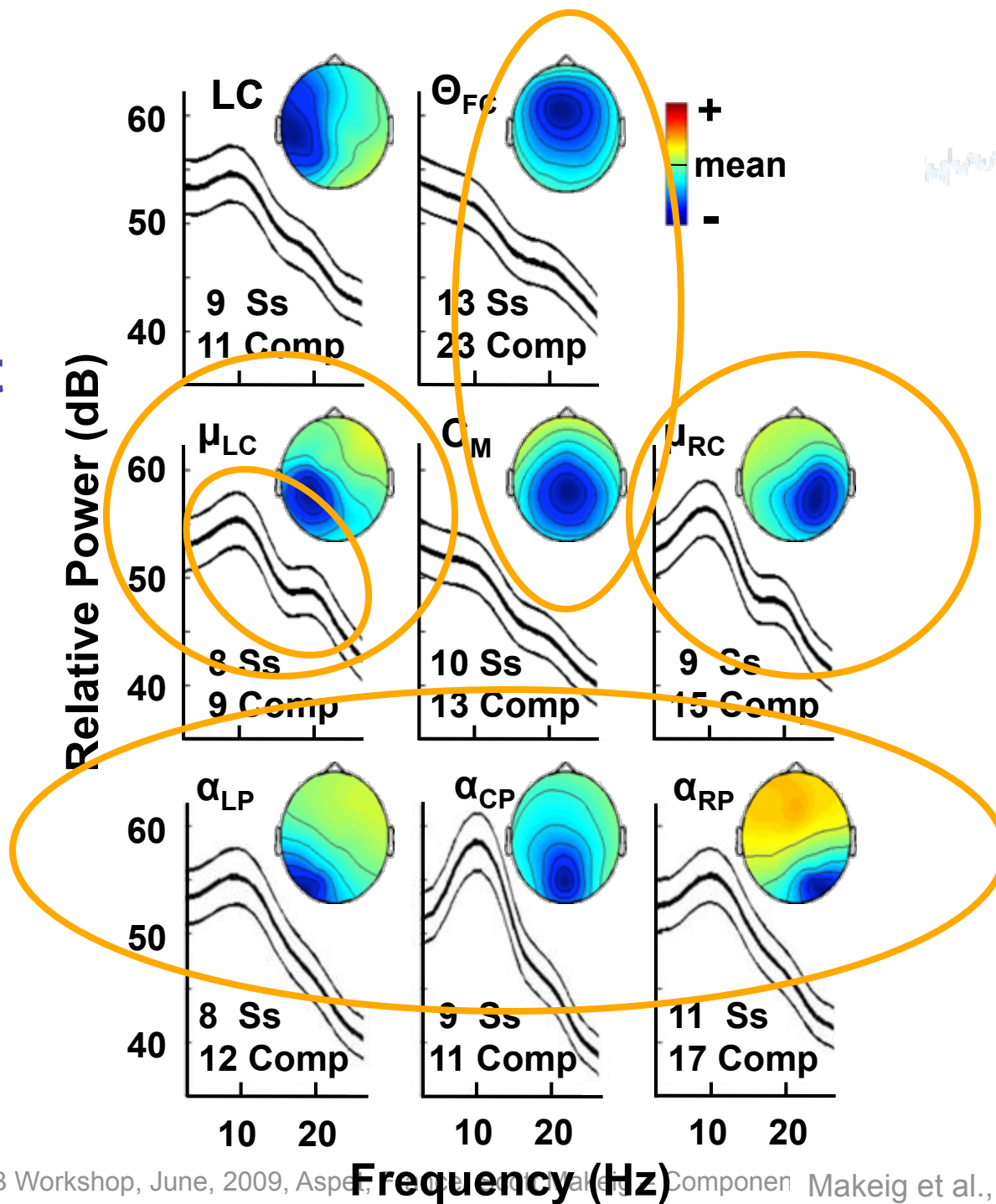


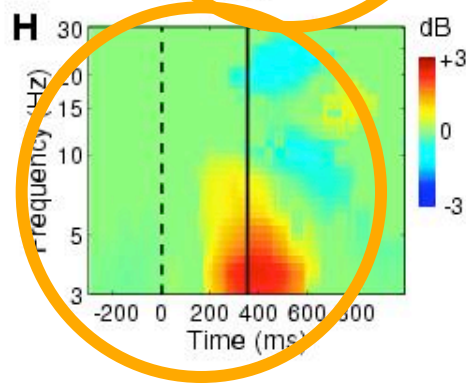
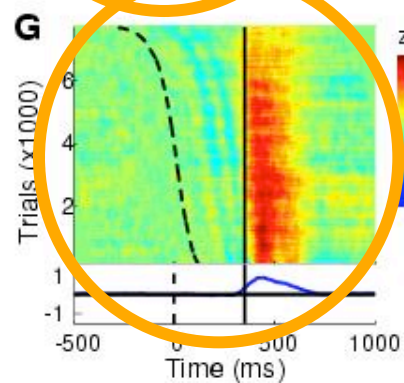
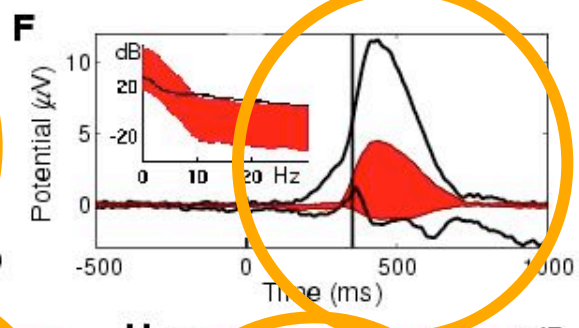
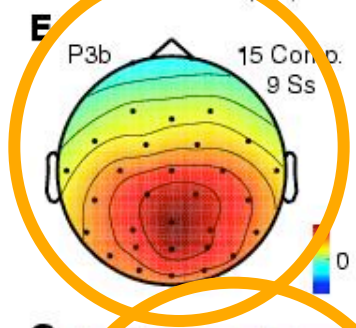
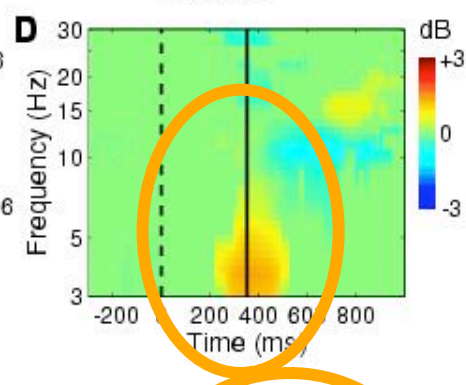
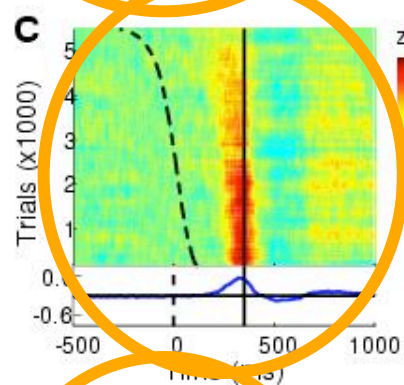
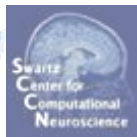
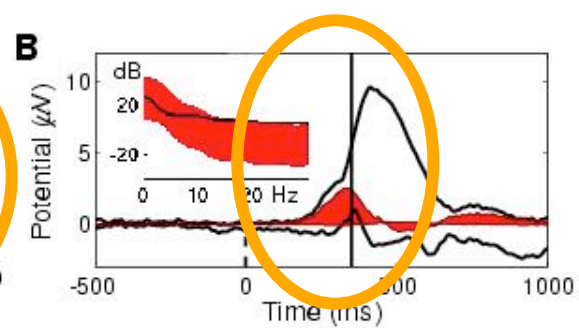
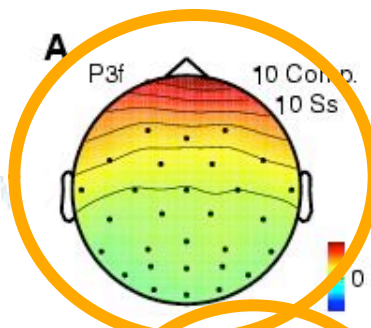
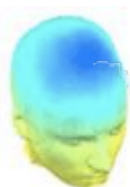
- Clustered components from 15 Ss using a 'component distance metric' incorporating differences between their (weighted) scalp maps, dipole locations, spectra, ERP, ERSP, and ITC patterns.
- Hand-adjusted clusters to remove outliers.
- Determined time/frequency regions of significant ERSP and ITC for each component using permutation-based statistics.
- Used binomial statistics to highlight time/frequency regions significantly active within clusters.

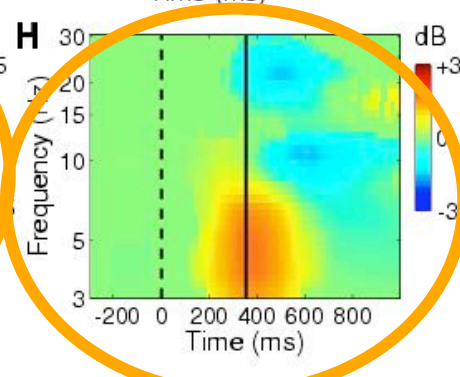
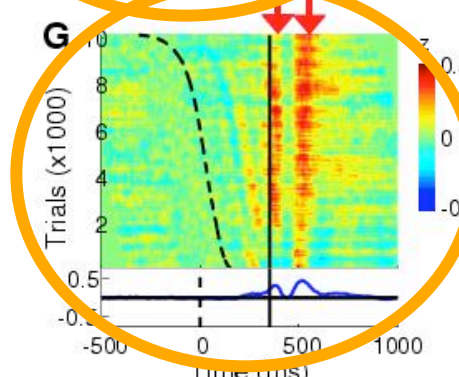
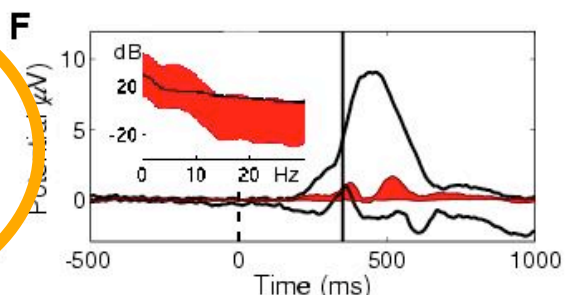
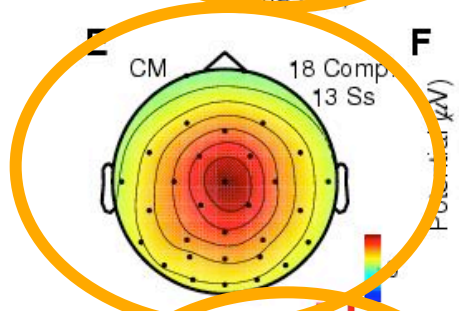
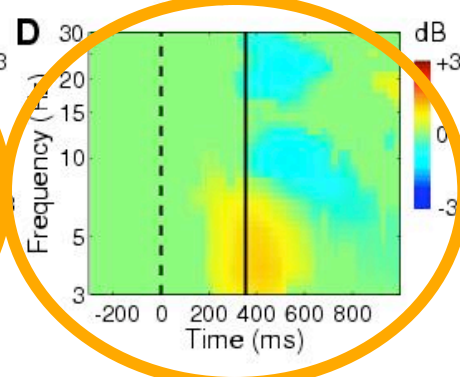
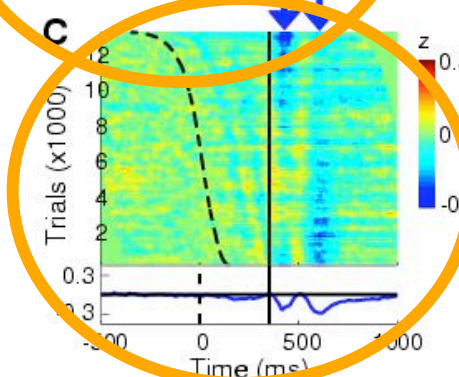
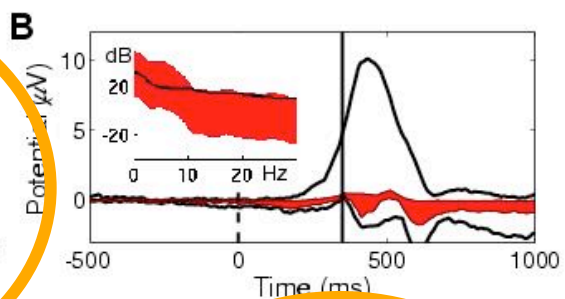
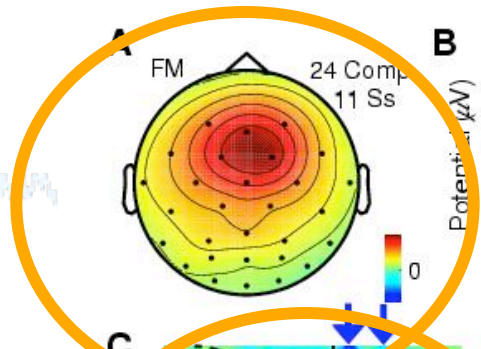
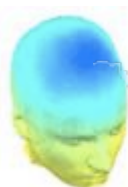


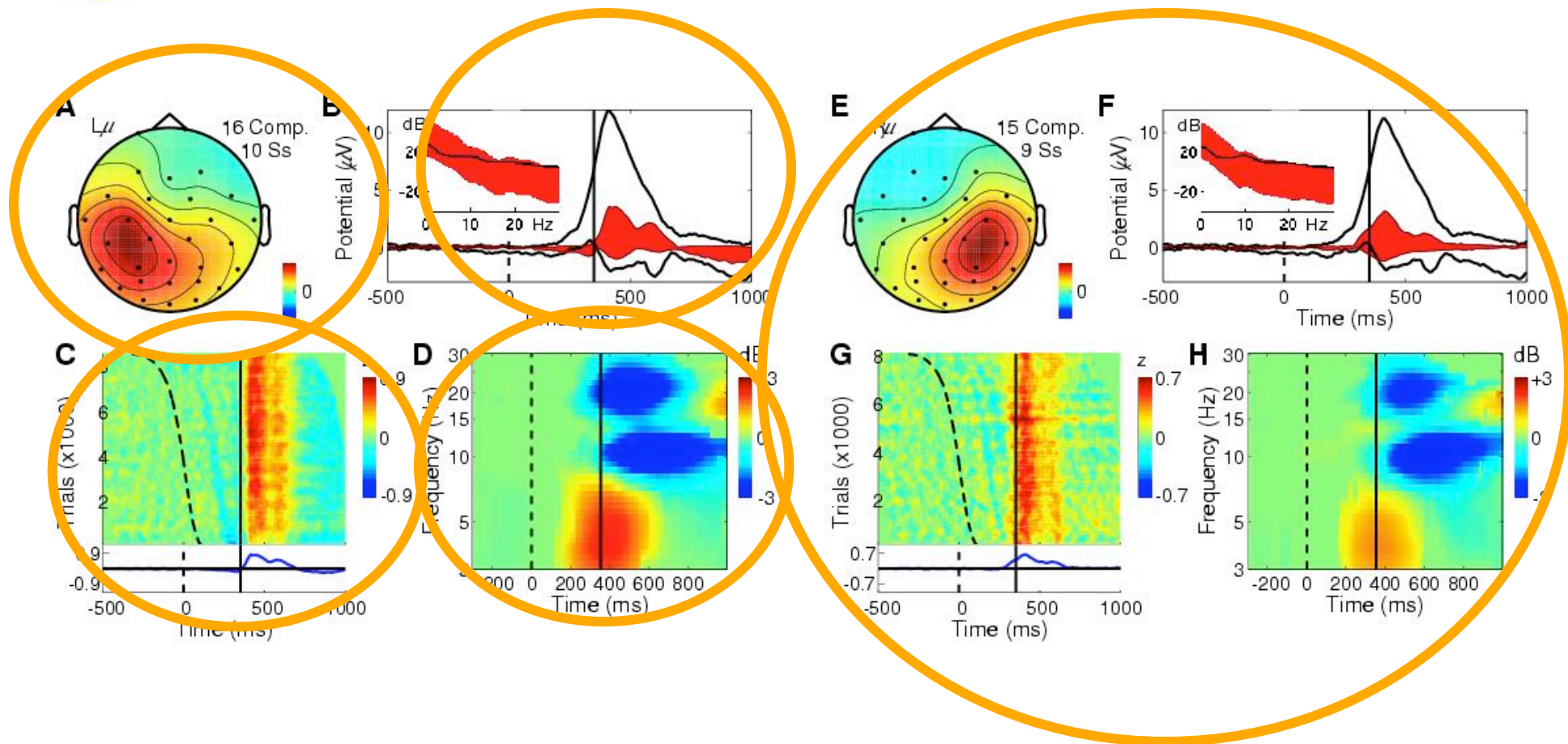
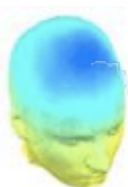


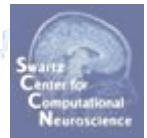
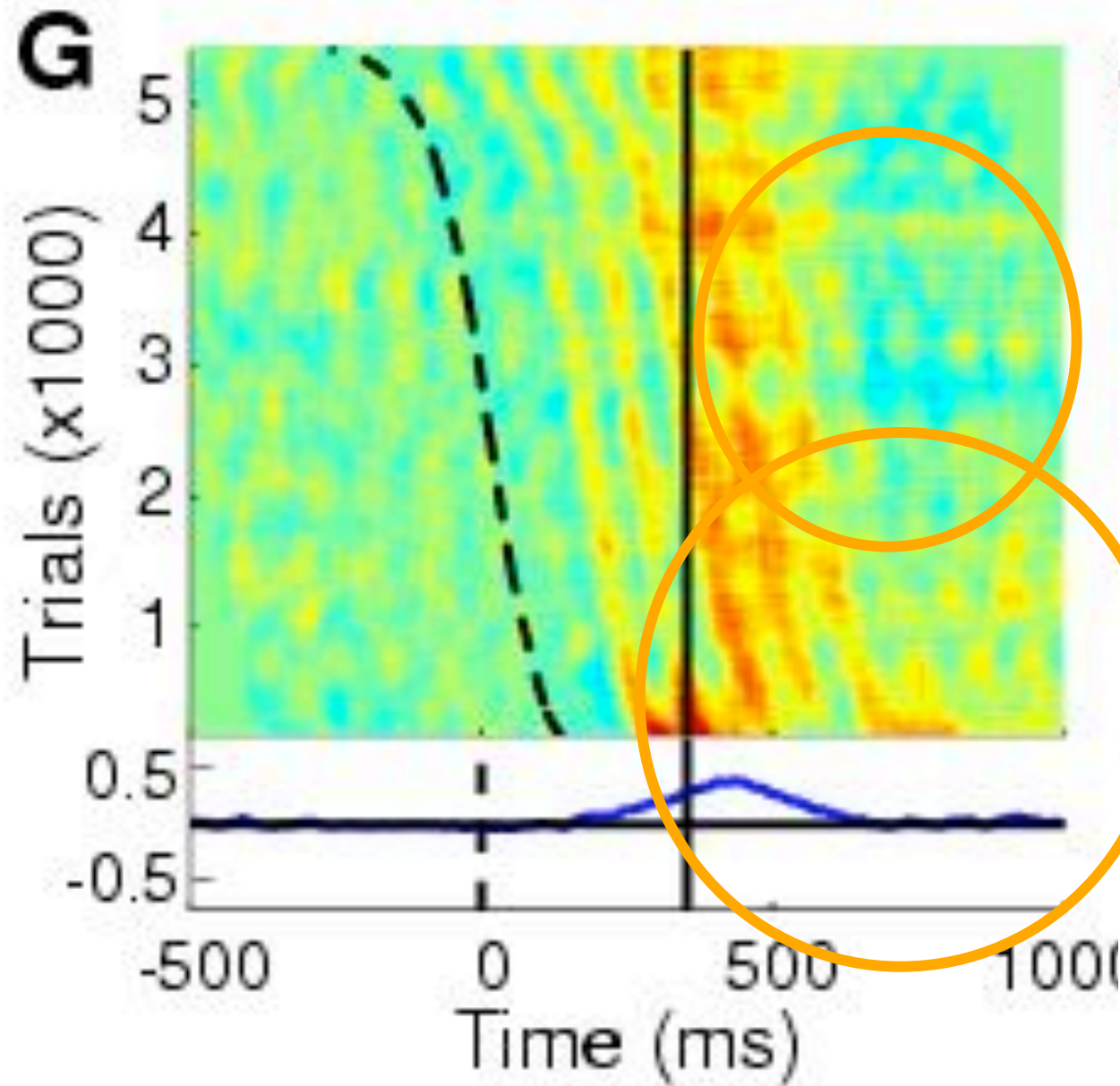
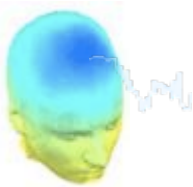
## N1 Study Component Clusters

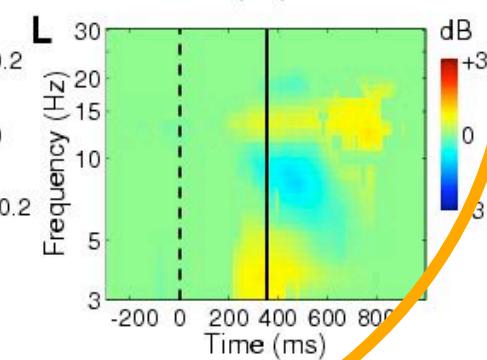
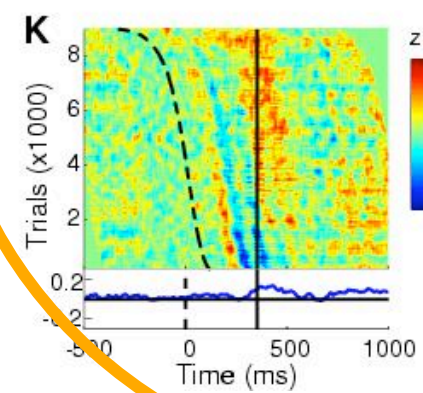
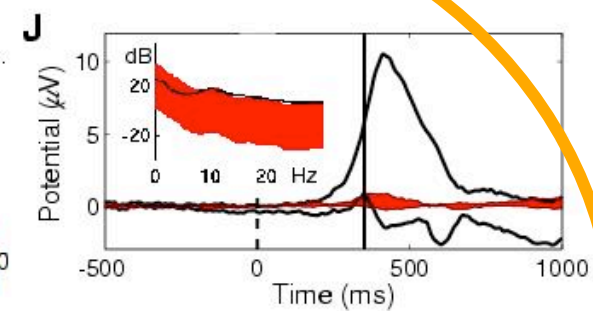
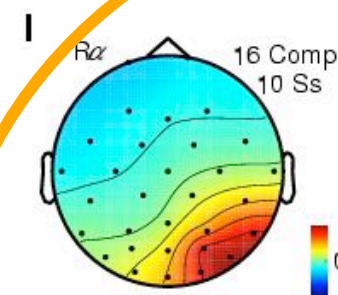
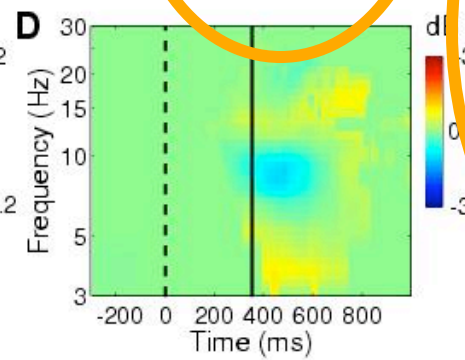
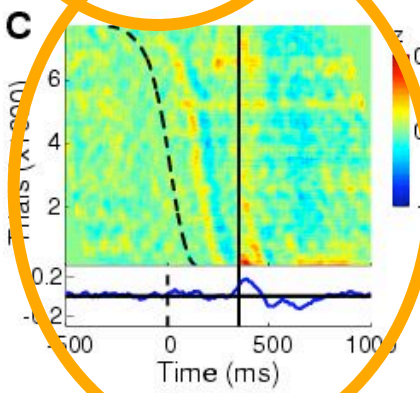
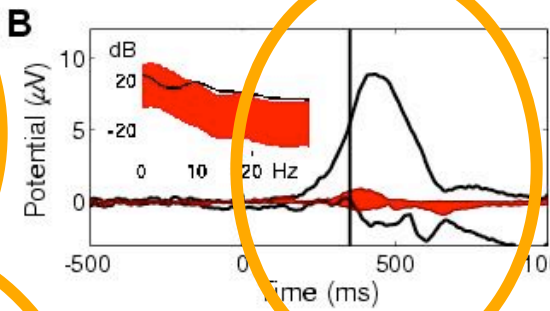
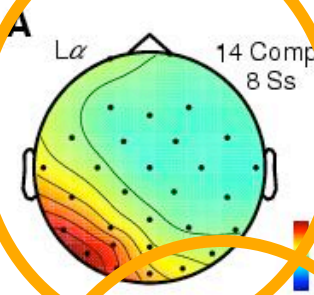
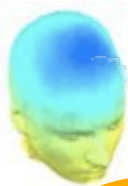




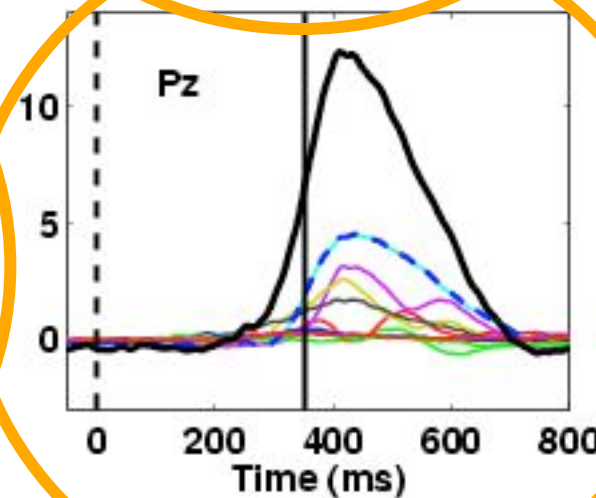
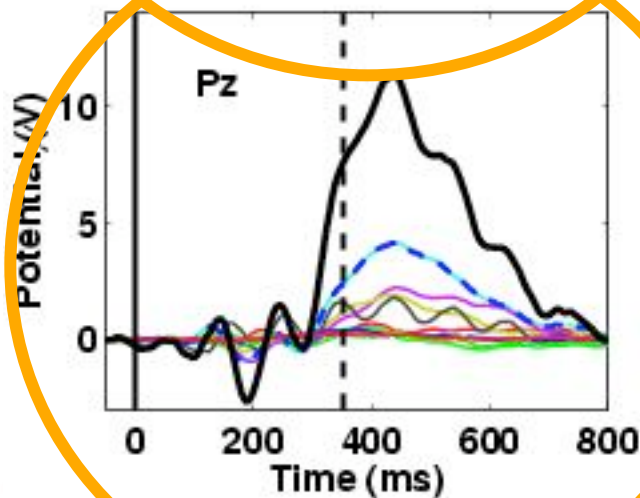
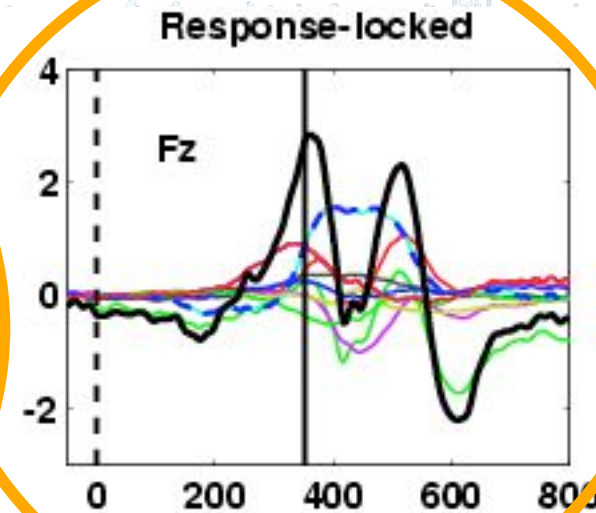
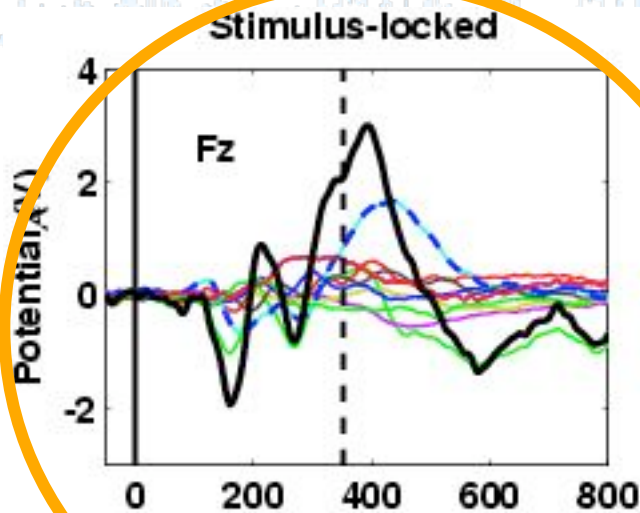
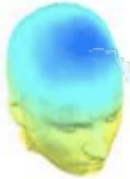




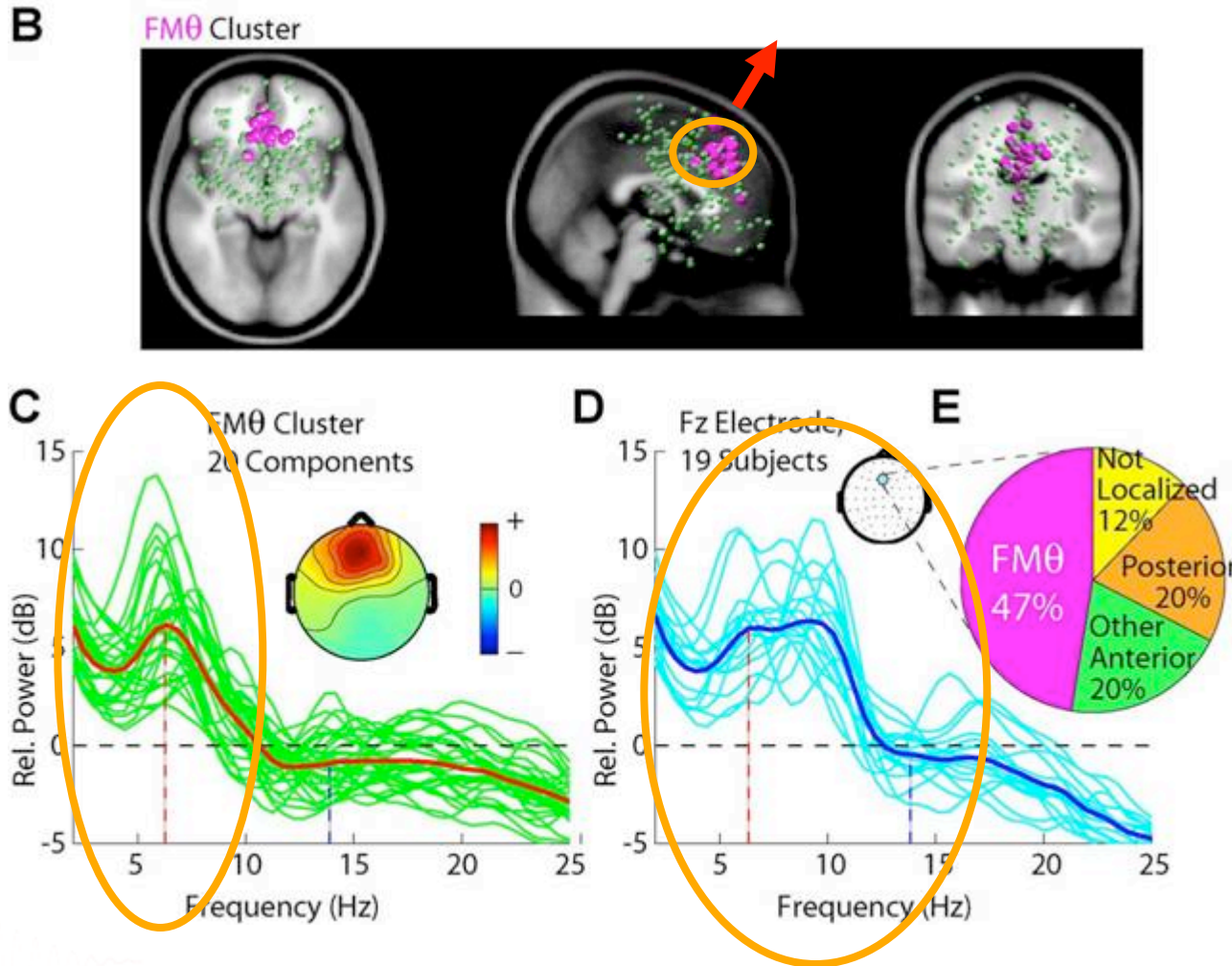
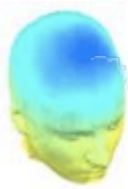


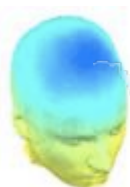


# Complex event-related dynamics underlie 'the' P300

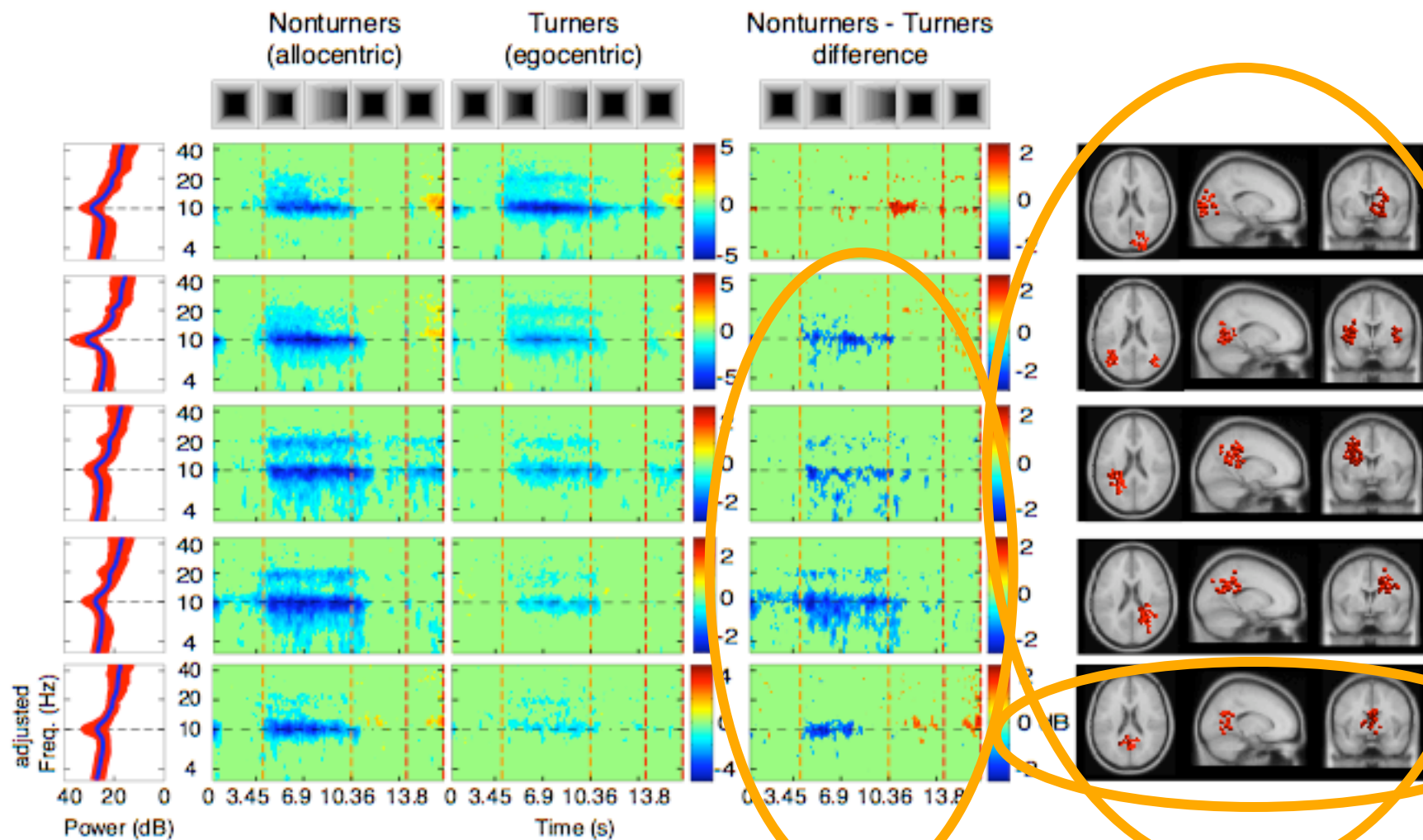


# A FM $\theta$ cluster during working memory

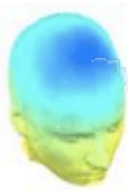




# IC Clusters distinguishing Turners & Nonturners



# Are obtained component clusters “real”?



- **Naïve realism** (a.k.a. “expertise”)

- “Yes! ... because I know one when I see one!”
  - “If it appears where Mu components appear,  
and acts like Mu components act,  
then it IS a Mu component!”

- **Convergent evidence** (a.k.a., “double-checking”)

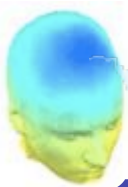
- Two possible approaches:
  - Cluster on PLACE → Check ACTIVITY consistency (re task)
  - Cluster on ACTIVITY → Check PLACE consistency

- **Absolute truth:**

- More ideal forward and inverse models
- Invasive multiscale recordings + modeling

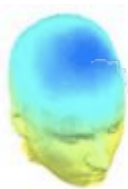


# EEGLAB clustering procedure

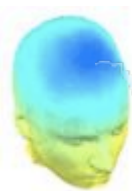


1. Identify a set of datasets as an EEGLAB **study** or '**studysset**'.
2. Specify the subject **group**, **subject** code, **condition** and **session** of each dataset in the study.
3. Identify **components to cluster** in each study dataset.
4. Decide on **component measures** to use in clustering the study and/or to evaluate the obtained component clusters.
5. Compute the component measures for each study dataset.
6. **Cluster the components** on these component measures.
7. Review the obtained **clusters** (e.g., their scalp maps, dipoles, and activity measures).
8. **Edit the clusters** (manually remove/shift components, make sub-clusters, merge clusters, re-cluster).
9. Perform **signal processing** within or between selected clusters.

# EEGLAB Clustering strategy



1. Cluster on **multiple measures** (dipole locations, scalp maps, spectra, ERPs, ITCs, ERSPs) in **one or more conditions**.
2. **Reduce the dimension** of each measure to a principal component subspace.
3. Compose a PCA-reduced **position vector** for each component.
4. **Cluster** the composed component vectors using **k-means** or other.
5. Use the computed component measures (not PCA-reduced) to **visualize the activities and spatial properties** of the clustered components.
6. Compute and visualize the **cluster-mean measures**.
7. Use the **clustered study set data** as input into **std\_** functions.



# Need ICs from all subjects be included in each cluster?



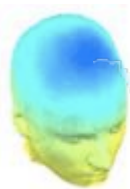
**Not all subjects contribute components to each cluster.**

## Why not?

- Different numbers of artifact components ( $\sim$ INR)
- Subject differences!?
- Is my subject group a Gaussian cloud??
  - 'subject space'

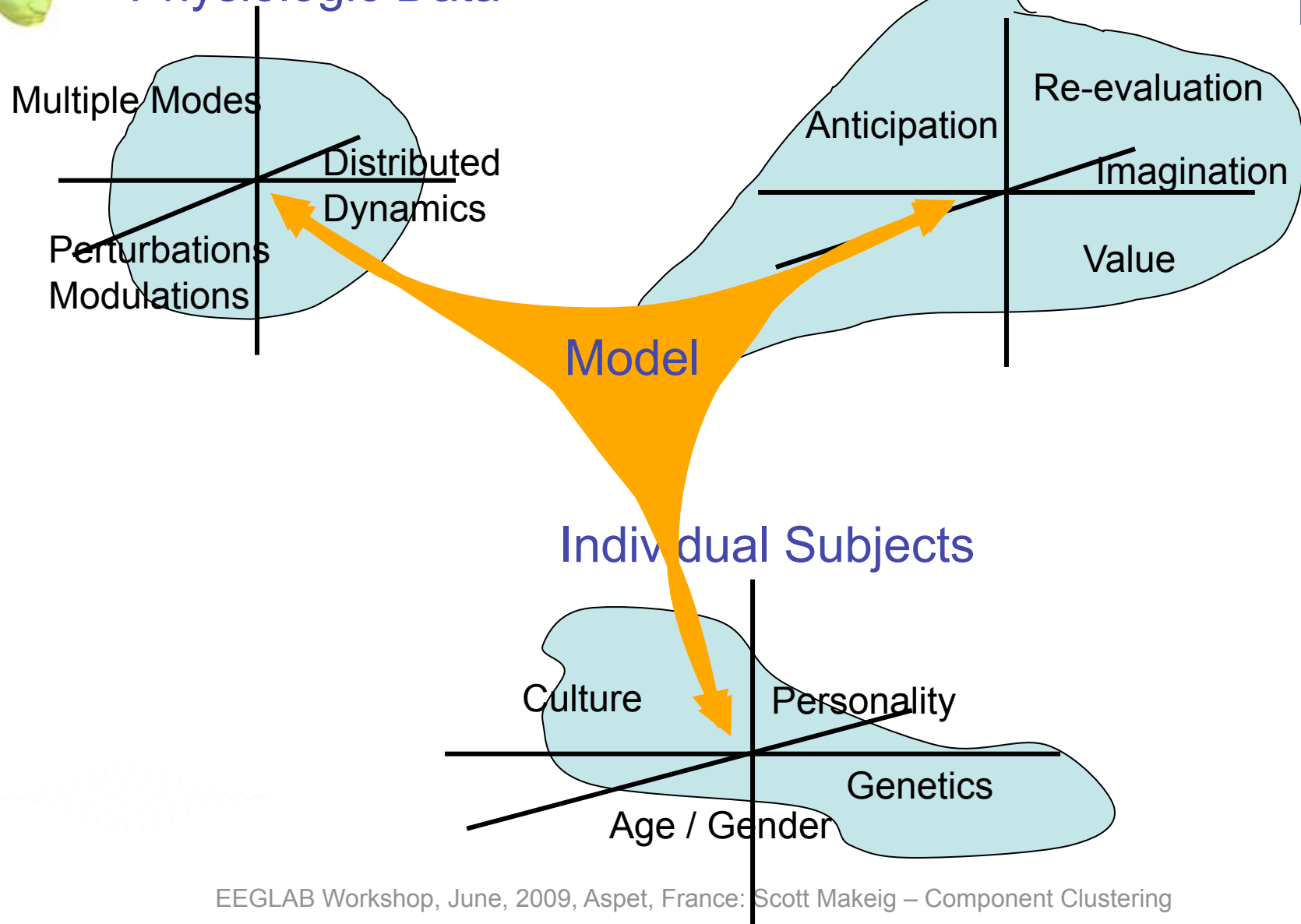


# Beyond Clustering

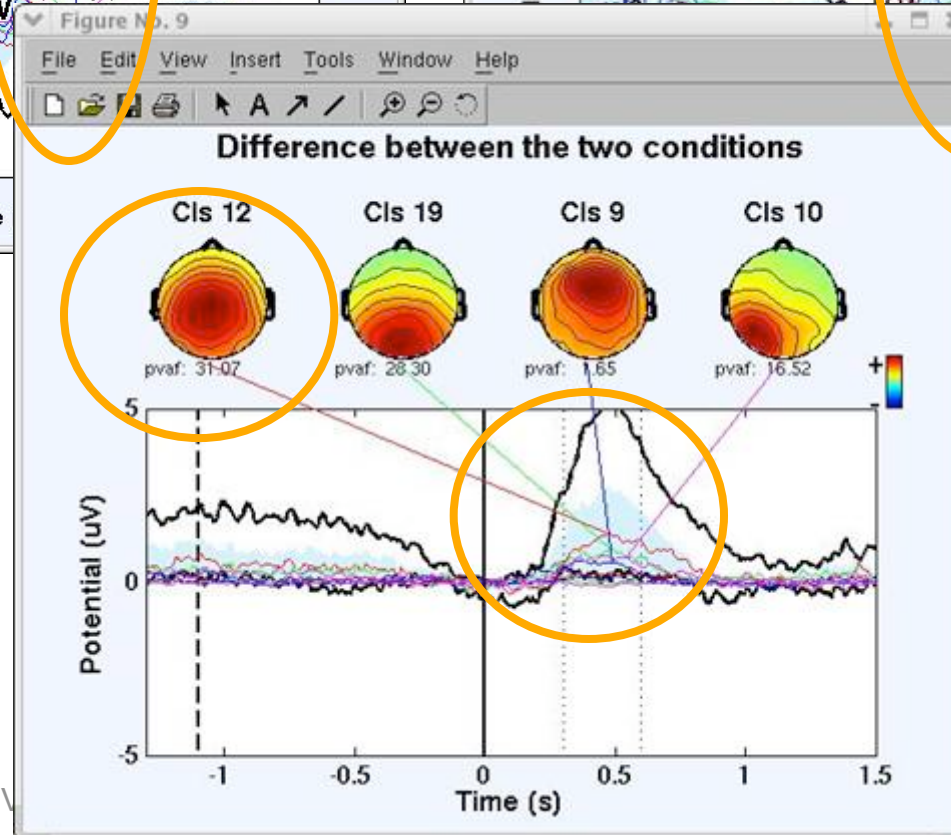
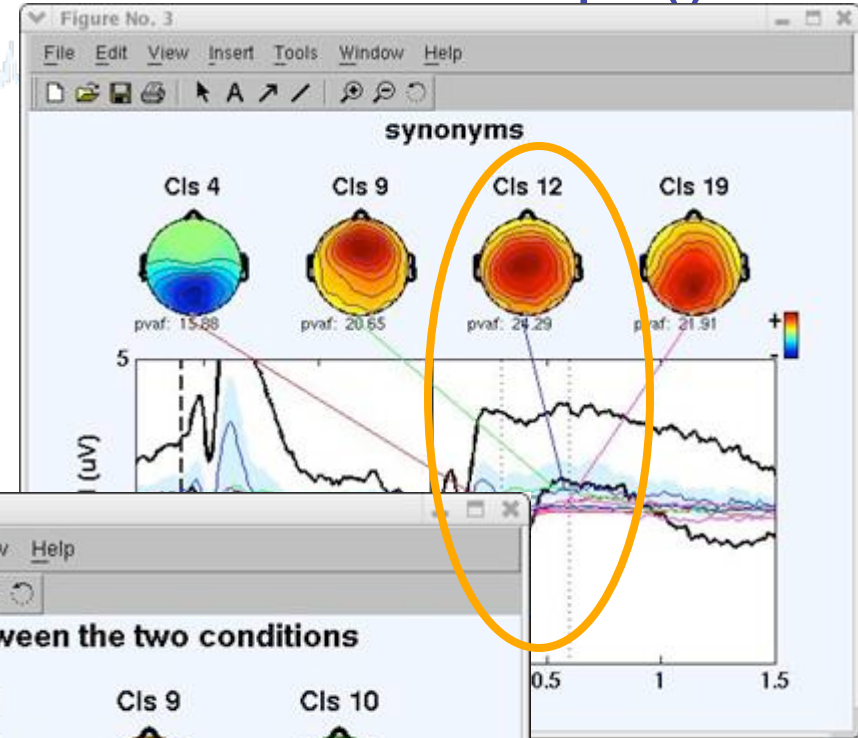
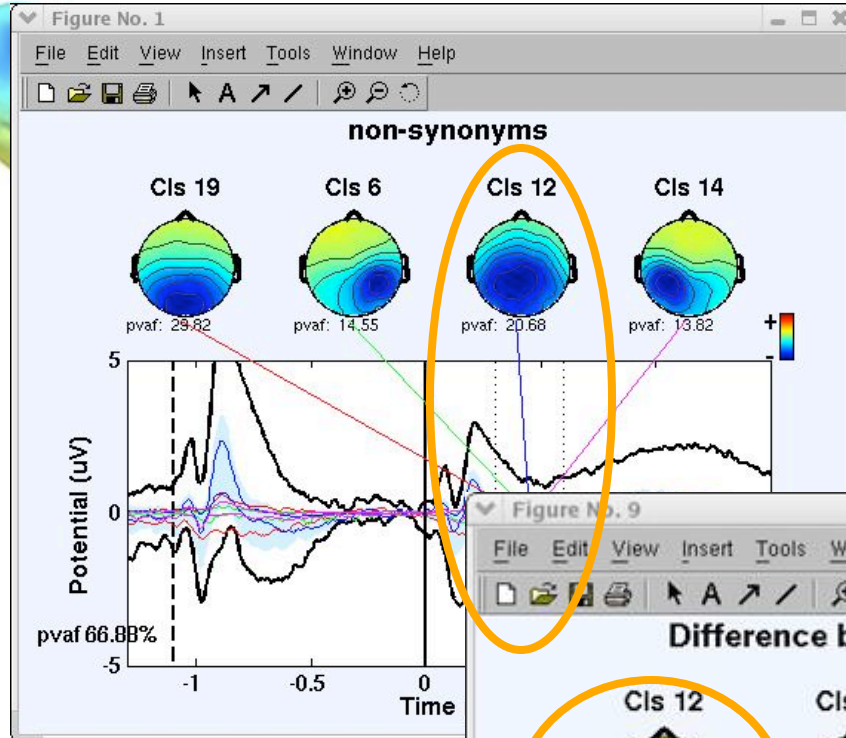


Physiologic Data

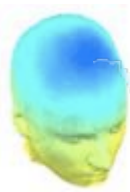
Cognitive Events



# Cluster ERP contributions - clust\_envtopo()



```
clust_envtopo(STUDY, ALEEG,
'clusters', [], 'subclus', [3 7 18 20],
'env_erp', 'all', 'vert', -1100,
'baseline', [-200 0], 'diff', [2 1],
'limits', [-1300 1500 -5 5],
'only_precomp', 'on', 'clustnums',
-4, 'limcontrib', [300 600])
```



# Component clustering research issues



## Research Issues:

- Alternative clustering methods (soon)
- Clustering goodness-of-fit
- Cluster plotting details
- Add new pre-clustering measures
- Study subject differences!?

## Development Plan:

- New EEGLAB 'STUDY.DESIGN' definition:
  - Vector of conditions → Uncoupled experimental STUDY.DESIGN structures, each with a vector of included conditions.
- More flexible and complete STUDY.DESIGN statistics