# DIPFIT and model co-registration



Co-register electrodes with model
Autofft, plot dipoles, fine fit
3D headplot co-registration

# Finding dipole locations using DIPFIT in EEGLAB

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		Grand average datasets					Au	tofit	(coarse fit, fine fit & plot)
		Locate dipoles using LORETA					Plo	t cor	mponent dipoles
		PCA plugin							



#### **Co-register to model**

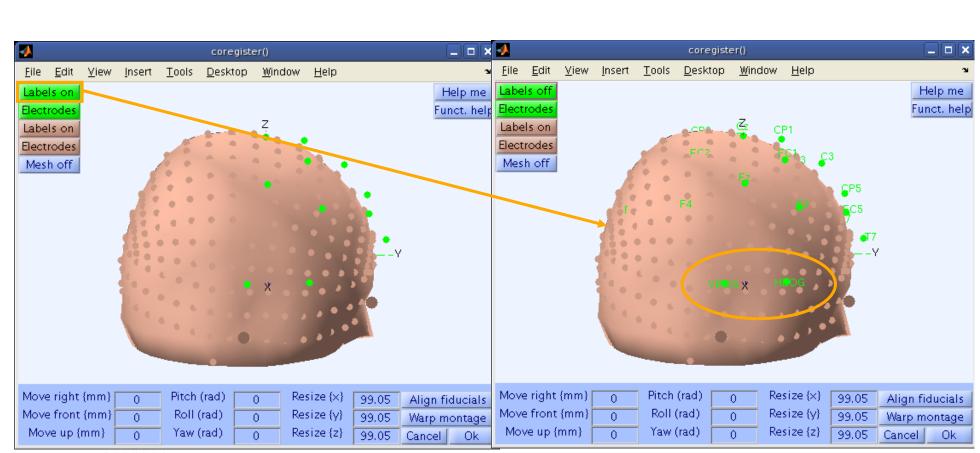


Head model (click to select)	Spherical Four–Shell (BESA) Boundary Element Model (MNI) CTF MEG Custom model files		
Head model file Ourput coordinates	glab/plugins/dipfit2.2/standard_BEM/standard_vol.mat MNI	Browse Click to select	Help
MRI file	lab/plugins/dipfit2.2/standard_BEM/standard_mri.mat	Browse	Help
Model template channel locations file	lugins/dipfit2.2/standard_BEM/elec/standard_1005.elc	Browse	Help
Co–register chan. locs. with head model		Manual Co-Reg.	_No Co−Reg.
Channels to omit from dipole fitting		List	
Note: For EEG, check that the channel locat (To do this: 'Set head radius' to about 85 in			

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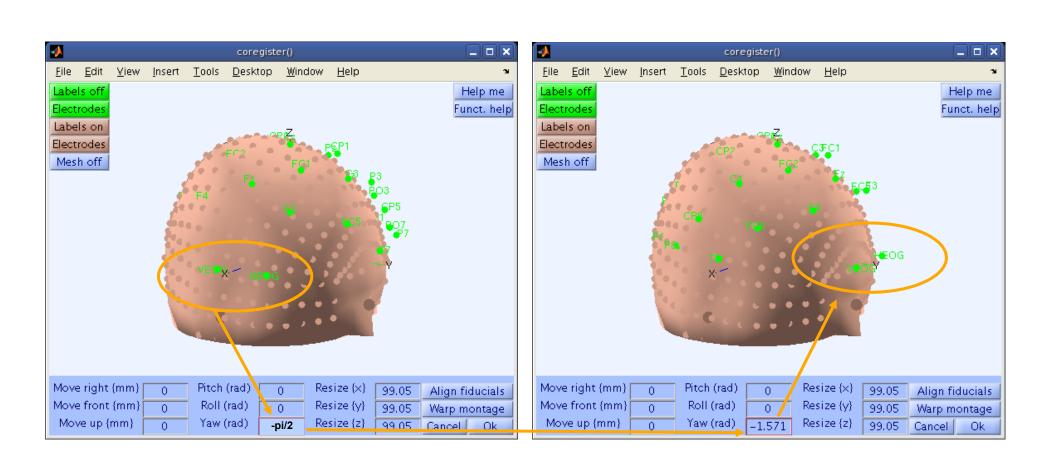
# Co-register to model, cont'd

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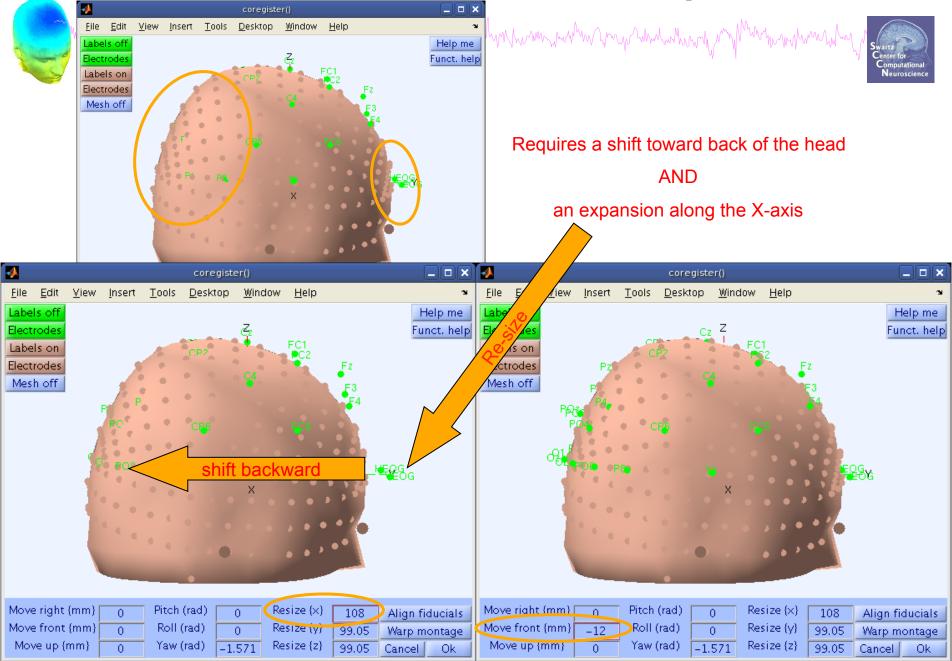


# Perform translation of electrode positions

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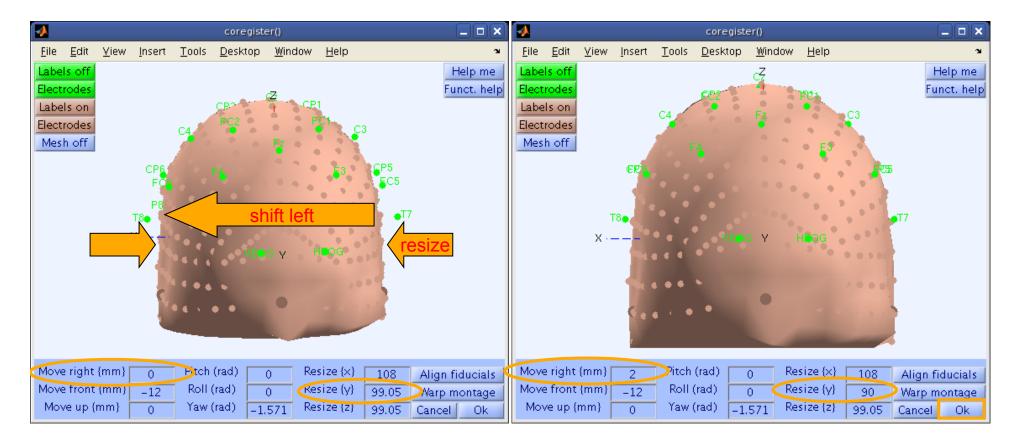
#### Perform translation of electrode positions



# Perform translation of electrode positions



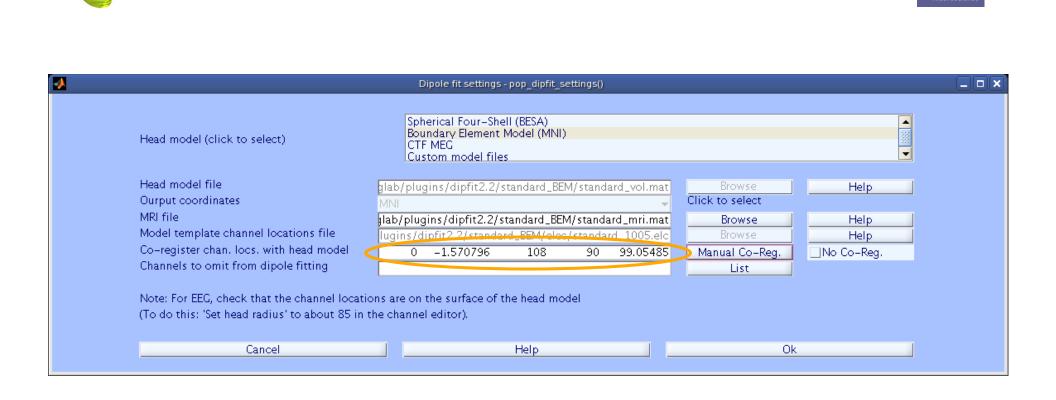






#### **Confirm electrode transformation**

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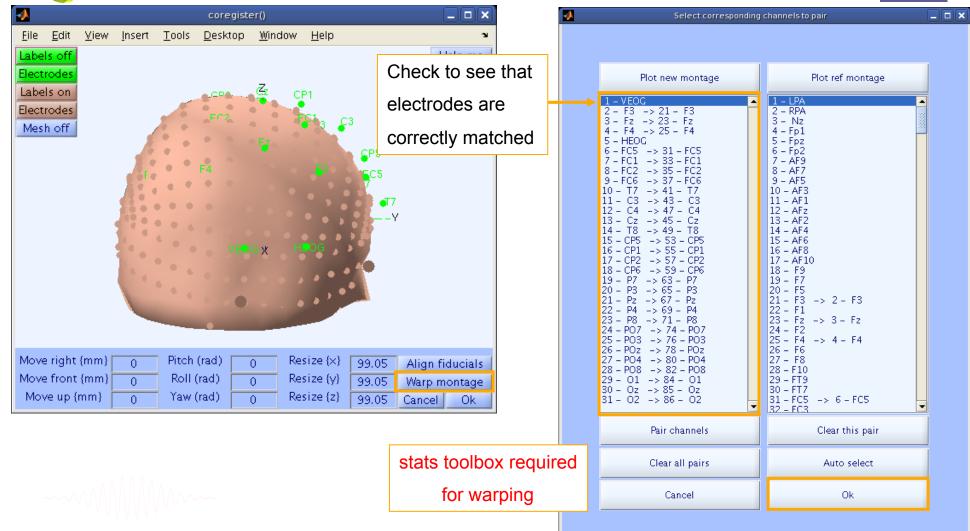


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#### Alternatively, warp to standard montage

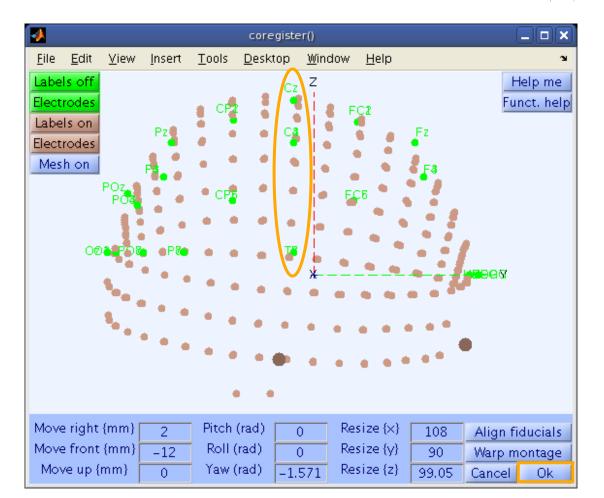
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# **Check coregistration with model**





#### **EEG.dipfit structure**

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hdmfile:	[1x76 char]
mrifile:	[1x71 char]
chanfile:	[1x83 char]
chansel:	[1x33 double]
coordformat:	'spherical'
model:	[1x33 struct]
current:	32
vol:	[1x1 struct]
coord_transform:	$[0 \ 0 \ -1.570796 \ 100 \ 76 \ 90.87264 \ 1 \ 1 \ 1]$

From head model transformations

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# DIPFIT and model co-registration

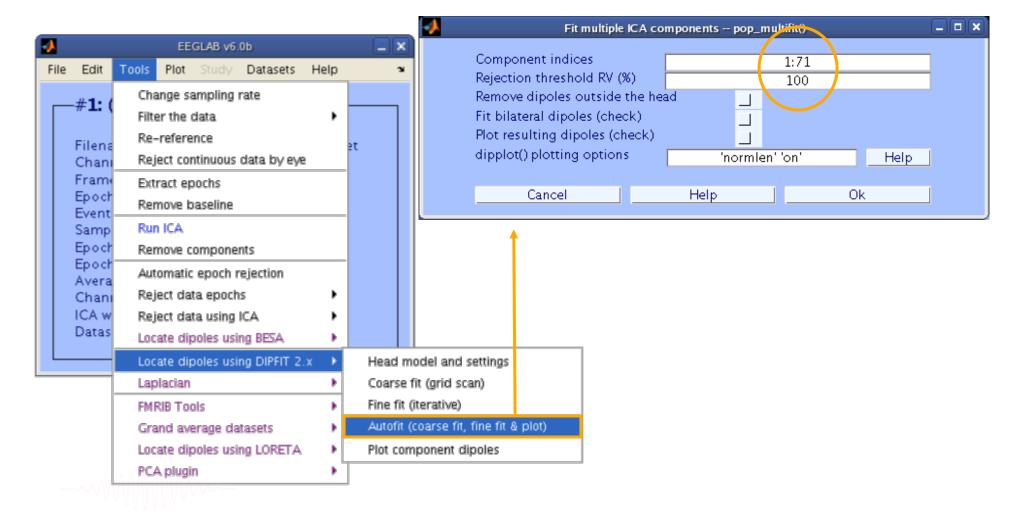


Co-register electrodes with model
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3D headplot co-registration

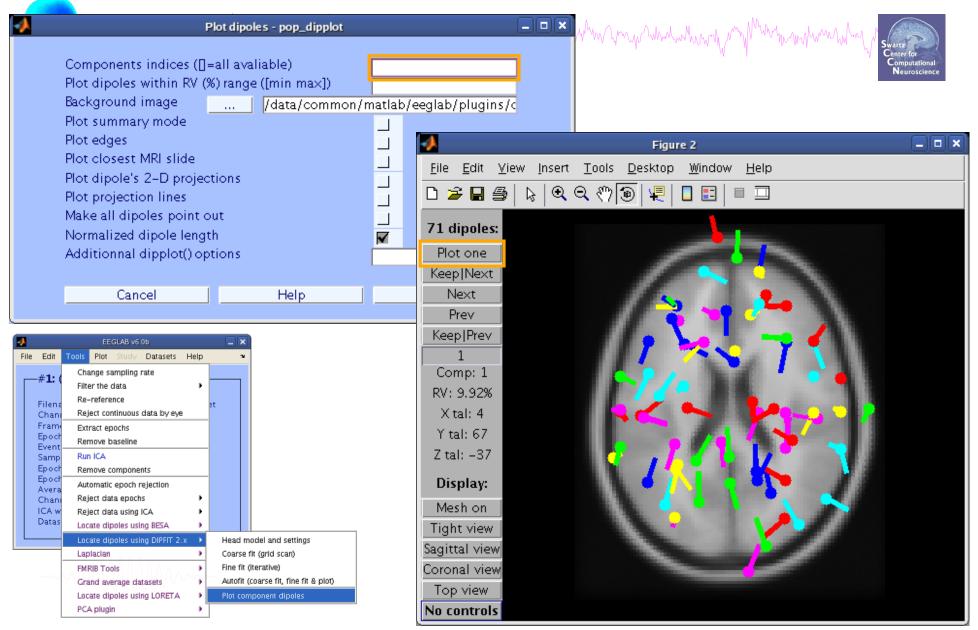
# Autofit equivalent dipoles

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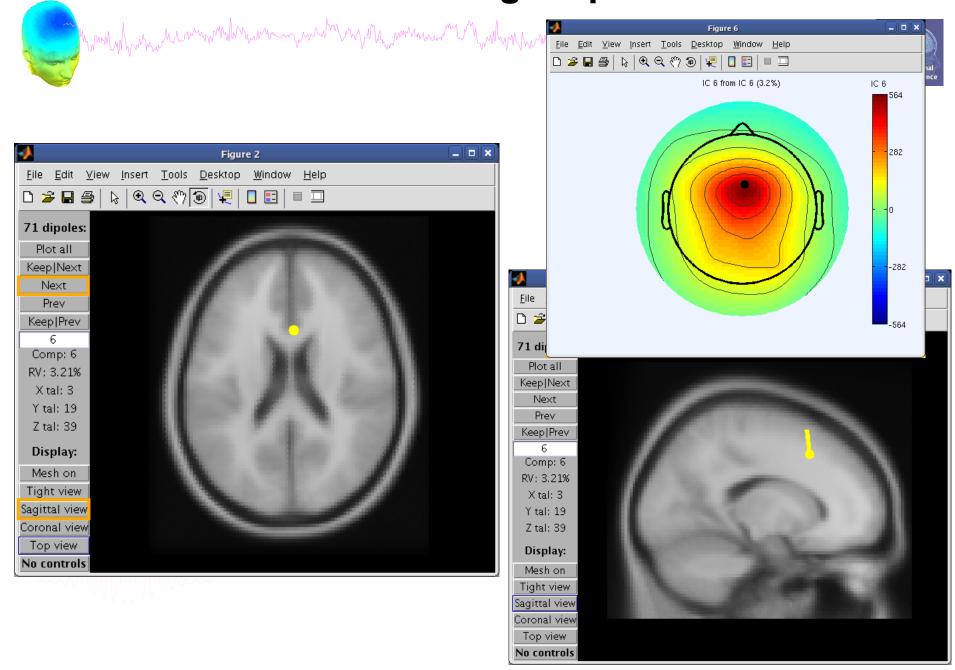




### **Plot dipoles**



### Scroll through dipoles



# Fine fit options in DIPFIT

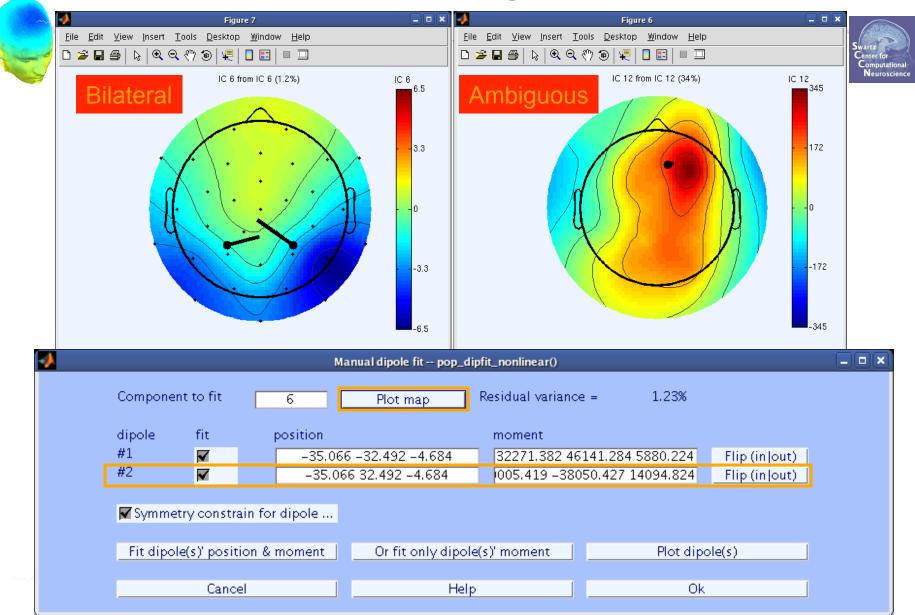
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		EME	RIB Too	ols		•	Fine	fit (	(iterative)	
		Grand average datasets					Autofit (coarse fit, fine fit & plot)			
		Loc	ate dip	oles usi	ng LORETA	•	Plot (	on	nponent dipoles	
		PCA	A plugii	n		• • • <sup>-</sup>				

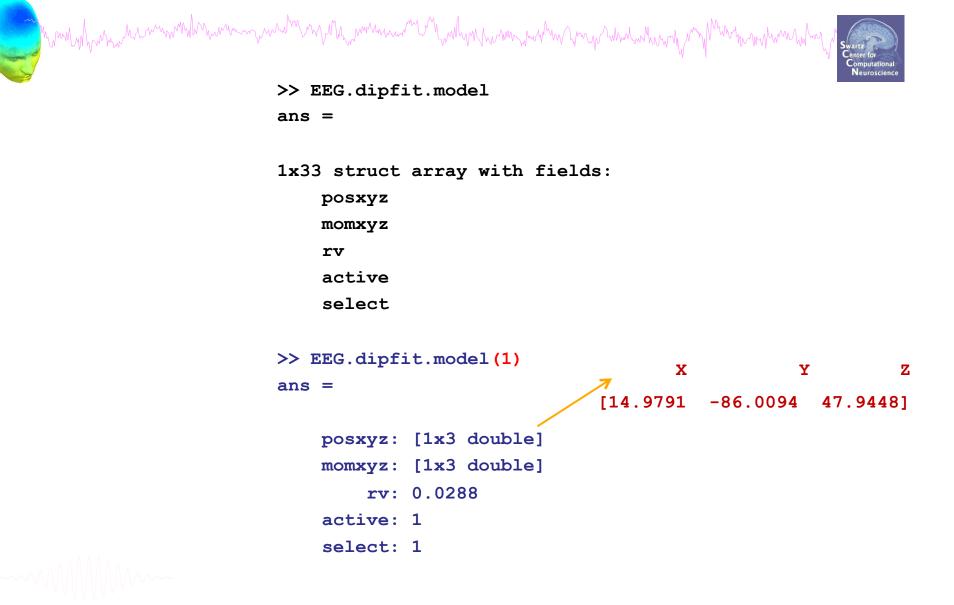


#### **Fine fit menu** how when the way when the work when have and have a show the show of the show the sh Figure 6 Eile Edit View Insert Tools Desktop Window Help 🗅 🗲 🖬 🞒 🖕 🔍 Q. (까 🕲 🐙 📘 📰 💷 IC 6 from IC 6 (3.2%) IC 6 564 282 -282 Manual dipole fit -- pop\_dipfit\_nonlinear() - 0 × Component to fit Residual variance = 3.21% Plot map 6 dipole position fit moment #1 28.222 -2.401 37.331 2,380 475942.653 3819304.288 Flip (in out) $\checkmark$ #2 0.000 0.000 0.000 0.000 0.000 0.000 Flip (in out) Symmetry constrain for dipole ... Fit dipole(s)' position & moment Or fit only dipole(s)' moment Plot dipole(s) Ok Cancel Help



#### **Bilateral dipoles**

#### **EEG.dipfit structure**



# DIPFIT and model co-registration



- 1. Co-register electrodes with model
- 2. Demonstration
- 3. 3D headplot co-registration

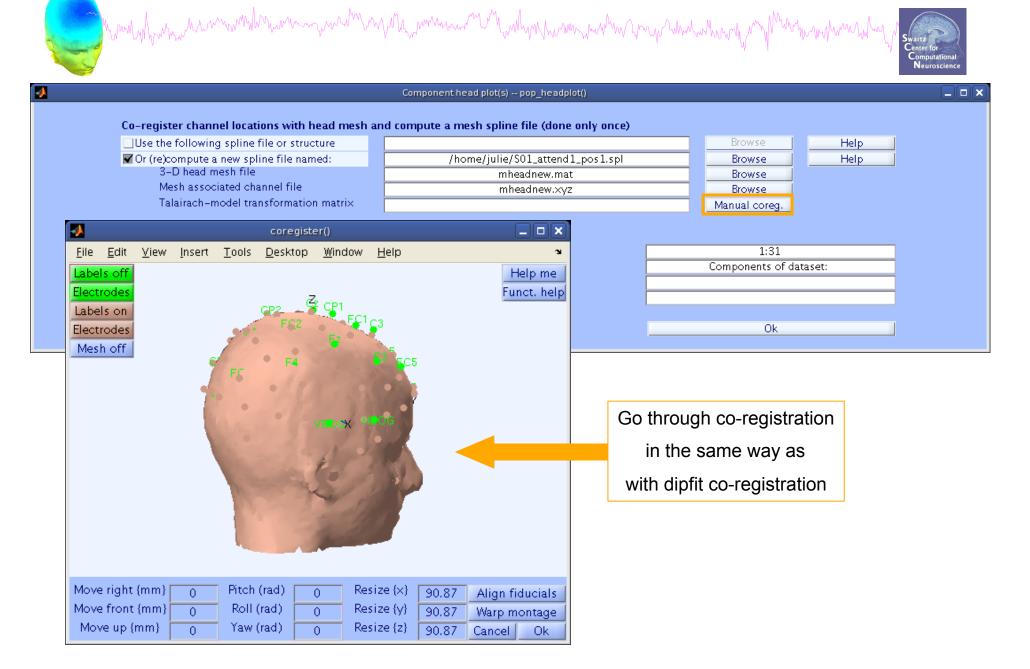
# Plot scalp maps in 3D



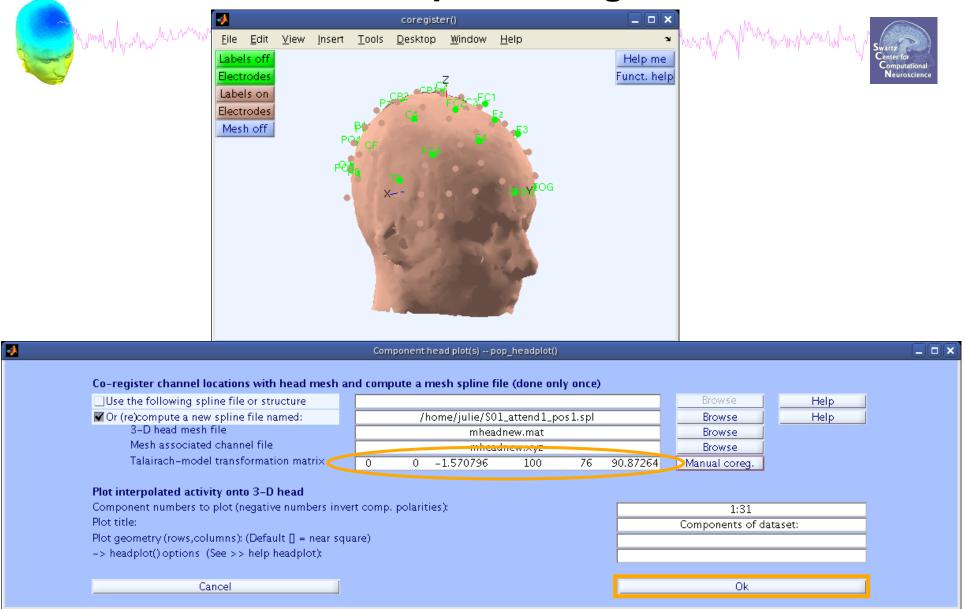


								Headplot() warning			
EEGLAB v6.0b File Edit Tools Plot Study Datasets Help					Help		tim	eadplot() must generate a spline file the first me it is called or after changes in the channel location file. ou must also co–register your channel locations with the			
	#1.7	n a d	Channel locations				head template.				
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#### **Headplot co-registration**



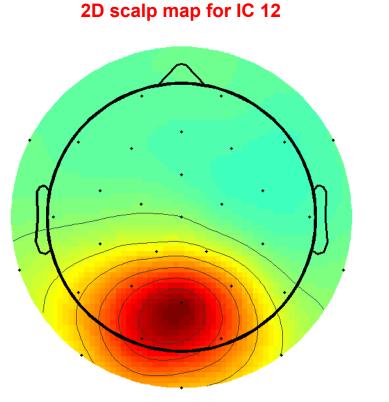
#### **Confirm headplot co-registration**

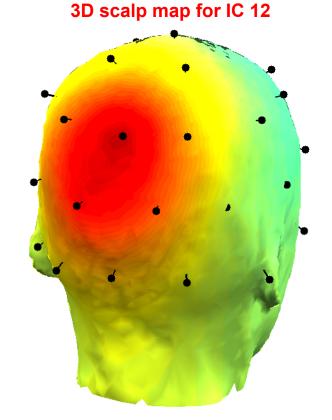


# **Spline file in EEG structure**









# Exercise

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- Load 'stern125.set'
- Practice co-registering electrodes with **BEM** model (choose 'Erase' because this dataset has co-registration done already)
- Autofit IC dipoles
- Fine fit dipoles
- Plot dipoles from the GUI; scroll through components individually
- Co-register the head model for 3D scalp map plotting. Then plot some ICs in 3D
- Advanced

- In the Finefit menu, try fitting a bilateral dipole, what happens to the residual variance?

- Try plotting a subset of dipoles in 'summary mode'

- Try purposely misaligning electrodes with model, how far off are the resulting dipoles from the original locations?

