

MEMORY: Close all your applications, including web browsers

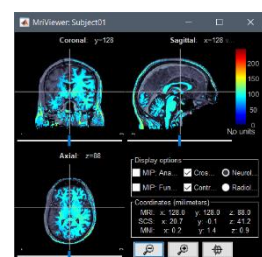
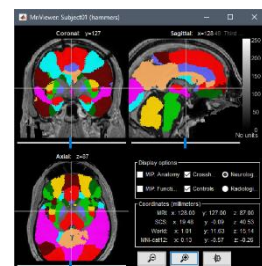
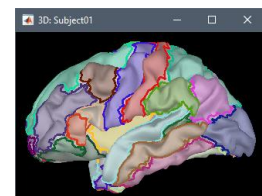
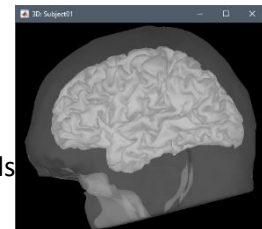
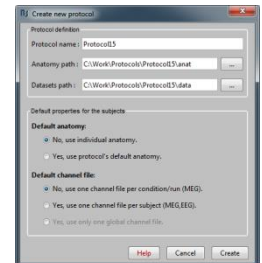
INTERNET: Internet connection needed for the MNI normalization

8:30-9:15 Onsite assistance in installing the material for the training session

9:15-9:45 Lecture: Introduction to Brainstorm

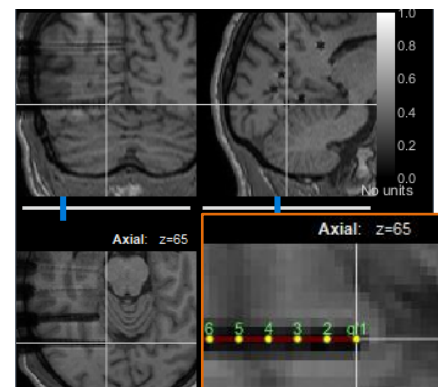
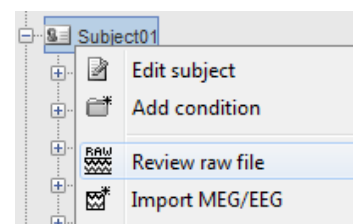
9:45-10:45 Import anatomy 1hr

- Start Brainstorm: from Matlab or stand-alone
- Create new protocol "Workshop":
 No, use individual anatomy
 No, use one channel file per acquisition run
- Create new subject:
 Introduction to database explorer: protocols, exploration modes, popup
 Switch to anatomy view: 1st button above database explorer
 Right-click on protocol top node > New subject: Subject01
- Import T1 pre-implantation + CAT12 segmentation:
 Right-click on Subject01 > **Import anatomy folder**
 Select format: **CAT12**
 Select folder: workshop_cuttingeeg/derivatives/cat12/sub-01_ses-pre
 Number of vertices: 15000
 Introduction to the MRI viewer: Click, mouse wheel, color bar, popup
 Compute MNI normalization / MAFF8: set default positions for the fiducials
 Coordinate systems: MRI, SCS, World, MNI
- How to run CAT12 segmentation from Brainstorm
- Display cortex:
 Close figure, double-click on cortex_15002V (low-resolution pial surface)
 3D figure: Rotation, zoom, predefined views
 Surface tab: Smooth slider, sulci, edges
 Scout tab: Parcellations of the surface vertices
- Volume parcellations: AAL3, Hammers, tissues
 Adjust transparency, change the atlas, non-linear MNI transformation
 Add MNI parcellation: Schaefer2018_100_7net
- Import T1 post-implantation:
 Right-click on Subject01 > **Import MRI**
 Select file: /sub-01/ses-postimp/anat/sub-01_ses-postimp_T1w.nii.gz
 Apply transformation: **YES**
 How to register: **IGNORE** (volumes are already co-registered with SPM)
 Reslice the volume: **YES** (so we can overlay them)
 Surface tab: Adjust threshold with slider "Data options: Amplitude"
- Rename the volumes **T1pre** and **T1post**
- **Close all:** Big cross on the top-right, close all the figures and empty memory



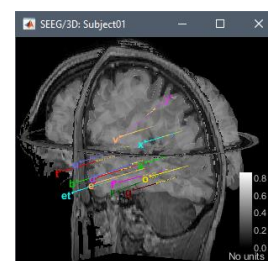
10:45-11:20 SEEG contact positions 35 min

- Switch to functional view : 2nd button above database explorer
- Create link to SEEG recordings:
 - Right-click on Subject01 > **Review raw file**
 - Select format: **SEEG: Deltamed/Micromed/...**
 - Select file: sub-01/ses-postimp/ieeg/sub-01_..._ieeg.eeg
 - Double click on the channel file: Missing 3D positions
- Mark electrode **g'**:
 - Right-click on channel file > MRI registration > Edit: **T1post**
 - Tab iEEG: Select **g'**: **DIXI D08-12AM Microdeep**
 - Set tip: [X=115, Y=64, Z=65] - Axial view: Lowest horizontal
 - Set entry: [X=76, Y=64, Z=64]
 - Menu Contacts > Save modifications
- Contacts > Compute atlas labels
 - File selection: Cancel (do not save) / Keep default options



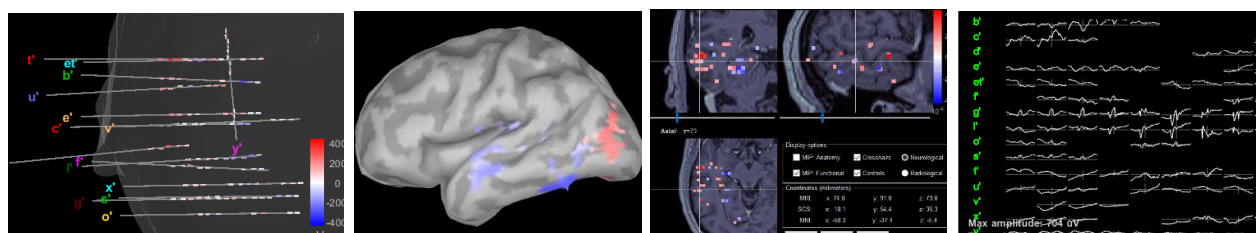
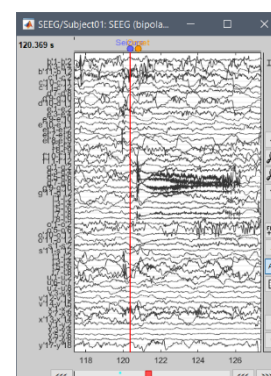
Channel	SCS	aal3	hammers	tissues_cat12	Schaefer2018_100_7net (MNI-cat12)
g'1	[-46.734,12.724,33.596]	Left Lingual gyrus	OL lingual gyrus L	Gray	Vis_3 L
g'2	[-46.713,16.224,33.526]	Left Lingual gyrus	OL lingual gyrus L	White	Vis_3 L
g'3	[-46.692,19.723,33.455]	Left Lingual gyrus	OL lingual gyrus L	White	Vis_3 L
g'4	[-46.671,23.222,33.385]	Left Lingual gyrus	OL lingual gyrus L	White	Vis_2 L
g'5	[-46.649,26.721,33.315]	Left Lingual gyrus	OL lateral remainder occipital lobe L	Gray	Vis_2 L
g'6	[-46.628,30.220,33.244]	N/A	OL lateral remainder occipital lobe L	White	N/A

- Import positions for all the electrodes
 - Right-click on channel file > Add EEG positions > Import from file
 - Select format: **EEG: BIDS electrodes.tsv, subject space mm**
 - Select file: sub-01/ses-postimp/ieeg/..._space-Other_electrodes.tsv
 - Scaling factor: 1
 - Double-click on channel file, Move slices, Add cortex from Surface tab



11:20-11:45 Review the recordings 25 min

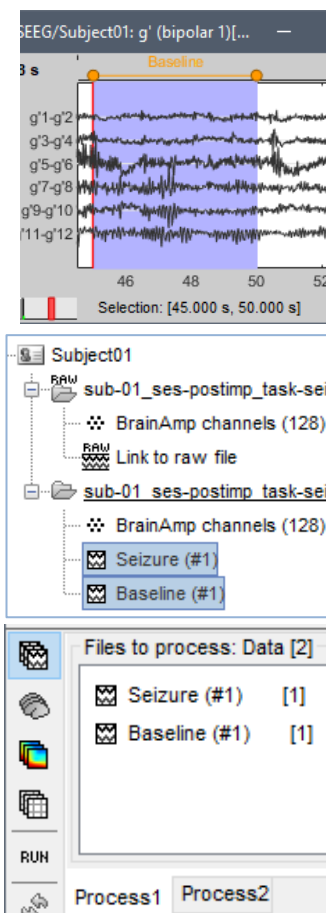
- Review SEEG: Right-click on "Link to raw file" > SEEG > **Display time series**
 - Display in columns: Button [~] in the Record tab
 - Amplitude: Buttons and shortcuts, Display menu
 - Time: Display windows of **10 seconds**, Scroll with F3, Auto-Scale button
 - Select bad channels (o'1) + right click > Channels > **Mark selected as bad**
 - Montages: **SEEG (Bipolar 1)** - Create personal montages + shortcuts
 - Online filters
- Right-click on "Link to raw file" > SEEG > **2D Layout**
- Change the window layout to "Tiled"
- Other menus: **Display on cortex**, **Display on MRI (MRI Viewer)**, **3D Electrodes (Head)**



Epileptogenicity maps

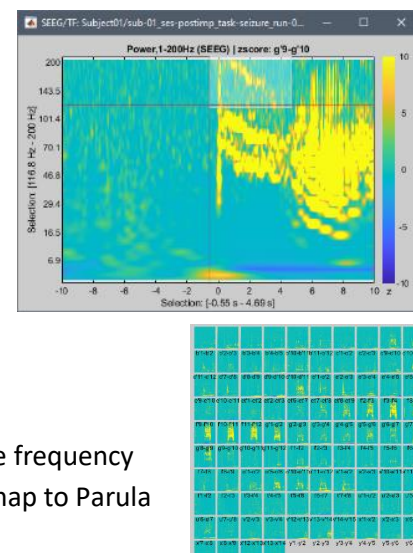
Import epochs of interest 15 min

- Baseline:
 - Change montage: **SEEG (bipolar 1)**
 - Create a new category of events: **Add group > Baseline**
 - Select time: **45s - 50s** (right-click > Time selection > Set selection)
 - Add baseline event
- Close figure, save modifications
- Import seizure:
 - Right-click on "Link to raw file" > Import in database
 - Use events: **Seizure**, Epoch time: **[-10000, +40000] ms**
 - NO Remove DC offset**
 - Do **NOT** select: Create a separate folder for each event type
- Import baseline:
 - Right-click on "Link to raw file" > Import in database
 - Use events: **Baseline**
- Apply bipolar montage:
 - Drag and drop the two imported files in the Process1 box
 - Run > Process Standardize > **Apply montage**:
 - Montage name: **Subject01: SEEG (bipolar 2)**
 - Create new folders: **YES**



Time-frequency analysis OPTIONAL 5min

- In Process1: Select "Seizure"
- Select process: **Extract > Extract time > [-10, +10]s**
- Add process: Frequency > **Time-frequency (Morlet wavelets)**:
 - Sensor types: **SEEG**
 - Log 1:40:200**, Central frequency: **1Hz**, Time resolution: **10s**
- Add process: Standardize > **Baseline normalization**:
 - Baseline: **[-7.5, -2.5] s**, **Z-score**, **Overwrite**
- Run, right-click on result > All channels
- Look for the highest frequency band with significant activity:
 - Hypothesis: the closest to the seizure onset zone, the highest the frequency
 - Smooth display, Set colormap max to **[-10,+10]z**, Change colormap to Parula
 - Frequency band of interest: **120-200 Hz**



Epileptogenicity maps 20min

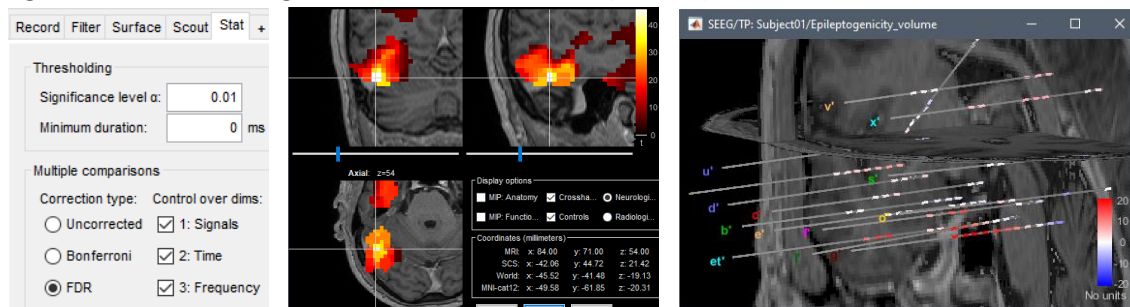
- Requires SPM to be installed
- Epileptogenicity maps in volume:

In Process2: A=Baseline, B=Seizure, Process **Epilepsy** > **Epileptogenicity maps**

Frequency band: **120-200 Hz**, Latency: **0**, **Volume**

Right-click on “source” > Cortical activations > Display on MRI (MRI viewer)

Right-click on “recordings” > SEEG > 3D electrodes (MRI 3D)



- Epileptogenicity maps on cortex surface:

In Process2: A=Baseline, B=Onset, Process **Epilepsy** > **Epileptogenicity maps**

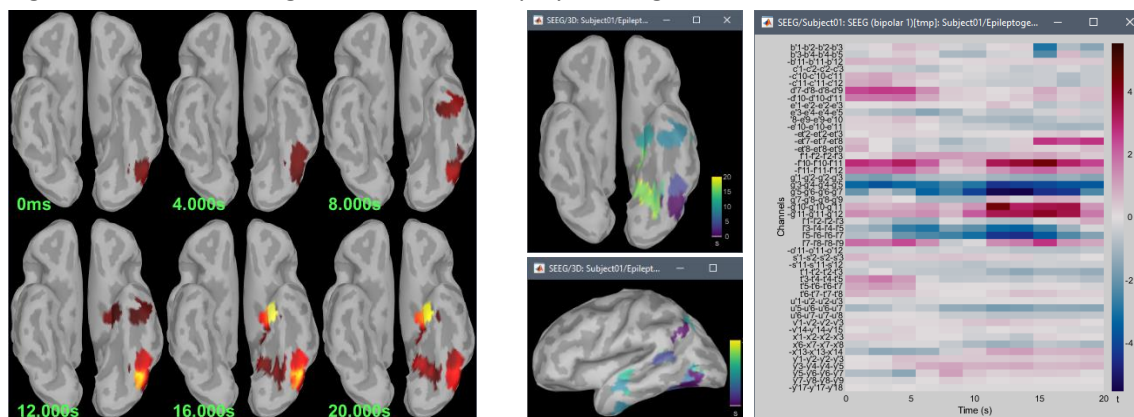
Frequency band: **120-200 Hz**, Latency: **0:2:20**, **Surface**

Double click on the two “source files”:

El_..._120_200_3: Epileptogenicity maps are now time resolved

Delay_..._120_200_3_50: Delay map = all the epileptogenicity maps ($p < 0.05$) overlaid for all the time points, colormap=seconds

Right-click on “recordings” file > SEEG > Display as image



- Additional topics, if any time left:
 - Guidelines panel: Simplified interface for this pipeline, designed for clinicians
 - ECOG example
 - Montage editor

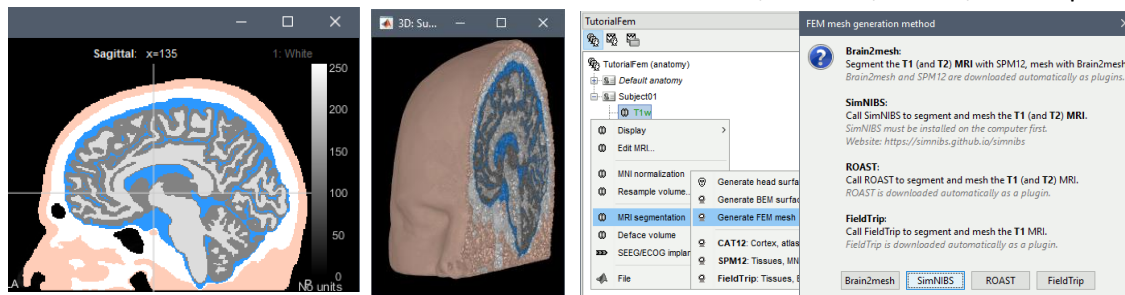
FEM Modeling 15min

Brainstorm FEM tutorial: MEG/EEG Median nerve stimulation

<https://neuroimage.usc.edu/brainstorm/Tutorials/FemMedianNerve>

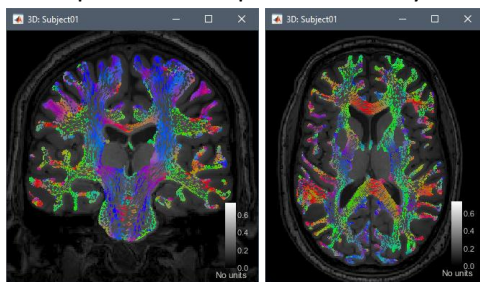
1) Segmentation

- Obtain a good segmentation of the head tissues : GM, WM, CSF, skull, skin
- Create a 3D tetrahedral mesh of the head tissues: Brain2mesh, SimNIBS, ROAST, FieldTrip



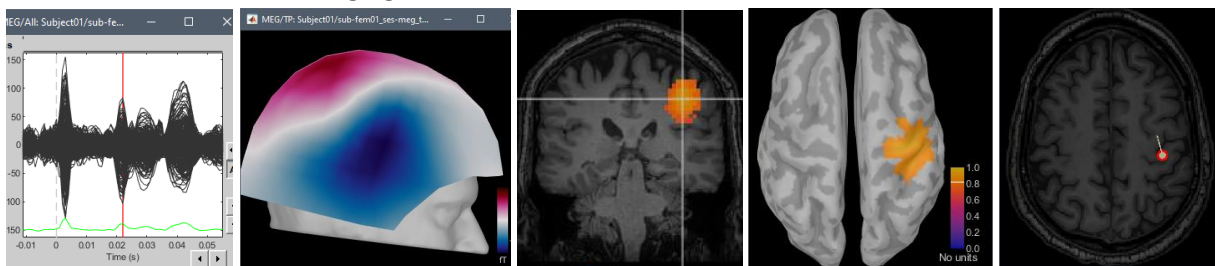
2) Evaluate tissue anisotropy

- Compute DTI tensors from DWI images (BrainSuite Diffusion Pipeline)
- Compute anisotropic conductivity tensors for the tetrahedral elements of the white matter



3) Estimate sources

- Compute FEM forward model with DUNEuro
- Minimum norm source imaging



Future applications:

- Modeling local currents around implanted electrodes for SEEG and DBS
- Modeling of intracranial currents for transcranial stimulations tACS/tDCS