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Propagation of epileptic spikes revealed by diffusion-based constrained MEG source reconstruction

Athena

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Goal: Study of the propagation of an epileptic spike.

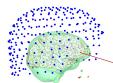
Method: 1- cortex parcellation via structural information coming from diffusion MRI (dMRI)

- 2- MEG inverse problem on a parcellated source space
- 3- study of the propagation of an epileptic spike via the active parcels

Results on real data allowing to study the spatial propagation of an epileptic spike.

1 Preprocessing

- 1- Co-registration of the T1wMRI and dMRI [soft FSL]
- 2- Surface meshes extraction [soft Freesurfer]
- 3-Computation of the leadfield matrix G [soft OpenMEEG]



Each vertice of the WM/GM boundary mesh is:

- Source of a magnetic field
- Seed for tractography

2 dMRI-based cortex parcellation

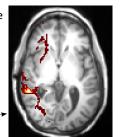
Connectivity profile of a source CP:

CP(i) = connectivity value between the source and the ith voxel.

4- For each source:

Computation of its connectivity image.

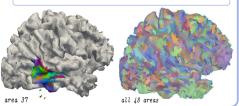
Tool: Probtrackx [soft FSL]



5- Pre-clustering via Brodmann's atlas

For each Brodmann area, clustering of sources having a close connectivity to all sources:

- 5.1- Computation of the CP-based correlation matrix R_[5]
- 5.2- K-means algorithm on R



3 MEG inverse problem with parcellated source space

We reduce the inital distributed source space S into s in accordance with the cortex parcellation:

$$S = P \times s$$
 with $P(i, j) = \begin{cases} 1, & \text{if source } i \text{ is in area } j \\ 0, & \text{if not} \end{cases}$

and compute the reduced leadfield $\,G_P$ for the parcellated source space $s\colon G_P=G\times P$

Thus, the MEG inverse problem on the parcellated source space using Tikhonov regularization becomes :

$$\min_{s} || \mathbf{M} - \mathbf{G}_{p} \mathbf{s} ||^{2} + \lambda || \mathbf{s} ||^{2}$$

where M contains the MEG measurements.

4 Study of the propagation of an epileptic spike

At each time sample, we want to determine the cortical areas at the origin of the activity. We call \mathscr{S}_t the set of these areas, for a time sample t

For each time sample t, we compute the power $\mathscr{P}_{p,t}$ of each area p on a sliding time window $[t-\alpha,t+\alpha]$:

$$\begin{split} \mathscr{P}_{p,t} &= \sum_{i=t-\alpha}^{t+\alpha} |s(p,i)|^{2}. \\ \mathscr{S}_{t} &= \{p_{a} : \mathscr{P}_{p_{a},t} > F * \max(\mathscr{P}_{p,t}), \forall p\} \end{split}$$

with F a pourcentage.

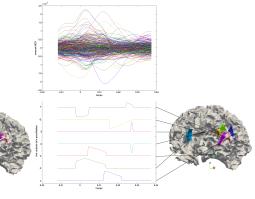
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5 Results & Conclusion

Almost the same parcels are activated for all epileptic spikes.

- The direction of propagation changes: from the back of the frontal lobe to the front or opposite direction.
- The time of activation of each parcel characterizes the spike.

Results on 3 epileptic spikes of a single subject.



- The parcellation allows an easier representation of source space.
- The method reveals differences between spikes (direction of propagation and time of activation of parcels).
- Future works will be to analyse the structural network supporting the propagation.