Exploring variability patterns across fMRI statistic maps

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Introduction

- Neuroimaging workflows are highly flexible with many possible choices at each step of an analysis (*Carp 2012*).
- Different analytical choices can cause variation in the results (Botvinik-Nezer et al. 2020).
- Exploring the **pipeline space** is essential to understand the relationships between results and to guide researchers with good practices.

 \rightarrow Identifying groups of pipelines that provide similar results.

Materials & Methods

HCP many pipelines dataset

- The Human Connectome Project (Van Essen et al. 2013): 1080 participants with MOTOR task fMRI data, with 6 base contrasts to study.
- Pipelines constrained on **specific parameters**:

Software package Smoothing kernel SPM or FSL

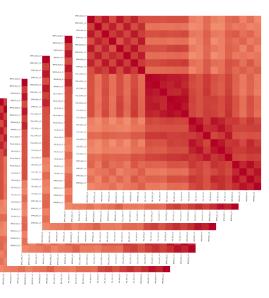
Motion regressors 0,6 or 24

HRF derivatives Included (1) or not (0)

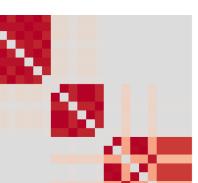
FWHM 5 or 8 mm

Louvain method for community detection

Extraction of **communities from large networks** based on the optimization of modularity (Blondel et al. 2008).



- 1. Correlation matrices between maps of different pipelines for each group.
- 2. Reorganize each group matrix with Louvain communities.



3. Global matrix of pipelines with occurrence of belonging to the same partition across all groups.

 \rightarrow Assessing the stability of these groups in different contexts.

\rightarrow 24 different pipelines

1000 groups of 50 randomly sampled participants.

Code available at: https://gitlab.inria.fr/egermani/hcp_pipelines



4. Reorganize global matrix with Louvain communities.

 \rightarrow Matrix and graph organized based on similarity of pipelines statistic maps across different groups.

SAME DATA, DIFFERENT PIPELINES

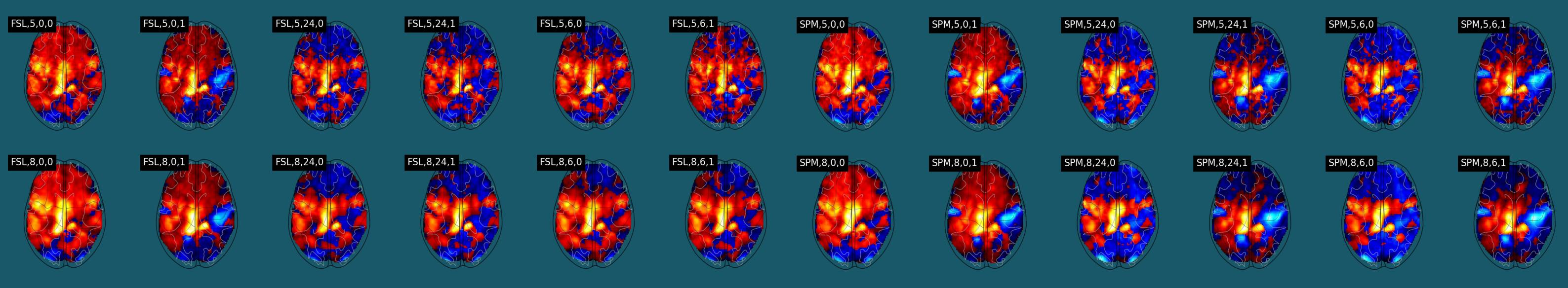


Fig1. Group-level statistic maps (N=50) obtained with 24 different pipelines for the contrast right-hand. Pipelines are named as followed: Software, Smoothing kernel, Motion regressors, HRF derivatives.

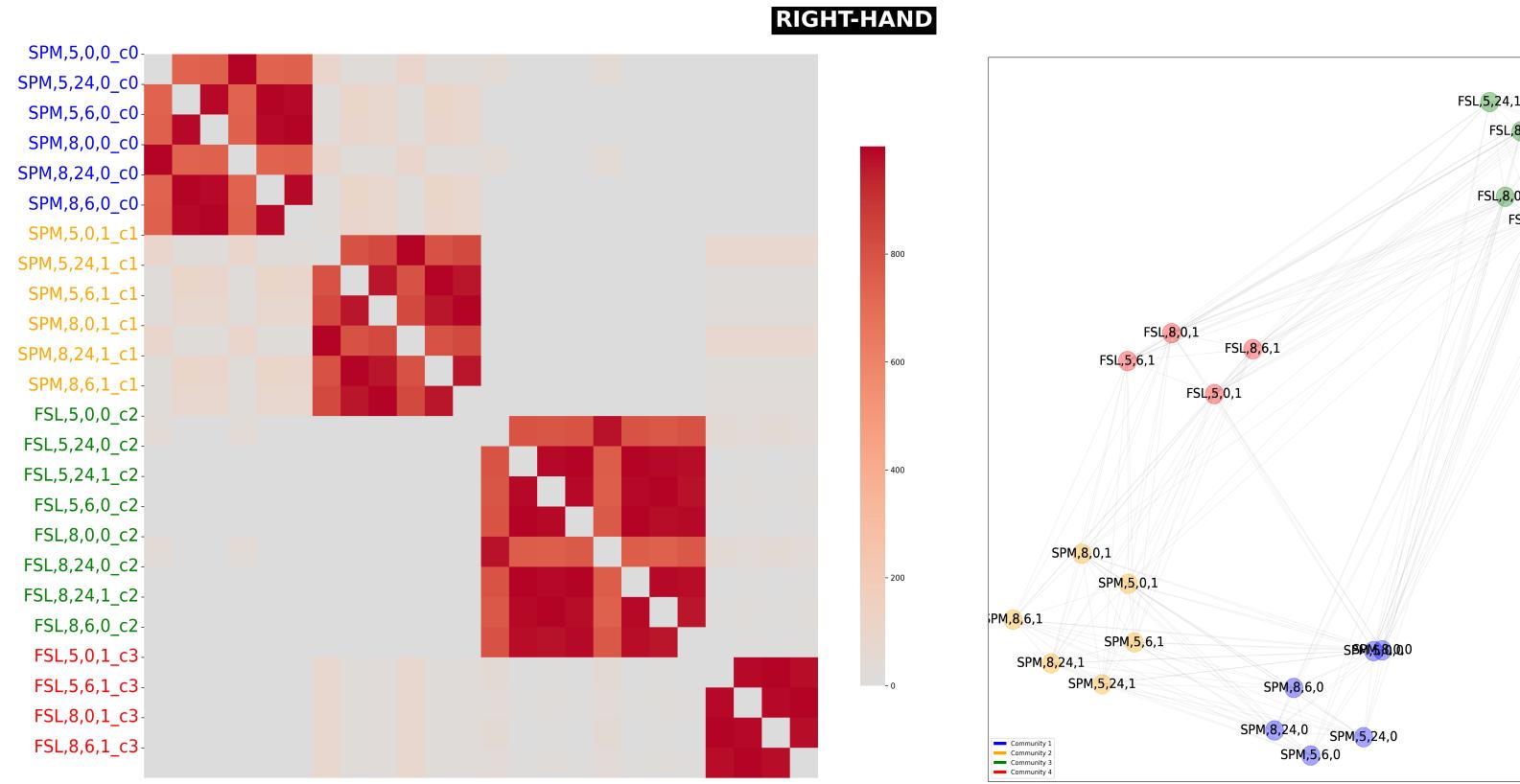
FSL,8,24,0

FSL,5,0,0

FSL,8,0,0

FSL,854516





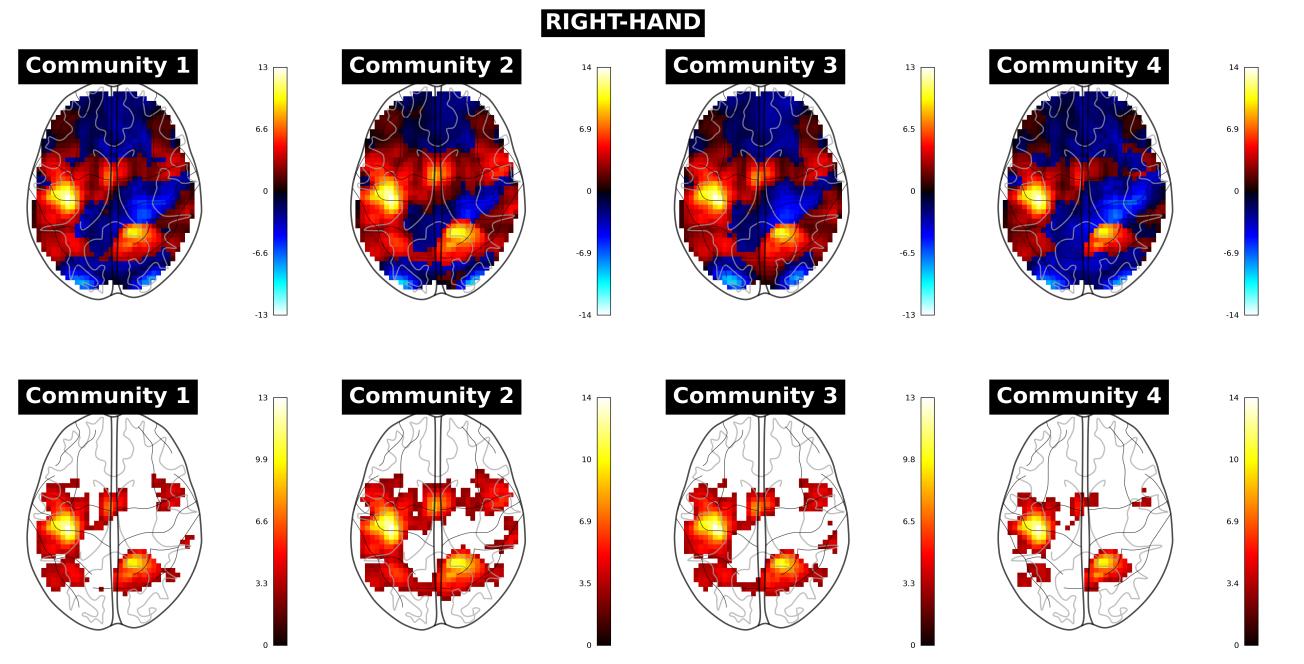


Fig3. Mean statistic maps of communities for the contrast right-hand. Unthresholded maps (*upper*) were thresholded using a FDR of p<0.05 (lower).

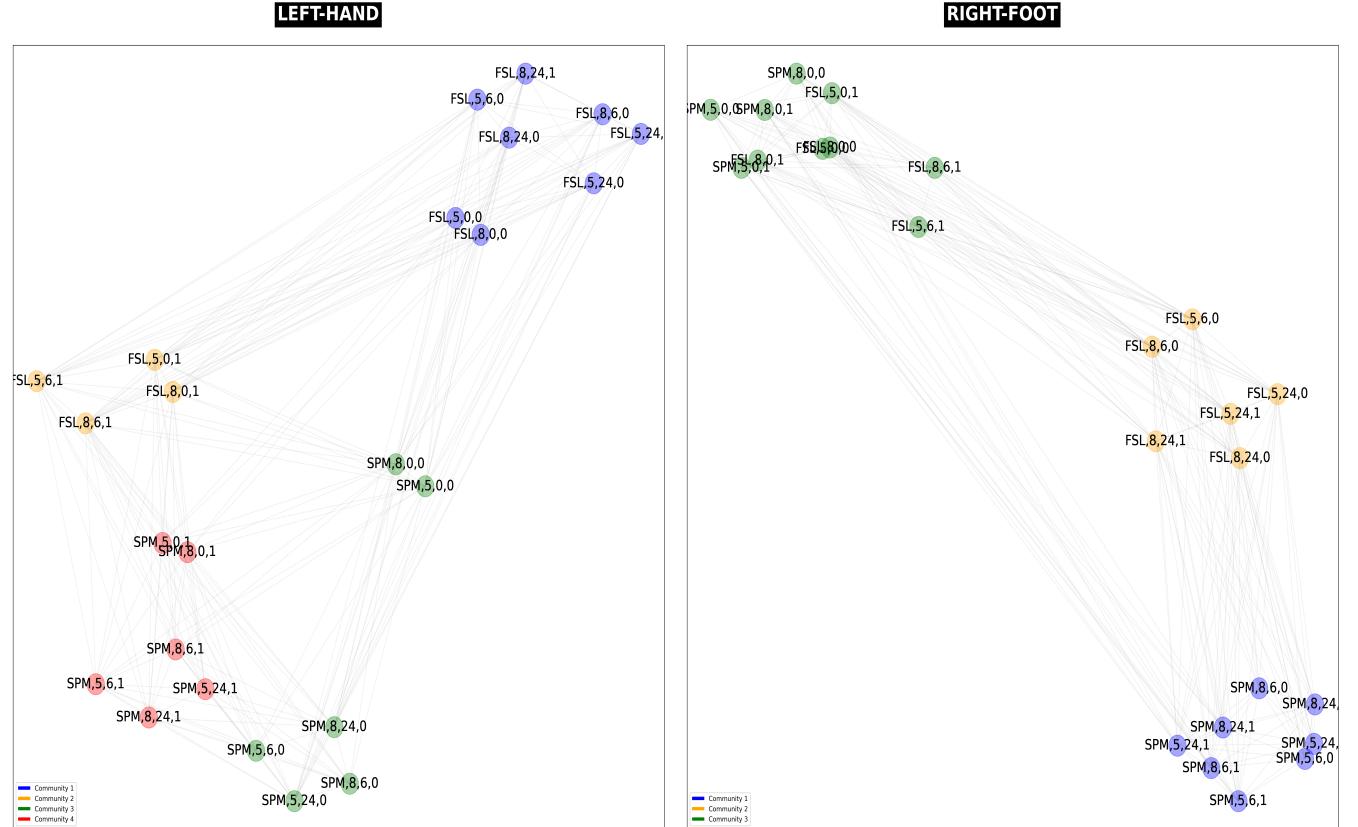


Fig2. Louvain matrix and graph representing the number of times pipelines belong to the same community for pipelines across different group-level statistic maps of the contrast right hand.

- different groups of participants.
- ✓ Stability of communities of pipelines across ✓ Small differences regarding the extent of the activation area.
- ✓ Pipelines grouped with those sharing specific different pipelines, parameters: same smoothing \rightarrow same community
- ✓ Similar number and groups of pipelines in communities for *right hand* and *left hand*: robustness to the global contrast.

\checkmark Global activation patterns are consistent \checkmark Communities are different for the contrast between communities. right foot.

Conclusion

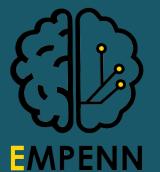
Pipelines have different behaviors depending on contrasts, probably due to specific properties of the data. Further work will be needed in order to determine specific patterns of the communities, understand why some pipelines behave differently and how to use these patterns to tackle **analytical variability** with, for instance, style transfer techniques.

Fig4. Louvain graphs representing number of times pipelines belong to the same community across different group-level statistic maps for the contrasts right foot (*right*) and left hand (*left*).

References

Botvinik-Nezer, R. et al. 2020. « Variability in the analysis of a single Van Essen, D. C., for the WU-Minn HCP Consortium. (2013). « The neuroimaging dataset by many teams ». Nature. WU-Minn Human Connectome Project: An overview ». NeuroImage. Carp, J. 2012. « On the Plurality of (Methodological) Worlds: Blondel, V. et al, 2008. « Fast unfolding of communities in large networks », Journal of Statistical Mechanics: Theory and Estimating the Analytic Flexibility of FMRI Experiments ». Frontiers in Neuroscience. Experiment, vol. 2008, no 10.

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