Rank-adaptive covariance changepoint detection for estimating dynamic functional connectivity from fMRI data

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Agenda

- Scientific Background
- Research Question
- Methodology
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Functional MRI (fMRI)

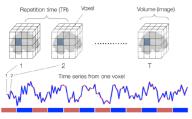
Magnetic resonance imaging (MRI) comes in two types

- Structural: anatomy and pathology of brain
- Functional (fMRI): brain activity

Brain activity is generally measured via **Blood Oxygen Level Dependent** (**BOLD**) signals. Safe to collect, 95%+ of fMRI studies use this.



(a) MRI scanner [1]



(b) Illustration of data collected for each voxel over time [2]

Functional Connectivity (FC)

"Functional connectivity is defined as the undirected association between two or more fMRI time series" [2].

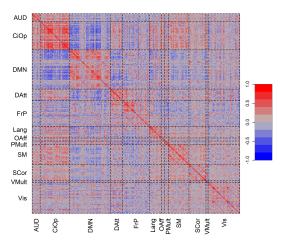


Figure: Functional connectivity matrix (sample correlation) of fMRI time series using all time points. A clear low-rank block structure is present.

Dynamic Functional Connectivity

- **Dynamic functional connectivity (DFC)**: functional connectivity which changes over time.
- Unknown if DFC best characterized as changing in discrete fashion (e.g. ABAABA) or continuously.

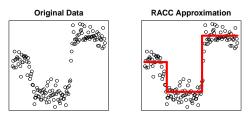


Figure: Illustration of RACC's approximation of FC

Potential biomarkers:

- Number of changes
- Time per state
- High connectivity vs. low connectivity



Time-Varying Sample Covariance Matrix - Windowed

Covariance matrix of rs-fMRI data for an example subject. Window width of 20 observations.

Research Question

Our research addresses how to best segment resting state fMRI (rs-fMRI) time series into segments with constant covariance structure. We take a changepoint detection approach and propose RACC: Rank Adaptive Covariance Changepoint detection.

Challenges of rs-fMRI Segmentation

- Type I error control & selective inference
- p ≫ n

Features of RACC

- Binary Segmentation Algorithm
- Permutation Test (and controlled Type I errors)
- Adaptive test statistic (low-rank covariance structure)

RACC Segmentation of rs-fMRI Data

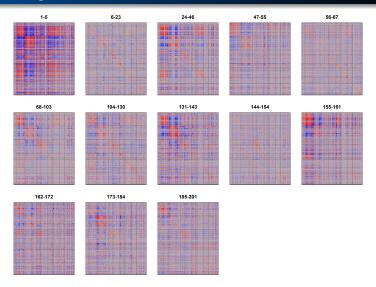


Figure: Empirical covariance matrices for the 13 segments of constant covariance after changepoints were identified by RACC.

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RACC Methodology

To test H_0 of constant covariance structure we use the following test statistic

$$T = \underbrace{\max_{\substack{i \in 2, \dots, n-2 \\ \text{max over} \\ \text{time}}} \max_{\substack{s \in \mathcal{S} \\ \text{max over} \\ \text{matrix norms}}} \frac{||\hat{\Sigma}_i^L - \hat{\Sigma}_i^R||_s - \hat{\mu}_{H_0, i, s}}{\hat{\sigma}_{H_0, i, s}}$$

$$\hat{\mu}_{H_0,i,s} = \widehat{E}_{H_0} \left[|| \hat{\Sigma}_i^L - \hat{\Sigma}_i^R ||_s \right] \quad \text{ and } \quad \hat{\sigma}_{H_0,i,s}^2 = \widehat{Var}_{H_0} \left[|| \hat{\Sigma}_i^L - \hat{\Sigma}_i^R ||_s \right]$$

Use permutations to find critical values of T, if reject null changepoint identified at $\hat{i} = \operatorname{argmin}_i \max_{s \in \mathcal{S}} \frac{||\hat{\Sigma}_i^L - \hat{\Sigma}_i^R||_s - \hat{\mu}_{H_0,i,s}}{\hat{\sigma}_{H_0,i,s}}$

Adaptive Ky-Fan(k) Statistic

$$T = \max_{i \in 2, \dots, n-2} \max_{s \in \mathcal{S}} \frac{||\hat{\Sigma}_i^L - \hat{\Sigma}_i^R||_s - \hat{\mu}_{H_0, i, s}}{\hat{\sigma}_{H_0, i, s}}$$

- Covariance matrix of rs-fMRI characterized by low-rank block structure
- Seek to improve power by looking at largest singular values of $\hat{\Sigma}_i^L \hat{\Sigma}_i^R$

$$||\hat{\Sigma}_i^L - \hat{\Sigma}_i^R||_{\mathsf{Ky-Fan}(k)} = \sum_{\ell=1}^k \sigma_\ell (\hat{\Sigma}_i^L - \hat{\Sigma}_i^R)$$

 $\bullet \ \mathcal{S} = \{\mathsf{Frobenius}^2, \mathsf{Ky}\text{-}\mathsf{Fan}(1), \mathsf{Ky}\text{-}\mathsf{Fan}(2), \ldots, \mathsf{Ky}\text{-}\mathsf{Fan}(\mathcal{K})\}$



Real Data Analysis - Single Subject

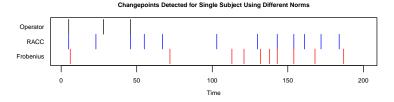


Figure: Final segmentation produced by RACC (utilizing Ky-Fan(k) norms), compared to just using operator or Froebnius norms.

Real Data Analysis - Multi-Subject

Previous research has investigated number of changes in FC differed in case vs. control

- More changes for subjects with Parkinson's [3]
- No differences for subjects with Mild Cognitive Impairment [4]
- More changes for subjects with moderate to severe non-suicidal self-injury [5]

We examine 135 subjects from the SPINs study (85 with schizophrenia, 45 healthy controls, 5 unknown). Find no statistically significant difference between number of changepoints identified by RACC (10.79, 10.59 average number of changepoints for case and control groups).

Extensions/Limitations

- How to choose an optimal K to choose
- Theory around consistency of detected changepoint in multiple changepoint case
- More comprehensive preprocessing to ensure exchangability of data
- Non-linear dimension reduction techniques
- Clustering dFC after segmentation

Thank you!

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References I

- [1] Mri scan https://www.nhs.uk/conditions/mri-scan.
- [2] Tor D Wager and Martin A Lindquist. Principles of fmri. *New York: Leanpub*, 2015.
- [3] Jinhee Kim, Marion Criaud, Sang Soo Cho, María Díez-Cirarda, Alexander Mihaescu, Sarah Coakeley, Christine Ghadery, Mikaeel Valli, Mark F Jacobs, Sylvain Houle, et al. Abnormal intrinsic brain functional network dynamics in parkinson's disease. *Brain*, 140(11):2955–2967, 2017.
- [4] Núria Mancho-Fora, Marc Montalà-Flaquer, Laia Farràs-Permanyer, Daniel Zarabozo-Hurtado, Geisa Bearitz Gallardo-Moreno, Esteban Gudayol-Farré, Maribel Peró-Cebollero, and Joan Guàrdia-Olmos. Network change point detection in resting-state functional connectivity dynamics of mild cognitive impairment patients. *International Journal* of Clinical and Health Psychology, 20(3):200–212, 2020.
- [5] Mark B Fiecas, Christian Coffman, Meng Xu, Timothy J Hendrickson, Bryon A Mueller, Bonnie Klimes-Dougan, and Kathryn R Cullen. Approximate hidden semi-markov models for dynamic connectivity analysis in resting-state fmri. *bioRxiv*, pages 2021–03, 2021.

References II

[6] Erich Leo Lehmann, Joseph P Romano, and George Casella. *Testing statistical hypotheses*, volume 3. Springer, 2005.

Theoretical Results

Assuming normal data distinct egenvalues, common eigenvectors $\Sigma_1 - \Sigma_2 = \sum_{j=1}^d u_{l_j} (\lambda_{\delta,j}) \tilde{u}_{l_j}^{\top}$, where \tilde{u}_{l_j} and $\lambda_{\delta_j} = |\lambda_{1,l_j} - \lambda_{2,l_j}|$

Theorem (Limits of Singular Values)

- $\lambda_{\delta,j} \neq 0 : \sqrt{n}(\hat{\lambda}_{\delta,j} \lambda_{\delta,j}) \stackrel{d}{\rightarrow} \mathcal{N}(0, 2(\lambda_{1,l_j}^2 + \lambda_{2,l_j}^2))$
- $\lambda_{\delta,j} = 0 : \sqrt{n} \hat{\lambda}_{\delta,j} \stackrel{d}{\to} \mathcal{F}(0, 2(\lambda_{1,l_j}^2 + \lambda_{2,l_j}^2))$ where \mathcal{F} a folded normal distribution

Theorem (Limits of Ky-Fan(k)-norm)

Suppose exists $k \in \{1, ..., p\}$ s.t. $j \le k_0 \ \lambda_{\delta,j} \ne 0$ and $j > k_0, \ \lambda_j = 0$

$$\bullet \ k \leq k_0 \ : \ \sqrt{n} \Big(\| \hat{\Sigma}_1 - \hat{\Sigma}_2 \|_{(k)} - \| \Sigma_1 - \Sigma_2 \|_{(k)} \Big) \xrightarrow{d} \mathcal{N} \left(0, \textstyle \sum_{j=1}^k 2(\lambda_{1,l_j}^2 + \lambda_{2,l_j}^2) \right)$$

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Identification of a Changepoint via Permutation Test

Distribution of T complex; permutation tests ensure Type I error control under null for *any* test statistic of the data [6]. Use permutations to set critical value of T and estimate $\hat{\mu}_{H_0,i,s}$, $\hat{\sigma}^2_{H_0,i,s}$.

Under null of equal covariances across time

$$\underbrace{\frac{\chi_{1}, \chi_{2}, \chi_{3}, \chi_{4}, \chi_{5}}{\Sigma_{1:5}}}_{\Sigma_{1:5}} | \underbrace{\frac{\chi_{6}, \chi_{7}, \chi_{8}, \chi_{9}, \chi_{10}}{\Sigma_{6:10}}}_{\Sigma_{6:10}} \Rightarrow \Sigma_{1:5} - \Sigma_{6:10} = 0$$

$$\underbrace{\chi_{8}, \chi_{4}, \chi_{1}, \chi_{7}, \chi_{10}}_{\Sigma_{1:5}^{*}} | \underbrace{\chi_{9}, \chi_{2}, \chi_{3}, \chi_{5}, \chi_{6}}_{\Sigma_{6:10}^{*}} \Rightarrow \Sigma_{1:5}^{*} - \Sigma_{6:10}^{*} = 0$$

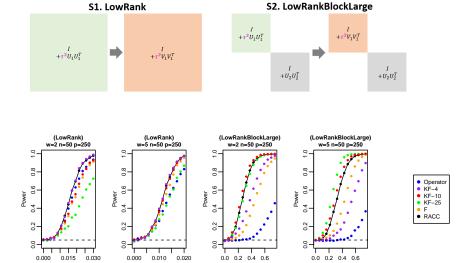
Under alternative of change in covariance at t=5

$$\underbrace{\frac{X_{1}, X_{2}, X_{3}, X_{4}, X_{5}}{\Sigma_{1:5}}}_{\Sigma_{1:5}} | \underbrace{\frac{X_{6}, X_{7}, X_{8}, X_{9}, X_{10}}{\Sigma_{6:10}}}_{\Sigma_{6:10}} \Rightarrow \underbrace{\Sigma_{1:5} - \Sigma_{6:10}}_{\Sigma_{6:10}} \neq 0$$

$$\underbrace{\frac{X_{8}, X_{4}, X_{1}, X_{7}, X_{10}}{\Sigma_{1:5}^{*}}}_{\Sigma_{5:8}^{*}} | \underbrace{\frac{X_{9}, X_{2}, X_{3}, X_{5}, X_{6}}{\Sigma_{5:8}^{*}}}_{\Sigma_{5:8}^{*}} \Rightarrow \Sigma_{1:5}^{*} - \Sigma_{6:10}^{*} \approx 0$$

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Simulation Results - Power



Here w represents the rank of the covariance matrix before and after a change-point.

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Simulation Results - Two Changepoint Case

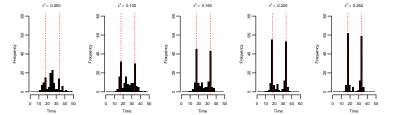


Figure: Power results for two equally spaced changepoints, (LowRank) w=5 setting for increasing magnitude of difference τ^2 .

Real Data Analysis - Single Subject

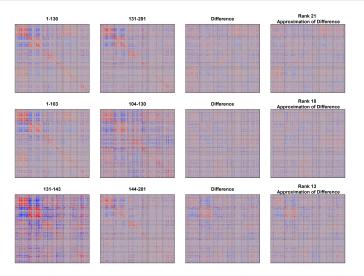


Figure: Visualization of first three steps of binary segmentation algorithm.