A parcellation scheme based on von Mises-Fisher distributions and Markov random fields for segmenting brain regions using resting-state fMRI

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2 Mathematical Formulation and modeling

Results and Inferences

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Mathematical Formulation and modeling

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- Modeling spatial correlations in rs-fMRI data to produce a parcellation even for non-contiguous regions of human brain

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- Parcellation of the cortex into subnetworks based on resting-state fMRI data opens up the possibility of developing novel functional atlases based entirely on cortical function

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- A second major limitation of current methods used to parcellate the brain is that they do not exploit spatial correlations that are inherent in rs-fMRI data

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Von Mises-Fisher Distribution

Probability Density Function

The probability density function of the von Mises–Fisher distribution for the random p-dimensional unit vector \mathbf{x} is given by $f_p(x;\mu,\kappa) = C_p(\kappa)e^{\kappa\mu^\top x}$, where $\kappa \geq 0$, $\|\mu\| = 1$ and the normalization constant $C_p(\kappa) = \frac{\kappa^{\frac{p}{2}-1}}{(2\pi)^{\frac{p}{2}}I_{\frac{p}{2}-1}(\kappa)}$, where I_{ν} denotes the modified Bessel function of the first kind at order ν .

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Relation to Isotropic Covariance MVG

Starting from a normal distribution, with isotropic covariance, $\kappa^{-1}I$, and a mean μ of length r>0 that has the density $G_p(x;\mu,\kappa)=(\sqrt{\frac{\kappa}{2\pi}})^p e^{-\kappa\frac{(x-\mu)^\top(x-\mu)}{2}}$ the Von Mises-Fisher distribution $f_p(x;r^{-1}\mu,r\kappa)$ is obtained by conditioning on $\|x\|=1$.

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- We will iterate over these steps until the energy function converges.

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- Let each T dimensional normalized voxel time series y_i given its cluster label $X_i = I$ follow T-variate VMF distribution given by $p(y_i|X_i = I, \mu_I, \kappa_I) = C(\kappa_I)e^{\kappa_I\mu^\top y_i}$

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- We model the prior distribution of unknown cluster labels X as discrete MRF given by $P(X) = \frac{1}{Z}e^{-(U_s(X)+U_l(X))}$, where $U_s(X)$ is the energy function which imposes spatial regularization and $U_l(X)$ is the label cost

• $U_s(X) = \sum_{i=1}^M \sum_{j \in N_i} V(X_i, X_j)$ where the potts potential function is defined as $V(X_i, X_j) = \beta_s (1 - \delta(X_i - X_j))$, where $\delta(z) = 1$ if z = 0 and $\delta(z) = 0$ otherwise

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Notation for this problem

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- $\hat{X} = arg min_X U$

Parameters and Labels Update

VMF Parameter Estimate

To get the optimal parameters, we maximise the likelihood. We get

$$\overline{r_l} = \frac{\sum_{\{i \mid X(i)=l\}} y_i}{N}$$
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α Expansion Algorithm

- 1. Start with an arbitrary labelling f
- 2. Set **success** := **0**
- 3. For each label $\alpha \in \mathcal{L}$
 - 3.1 Find $\hat{\mathbf{f}} = \arg\min \, \mathbf{E}(\mathbf{f}')$ among \mathbf{f}' within one α expansion of α
 - 3.2 If $\mathbf{E}(\mathbf{\hat{f}}) < \mathbf{E}(\mathbf{f})$, set $\mathbf{f} = \mathbf{\hat{f}}$ and success:=1
- 4. If success = 1 goto 2
- 5. Return f

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 - **6** Low correlation within cluster ($\rho_{in} = 0.3$) and across the clusters ($\rho_{ac} = 0.2$)

Results on Simulated Data

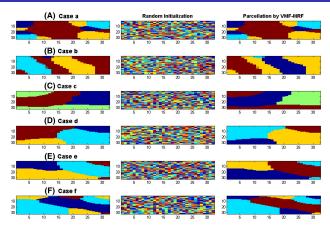


Figure: All 6 Simulated data samples

Iterations in case 4

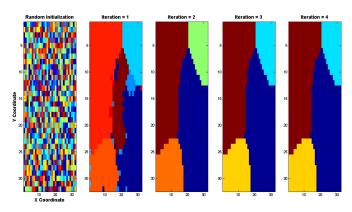


Figure: Clusters estimated by this algorithm in case 4

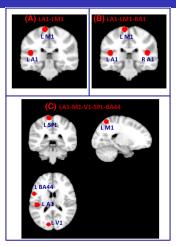


Figure: Clusters constructed from fMRI data(3 Input Data Sets)

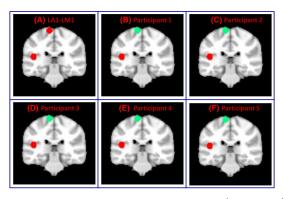


Figure: Left auditory and motor cortices (LA1-LM1)

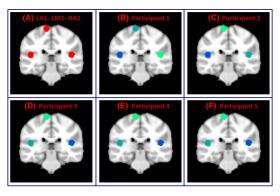


Figure: Left auditory, motor and right auditory cortices(LA1-LM1-RA1)

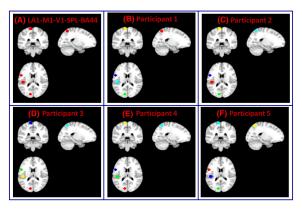


Figure: Left auditory, motor, and visual cortices, the superior parietal lobule (SPL) and inferior frontal gyrus (Brodmann Area BA 44) (LA1-M1-V1-SPL-BA44)

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- The usage of α expansion significantly improved the running time over ICM or simulated annealing based approaches. It does so by allowing updates on several voxels simultaneously instead of just one as in other cases.
- Now, since this approach worked for cases where the ground truth was already known(as depicted in the above results), we can run this algorithm to develop finer insights into functional brain segmentation

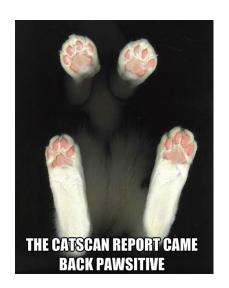
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- We expect the consensus matrix to average out the opinions of all the initializations for all voxel pairs and hence be a bimodal distribution with modes at 0 and 1.
- We could also generate updates for β_s and β_l , which may be done by several methods such as EM optimization with priors on β_s and β_l having latent variables.

In a Lighter Vein



Bibliography

- For the research paper click here
- ullet For lpha expansion click here
- For Von Mises-Fisher Distribution click here

Thanks for your attention! Any questions?

Hope you slept comfortably!