

Identifying Predictive Features in Brain Connectome Data

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Brain Connectome Data

What is brain connectome data?

Brain networks can be described as graphs where edges are between regions that "fire together".

- fMRI measures activation of voxels over time
- Calculate time dependent activation correlations
- Pick threshold correlation to define "fire together"
- Create graph

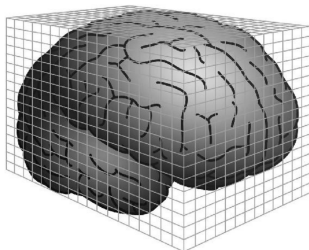


Figure 1-7: The human brain divided into voxels

(1)

Goals

Our goal is to find **novel features** of the connectome that can predict various behavioral, psychological, and demographic information.

E.g. Can we predict whether a participant is male or female from their connectome data?

Our analysis is partitioned into two themes:

- **local** connectome features (via motifs)
- **global** connectome features (via persistent homology)

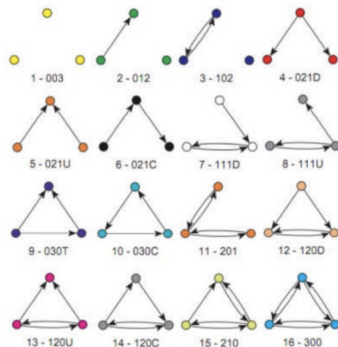
Motifs

Motifs: Sub-graphs that occur more often than would be expected in a random graph

- Triangle graph in social networks
- Triad motifs predict conscious state⁽²⁾

Motifs are going to be a proxy for local structure.

- Frequency of particular motifs
- Proportion of structural-function motifs



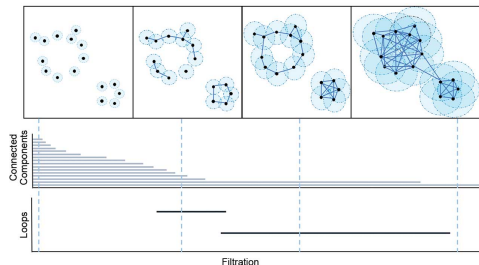
(3)

Persistent Homology

We want to learn how to abstract the global connectivity for graph learning

Process

- obtain time series data of fMRI scans
- construct graphs for each data set
- construct the node representation vectors
- identify a proper filtration
- apply persistent homology methods to obtain persistent bar codes → help expose significant features



(4)

References

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