Topological Data Analysis for Brain Networks Relating Functional Brain Network Topology to Clinical Measures of Behavior

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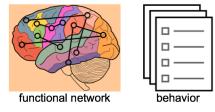
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Correlating Brain Network Topology with Autism Severity

Goal: Quantify the relationship between brain functional networks and behavioral measures.

Our Contribution: Use topological features based on persistent homology.

Result: Combining correlations with topological features gives better prediction of autism severity than using correlations alone.



Correlating Brain Network Topology with Autism Severity

About Autism Spectrum Disorders (ASD):

- No cure, causes unknown
- Diagnosis:
 - No systematic method
 - ADOS (Autism Diagnostic Observation Schedule)

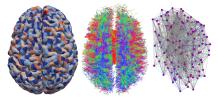
Correlate functional brain network to ADOS scores

- Early diagnosis
- Treatment tracking

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What is a Brain Network?

- Represents brain regions and pairwise associations
- Computation of Correlation Matrices:
 - Resting state functional MRI (R-fMRI)
 - Preprocessing
 - Define regions of interest (ROIs)
 - Estimate time series signals
 - Compute pairwise associations Pearson Correlation



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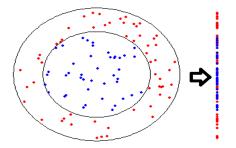
How to use this data?

- Graph and graph theoretic measures (e.g. small worldness)
 - Require binary associations (thresholding)
- Correlations as features
 - High dimensionality, not enough samples
- Dimensionality reduction: PCA, random projections
 - May lose structures in higher dimensions

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Why Topology

Projection - may lose structures in higher dimensions



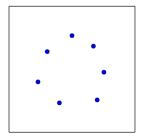
Topology captures structure

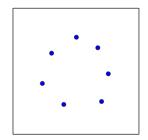
- In higher dimensions
- Across all continuous thresholds

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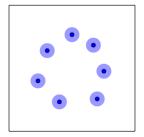
- What are topological features? Homological features:
 - Dim 0 Connected Components
 - Dim 1 Tunnels / Loops
 - Dim 2 Voids
- How to compute them (in a nutshell)?
 - Begin with point cloud
 - Grow balls of diameter t around each point
 - Track features of the union of balls as t increases

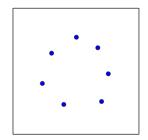
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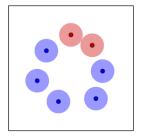


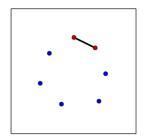
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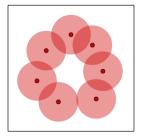


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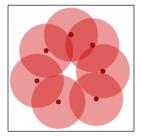


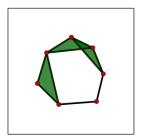
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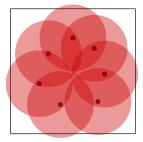


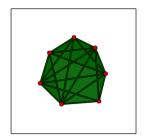
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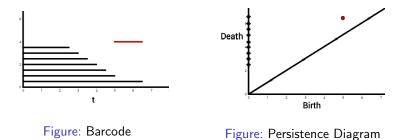
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Persistent homological features - encoded as barcodes or persistent diagrams

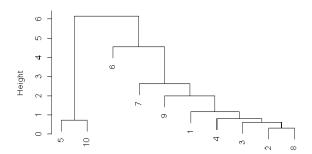


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Interpretation of Connected Components

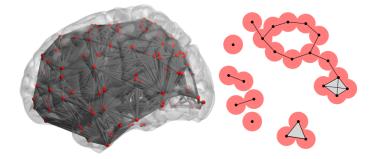
• Dim 0 features - hierarchical clustering



Cluster Dendrogram

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Computing Topological Features for Brain Networks



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A dimensionality reduction technique that finds two sets of latent dimensions from datasets X and Y such that their **projections** on the latent dimensions are **maximally co-varying**.

- X features from brain imaging: correlations, topological features (zero mean)
- Y clinical measure of behavior: ADOS scores (zero mean)

PLS models the relations between X and Y by means of **score vectors**.

PLS Regression

- *n* number of data points
- X predictor/regressor $(n \times N)$, Y response $(n \times M)$
- PLS decompose X, Y such that:

$$X = TP^T + E$$
$$Y = UQ^T + F$$

Where

- T, U latent variables/score vectors $(n \times p)$, factor matrices
- $P(N \times p)$, $Q(M \times p)$ orthogonal loading matrices
- $E(n \times N)$, $F(n \times M)$ residuals/errors
- *T*, *U* are chosen such that projections of *X*, *Y*, that is, *T* and *U*, are maximally co-varying.

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Iterative NIPALS¹ algorithm

- Find first latent dimension
 - i.e. find vectors w, c such that

$$t = Xw$$
, $u = Yc$

have maximal covariance

• Deflate previous latent dimensions from X, Y and repeat

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Kernel form of NIPALS algorithm (kPLS)

- 1. Initialize random vector u
- 2. Repeat until convergence

(a)
$$t = Ku/||Ku||$$

(b)
$$c = Y't$$

$$(c) \quad u = Yc / \|Yc\|$$

- 3. Deflate $K = (I tt^T)K(I tt^t)$
- 4. Repeat to compute subsequent latent dimensions

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- 87 Subjects: 30 Control, 57 ASD
- ADOS scores: 0 to 21
- 264 ROIs (Power regions)
- 264×264 correlation matrix.
- 34,716 distinct pairwise correlations per subject.

- Given: Correlation matrices
- Map to metric space

$$d(x,y) = \sqrt{1 - \operatorname{Cor}(x,y)}$$

- Compute persistence diagrams
- Define inner product of persistence diagrams² (i.e. kernel): Given two persistence diagrams *F*, *G*

$$k_{\sigma}(F,G) = \frac{1}{8\pi\sigma} \sum_{p \in F} \sum_{q \in G} e^{-\frac{\|p-q\|^2}{8\sigma}} - e^{-\frac{\|p-\bar{q}\|^2}{8\sigma}}$$

where for every $q=(x,y)\in$ G, $ar{q}=(y,x)$

²[Reininghaus Huber Bauer Kwitt 2015].

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Performed experiments with 3 kernels:

1. K^{Cor} - Euclidean dot product of vectorized correlations

2.
$$K^{TDA} = w_0 K^{TDA_0} + (1 - w_0) K^{TDA_1}$$

- K^{TDA_0} using only Dim 0 features
- K^{TDA_1} using only Dim 1 features

3.
$$K^{TDA+Cor} = w_0 K^{TDA_0} + w_1 K^{TDA_1} + (1 - w_0 - w_1) K^{Cor}$$

Baseline predictor - mean ADOS score

• Leave one out cross validation over parameters

- $\sigma_0, \sigma_1 (\log_{10} \sigma)$ from -8.0 to 6.0 by 0.2
- w₀, w₁ from 0.0 to 1.0 by 0.05
- k^{TDA} parameters: $\sigma_0 = -6.6$, $\sigma_1 = 1.8$, $w_1 = 0.95$
- $k^{TDA+Cor}$ parameters: $\sigma_0 = -7.8$, $\sigma_1 = 2.8$, $w_0 = 0.1$, $w_1 = 0.4$
- Compute RMSE
- Permutation test for significance

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Result Highlights:

- Baseline RMSE: 6.4302
- K^{TDA+Cor}:
 - Only method statistically significant over baseline
 - Permutation test p-value: 0.048
 - RMSE: 6.0156

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- Augmenting correlations with topological features gives a **better** prediction of autism severity than using correlations alone
- Topological features derived from R-fMRI have the **potential** to explain the connection between functional brain networks and autism severity

Many things to try

- Alternatives to correlation
- Different distance metric
- Different kernel
- Multi-site data

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Kernel Partial Least Squares Regression for Relating Functional Brian Network Topology to Clinical Measures of Behavior

Authors: Eleanor Wong, Sourabh Palande, Bei Wang, Brandon Zielinski, Jeffrey Anderson and P. Thomas Fletcher

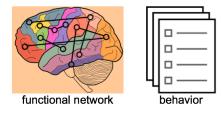
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