

Lightning Talk

Low Rank Mixup Augmentations for Contrastive Learning of Phenotypes from Functional Connectivity

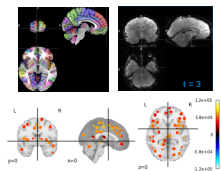
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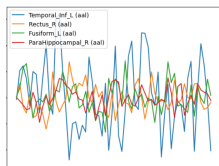
MCBIOS 2024, Emory University, March 22-24



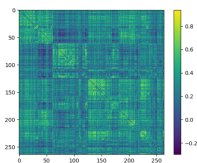
fMRI Machine Learning Pipeline



fMRI and Atlas



BOLD Timeseries



Functional Connectivity

Phenotypes



1. Age
2. Sex
3. Race
4. Schizophrenia
5. Neurodegen Disease



- How can we improve the accuracy of phenotype predictions?
- **Try contrastive learning**
 - Maximize similarity between **positive pairs**
 - Minimize similarity between **negative pairs**



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Motivation

Problem

Most contrastive learning frameworks require large numbers of subjects or data augmentations

- Most fMRI studies recruit fewer than 100 subjects
- Not clear how to augment fMRI-derived metrics such as functional connectivity (FC)

Solution

We create an augmentation strategy for FC based on mixup of the rank-1 approximation of FC (first component)

- First component is not effective for phenotype prediction



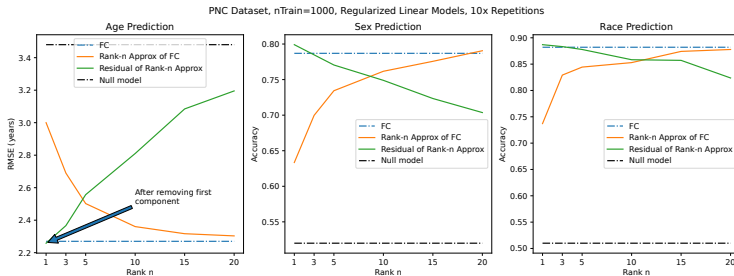
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- Construct low rank approximation to FC \mathbf{X} using the eigendecomposition
- Only keep the top N eigenvalues; set the rest to zero
- Since FC is a symmetric (PSD) matrix, it is orthogonally diagonalizable

$$\begin{aligned}\mathbf{X} &= \mathbf{V}\mathbf{\Lambda}\mathbf{V}^T \\ \hat{\lambda}_{ii} &= \begin{cases} \lambda_{ii}, & i \leq N \\ 0, & \text{otherwise} \end{cases} \\ \mathbf{X}^{(N)} &= \mathbf{V}\hat{\mathbf{\Lambda}}\mathbf{V}^T\end{aligned}\tag{1}$$

Removing the First Component

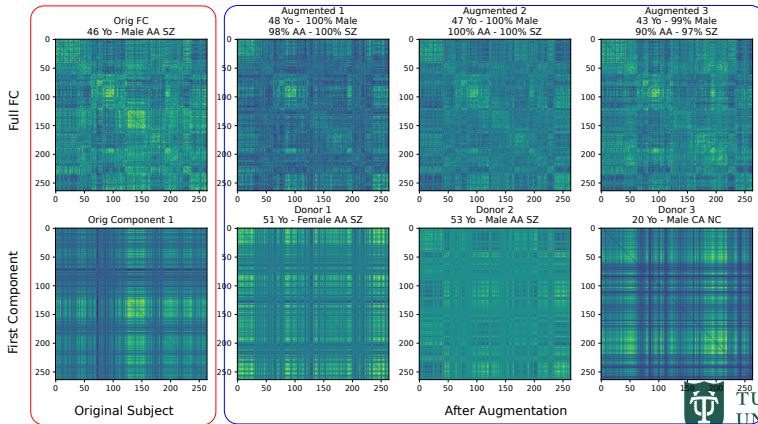
- Removing the first component doesn't reduce prediction accuracy (at all)
- Green curve at left in all graphs



Mixup Augmentations

- Use the ineffective first component in mixup augmentations to create positive pairs

Predictions are Largely Invariant to First Component of FC



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Moderately Improved Prediction Results

- Prediction results are improved 2-10% over MLP and GCN models not using augmentations
- Using PNC and BSNIP datasets

Dataset	Phenotype	Metric	MLP	GCN	CL+LRAug	p-value
BSNIP	Age	RMSE (yr)	11.82 ± 0.62	11.07 ± 0.67	10.25 ± 0.64	<0.001
BSNIP	Sex	Accuracy	68.1 ± 4.6	66.4 ± 6.1	71.7 ± 5.1	0.003
BSNIP	Race	Accuracy	76.0 ± 3.8	72.4 ± 6.1	77.7 ± 4.4	0.026
BSNIP	SZ	Accuracy	75.2 ± 3.6	71.2 ± 6.7	76.9 ± 4.4	0.017
PNC	Age	RMSE (yr)	2.62 ± 0.14	2.44 ± 0.12	2.18 ± 0.07	<0.001
PNC	Sex	Accuracy	77.9 ± 2.0	73.7 ± 9.8	79.7 ± 2.2	<0.001
PNC	Race	Accuracy	87.7 ± 1.6	86.6 ± 3.8	89.8 ± 1.9	<0.001

Additional Application to Brain Network Fingerprinting

- Removing the first component of FC helps identify same subject from different scan better than raw FC
- 97.3% identification accuracy versus 62.5%

