BISCoT: Behavior-Informed Subgroup-Consistent Connectome Template for Interpretable Brain Network Analysis

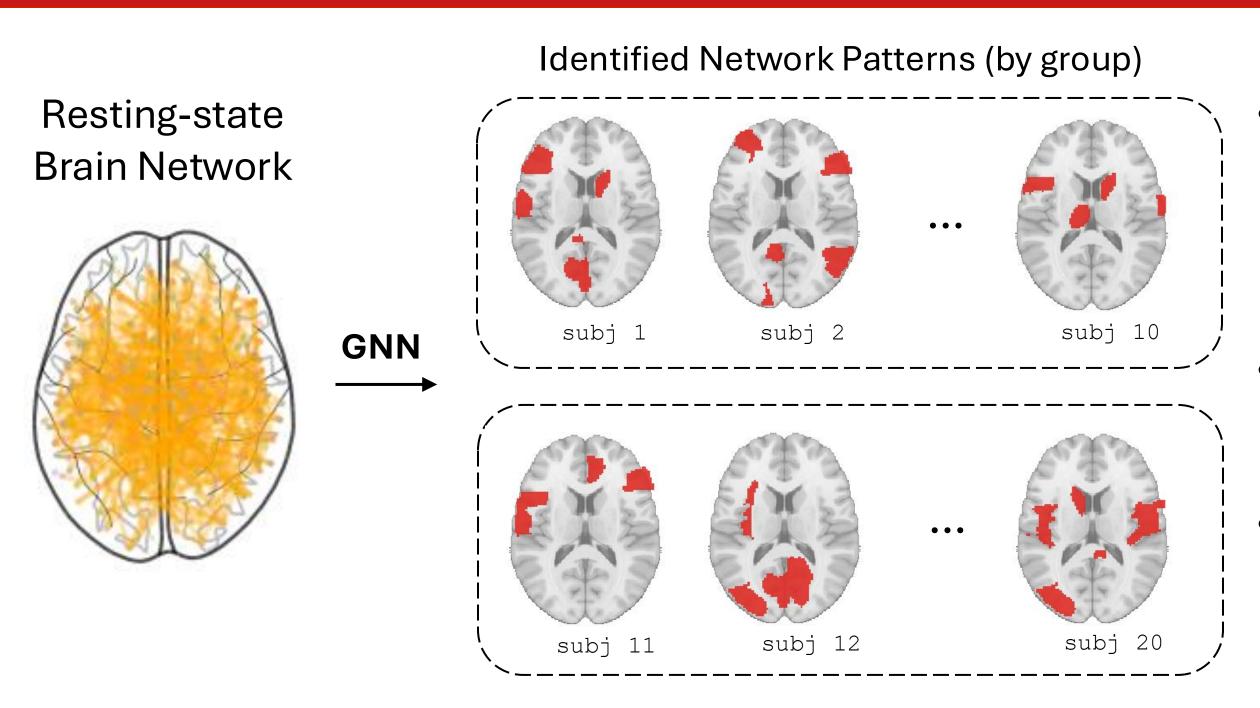


Zijian Chen¹, Stefen Beeler-Duden², Sophie Lawson², Zachary Jacokes², John Darrell Van Horn², Kevin A. Pelphrey² and Archana Venkataraman¹

¹Boston University; ²University of Virginia. email: zijianc@bu.edu



Motivation

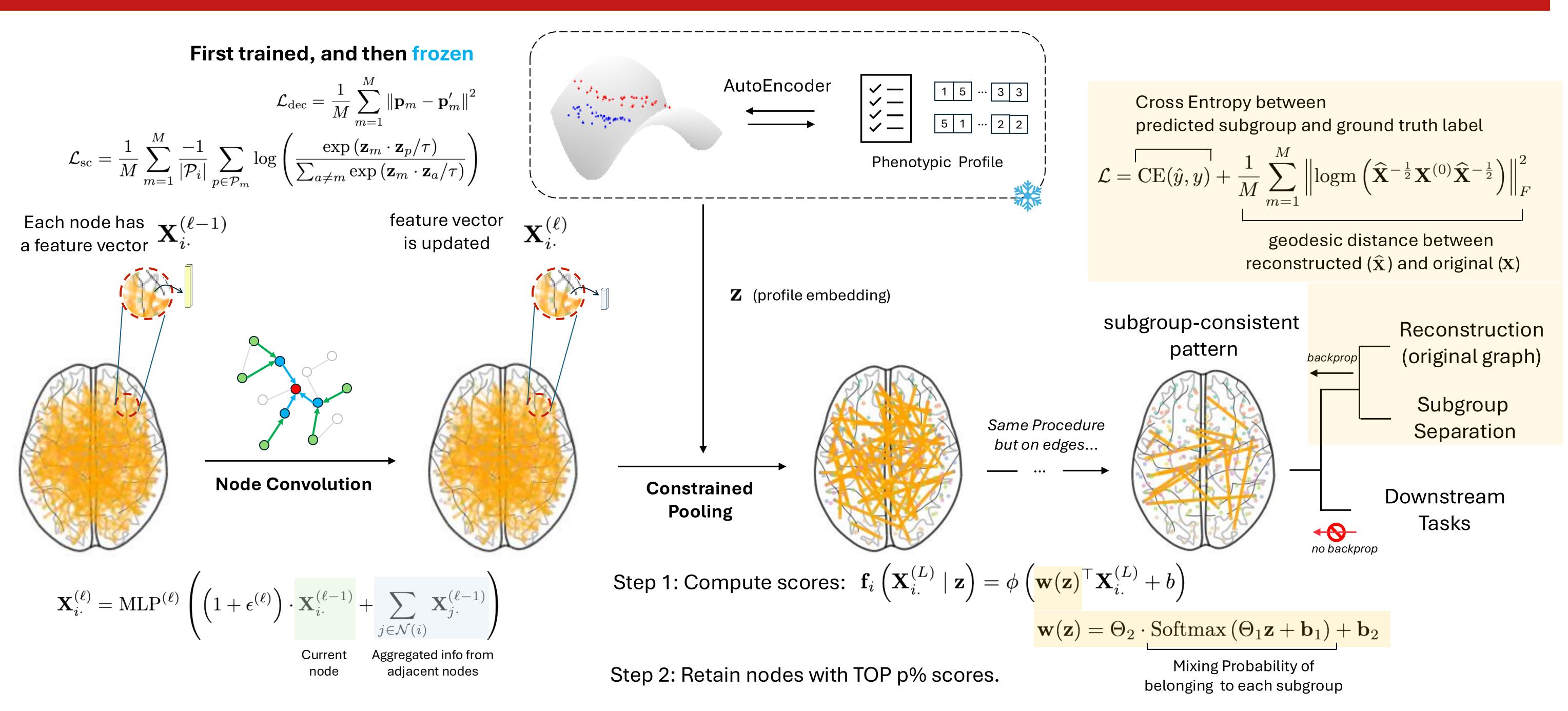


- Most GNN models do not control for a consistent feature map, nor do they account for heterogeneous phenotypes in patient cohorts
- These undermine the validity of interpretations of the identified patterns.
- These models also center at one-target prediction, limiting the generalizability of the patterns found.

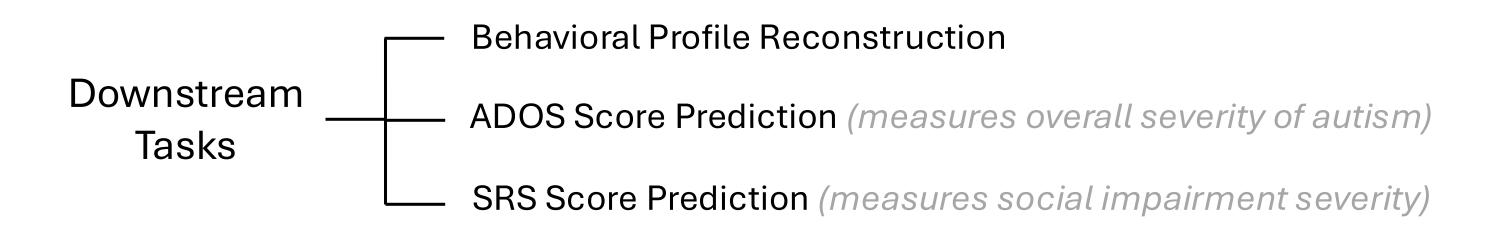
Highlights

- BISCoT is a self-supervised model for graph sparsification.
- Phenotypic data guides the pooling.
 Similar profiles lead to similar patterns
- Consistent pattern across subjects with similar phenotypes
- Better generalizability to various downstream tasks

Our Solution: BISCoT



Experiments and Results



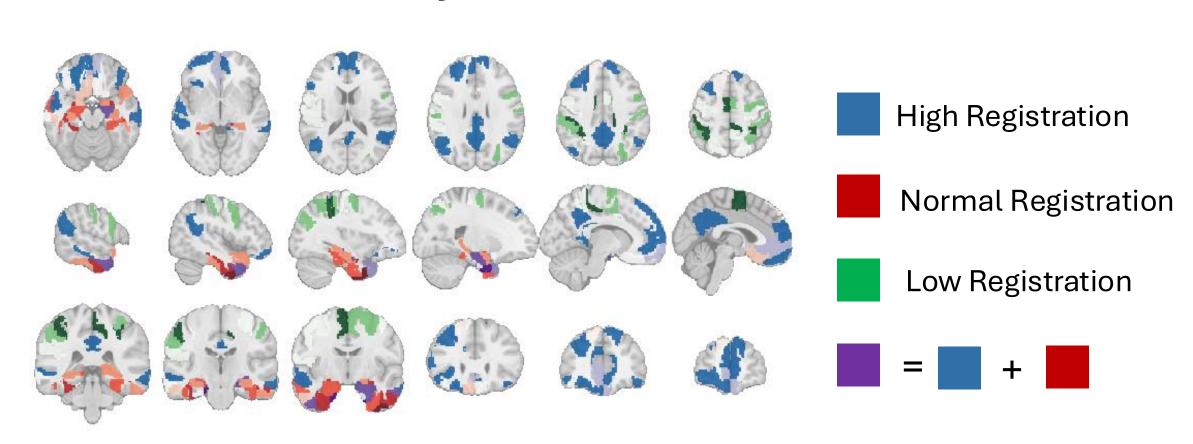
Original Network Reconstruction Performance

Method	Geodesic ↓	Frobenius ↓	Separation †	EigDev
BISCoT	112.48 ± 9.28	65.55 ± 4.04	0.77±0.12	0.35±0.07
GIN+TopK	181.09 ± 2.97*	65.64 ± 4.62	0.59 ± 0.07 *	0.37 ± 0.09
GIN+SAGPool	180.55 ± 0.98*	67.94 ± 5.79*	0.60 ± 0.11 *	0.42 ± 0.13 *
GIN+EdgePool	179.51 ± 2.22*	79.16 ± 3.13*	0.56 ± 0.10 *	0.61 ± 0.07 *
GraphSAGE+TopK	179.12 ± 1.64*	66.89 ± 4.66	0.52 ± 0.14 *	0.41 ± 0.10*

Downstream Tasks Performance

Method		Profile (0-60)		ADOS (0-30)		SRS (0-100)	
		MAE	RMSE	MAE	RMSE	MAE	RMSE
	BISCoT	4.53±0.48	5.80±0.59	3.58±0.83	4.77±0.69	13.43±1.29	16.19±0.92
GIN:	GIN+TopK	7.15±0.82*	9.20±1.26*	4.10±1.08	5.38±1.43	16.81±1.87*	20.93±2.50*
(GIN+SAGPool	6.79±0.75*	9.00±0.82*	4.27±0.51*	5.36±0.81*	23.93±2.87*	27.78±3.01*
C	SIN+EdgePool	6.90±0.83*	9.10±0.91*	4.62±1.44*	5.69±1.77*	23.47±4.94*	27.59±0.93*
Gr	aphSAGE+TopK	6.74±0.55*	8.63±0.76*	4.72±0.47*	5.92±0.59*	25.03±2.84*	28.91±2.89*
SOTA	BrainGNN	6.25±0.78*	7.97±0.99*	3.77±0.82	4.80±0.97	16.34±1.92*	20.97±1.98*
	MGCN	5.14±0.64*	6.54±0.79*	4.20±0.69	5.09±0.85	14.01±1.54	18.24±1.81
	BPI-GNN	19.19±1.31*	20.71±9.21*	7.21±0.71*	8.44±3.65*	18.45±4.46*	26.54±9.89*

Patterns identified by BISCoT:



Appear in at least 85% of the patients in each subgroup!





