#### Subgraph Neural Networks

#### Emily Alsentzer, Samuel G. Finlayson, Michelle M. Li, Marinka Zitnik

#### Presenter: Arshdeep Sekhon https://qdata.github.io/deep2Read

#### Subgraph Property Prediction

- Given Graph G = (V, E) and subgraph G' = (V', E') where  $V' \subseteq V$  and  $E' \subseteq E$ .
- Each subgraph S has a label y<sup>S</sup> and many S<sup>C</sup> which is a set of nodes in S that are connected to each other by a path.
- The task : if a subgraph has a specific property or not Graph G: subgraphs defined by node membership



Figure: colors indicate labels

Why are subgraph property prediction challenging?

- make joint predictions over larger structures of varying sizes: do not correspond to simple k-hop, possibly disconnected and far off
- higher-order connectivity patterns: how nodes within the subgraph interact and how they interact with nodes outside the subgraph (border and extenral nodes)
- subgraphs can be localized within a region of the graph or spread out: learn about the subgraph positions within the graph

subgraphs share edges and non edges

# Formulating Subgraph Prediction

#### subgraph problem

Given subgraphs  $(S_1, \ldots, S_n)$  the task is to get embeddings  $z_S \in R^{d_S}$  for every subgraph S. SUB-GNN uses a GNN to learn a classifier  $f : S \to \{1, \ldots, C\} f(S) = \hat{y}_S$ .

Difference from other gnns: operates directly on components

# Subgraph Properties to encode

network properties that are not necessarily defined for either nodes or graphs.

		-	
SUB-GNN Channel	SUB-GNN Subchannel		
	Internal (I)	Border (B)	
Position (P)	Distance between $S_i$ 's components	Distance between $S_i$ and rest of $G$	
Neighborhood (N)	Identity of $S_i$ 's internal nodes	Identity of $S_i$ 's border nodes	
Structure (S)	Internal connectivity of $S_i$	Border connectivity of $S_i$	

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#### subgraph properties: position

- border position: distance to the rest of G
- ▶ internal: distance between the components of G

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# subgraph properties: neighborhood

border neighborhood: nodes within k-hops of any node in S, each component has its own border neighborhood

internal neighborhood

#### subgraph properties: structure

- internal : internal connectivity of each subgraph
- border: edges connecting internal nodes to border neighborhood

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sub graph level message passing: Anchor Patches

$$\blacktriangleright \mathbb{A} = (A^1, \dots, A^Q)$$

anchor patches are subgraphs sampled from G specific to each channel : P, N and S

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sub graph level message passing: Anchor Patches to subgraph components

$$\blacktriangleright \mathbb{A} = (A^1, \dots, A^Q)$$

anchor patches are subgraphs sampled from G specific to each channel : P, N and S

$$\blacktriangleright MSG_{X,C} = \gamma_x(A_X, S^C)p_X$$

•  $\gamma$  is a similarity function for channel X

• 
$$\boldsymbol{a}_{X,c} = AGG_M(MSG_X(S^C, A_X, p_X)) \forall A_X in \mathbb{A}_X$$
  
•  $\boldsymbol{h}'_{X,c} = \sigma(\boldsymbol{W}_h[\boldsymbol{a}_{X,c}; \boldsymbol{h}'^{-1}_{X,c}])$ 

#### Property-aware output representations

- a matrix *M<sub>X</sub>* where each row is an anchor set message computed by *MSG<sub>X</sub>*
- **>** pass through a non linear activation function to get  $z_{x,c}$

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► for neighborhood: use  $z_{N,c} = h_{N,c}$ 

#### Agregating Property-aware output representations

- $\triangleright$   $z_{x,c}$  for a channel x and a subgraph component c
- First aggregate using channel aggregator AGG<sub>C</sub>
- Then aggregate using layer aggregator AGG<sub>L</sub>
- now we have z<sub>c</sub>
- **•** READOUT from  $z_c$  to  $z_s$
- Finally, SUB-GNN routes messages for internal and border properties (i.e., {P<sub>I</sub>, P<sub>B</sub>}, {N<sub>I</sub>, N<sub>B</sub>}, {S<sub>I</sub>, S<sub>B</sub>}) within subchannels for each channel P, N and S, and concatenates the final outputs

# A, $p_X$ , $\gamma$

- Sampling anchor patches
- Neural encoding of anchor patches
- Estimating similarity of anchor patches.

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## Sampling anchor patches

$$\blacktriangleright \phi_X : (G, S^C) \to A_X$$

- ► Internal position sampler node u<sub>P</sub> ∈ S, shared across all components in S
- ▶ Border position sampler node  $v_P \in G$  shared across all S
- ▶ Neighborhood internal sampler node  $u_N \in S^C$
- ▶ Neighborhood border sampler node  $v_N \in k hop \text{ of } S^C$
- structure anchor sampler : connected component sampled from G via triangular random walks

#### Neural encoding of anchor patches

For position and neighborhood anchor patches, same as initial node embeddings :

$$\psi_{N} = \psi_{p} = N_{i} \tag{1}$$

For structure nodes:

$$\psi_{\mathcal{S}}: \mathcal{A}_{\mathcal{S}} \to \boldsymbol{p} \in \mathbb{R}^d \tag{2}$$

- w fixed length triangular random walks (u<sub>πw(1)</sub>,..., u<sub>πw(n)</sub>)
   The triangular random walk samples triangular successors with probability β and non-triangular successors with probability
  - $1-\beta$ .
- input to LSTM, use sum of hidden states = p

## Neural encoding of anchor patches

For structure nodes, random walk strategy:

- internal: random walks over set  $I \{u | u \in A_s\}$
- neighborhood: random walks over set N {v | v ∉ A<sub>s</sub>} limited to neighborhood k hops

- ▶ border: random walks over set  $\{u | u \in I, v \in N, uv \in E\}$
- multiple random walks but single p

Estimating similarity of anchor patches and subgraph components

- similarity between subcomponent of subgraph and anchor patch
- $\blacktriangleright \gamma_X : (S^C, A_S) \to [0, 1]$
- for the position channel,  $\gamma_P = \frac{1}{d_{SP}(A_S, S_C) + 1}$

*d<sub>SP</sub>* is the shortest path between connected components *S<sup>C</sup>* and anchor path *A<sub>S</sub>*

 for structure channel, use the normalized Dynamic Time Warping (DTW)<sup>1</sup>

► d<sub>A<sub>S</sub></sub>, d<sub>S<sub>C</sub></sub>: ordered degree sequences for the subgraph component and anchor patch

# Algorithm summary

#### Algorithm 1: SUBGRAPH NEURAL NETWORK.

**Input:** Graph G = (V, E); Node representations  $\{\mathbf{x}_u | u \in V\}$ ; Subgraph S consisting of connected components  $S^{(C)}$  for c = 1, ..., R; Channels N, S, and P corresponding to neighborhood, structure, and position; Subchannels I and B corresponding to internal and border subgraph topology; Anchor patch sampling function  $\phi_X : (G, S) \to A_X$ ; Anchor patch encoder  $\psi_X : A_X \to \mathbb{R}^d$ ; Trainable weight matrices  $\mathbf{W}_{x,z}^{(l)}$  and  $\mathbf{W}_{x,z}^{(l)}$  for each layer  $l \in [1, L]$  and each channel x; Nonlinear activation function  $\sigma$ .

**Output:** Subgraph representation  $h_S$  for subgraph S

$$\begin{array}{ll} \mathbf{z}_c^0 = \sum_{u \in S^{(C)}} \mathbf{x}_u \\ \mathbf{h}_{X,c}^0 = \mathbf{z}_c^0 \text{ for channel } \mathbf{X} \in \{\mathsf{N},\mathsf{S},\mathsf{P}\} \end{array} \qquad \qquad // \text{ Channel-independent initialization} \\ \end{array}$$

æ // Aggregate components

## Sub-GNN Figure



# Computational Complexity and model extensions

- function of (number of subgraphs, size of the subgraphs)
- also depends on number of anchor patches: prespecified and fixed
- possible to use other types of similarity or joint learning of node embeddings

## Synthetic Experiments

- subgraph properties: density, cut ratio, coreness, component
- density: internal structure(250 subgraphs of size 20)
- cut ratio: border structure(250 subgraphs of size 20)
- coreness: average core number of the subgraph, tests border structure and position (221 subgraphs of size 20)
- component: the number of subgraph components,(250 subgraphs with 15 nodes per component) tests internal and external position.

#### Real World Datasets

- PPI-BP : 1591 subgraphs with 6 labels, labeled using Biological Process Ontology from MSigDB, Subgraphs are collections of proteins in the PPI network that are involved in the same biological process
- HPO-METAB : graphs of causal genes and symptoms, with subgraphs defined by symptoms. 2400 subgraphs with 6 labels from metabolic disorders: lysosomal, energy, amino acid, carbohydrate, lipid, and glycosylation.
- HPO-NEURO : about neurological disorders
- EM-USER : subgraphs about work out routines with 1343 sugraphs. Label is gender

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## HPO dataset challenges

- distinguishing subcategories of similar diseases (a challenge for averaging-based methods),
- exhibit class distributional shift between train and test, and have been designed to require inductive inference to nearby phenotypes using edges in the graph.
- require distinguishing subcategories of similar diseases (a challenge for averaging-based methods),

#### **Baselines**

- AVG: average of the node embeddings of the subgraph
- MN-GIN and MN-GAT: use virtual node to represent a subgraph
- ► s2v-N, s2v-S, s2v-NS: suhgraph 2 Vec
- GC: treat each subgraph as standalone graph using average of node embeddings

pretrained: GIN on link prediction

# Simulation microF1

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Method	DENSITY	CUT RATIO	CORENESS	COMPONENT
SUB-GNN (Ours)	$0.919{\scriptstyle\pm0.016}$	$0.629{\scriptstyle\pm0.039}$	0.659±0.092	0.958±0.098
Node Averaging	$0.429{\scriptstyle\pm0.041}$	$0.358{\scriptstyle\pm0.055}$	$0.530 \pm 0.050$	$0.516 \pm < 0.001$
Meta Node (GIN)	$0.442{\scriptstyle\pm0.052}$	$0.423{\scriptstyle\pm0.057}$	$0.611 \pm 0.050$	$0.784{\scriptstyle \pm 0.046}$
Meta Node (GAT)	$0.690{\scriptstyle\pm0.021}$	$0.284{\scriptstyle\pm0.052}$	$0.519 \pm 0.076$	$0.935 \pm < 0.001$
Sub2Vec Neighborhood	$0.345{\scriptstyle\pm0.066}$	$0.339{\scriptstyle \pm 0.058}$	$0.381 \pm 0.047$	$0.568 \pm 0.039$
Sub2Vec Structure	$0.339{\scriptstyle\pm0.036}$	$0.345{\scriptstyle\pm0.121}$	$0.404 \pm 0.097$	$0.510 \pm 0.013$
Sub2Vec N & S Concat	$0.352 {\pm} 0.071$	$0.303{\scriptstyle\pm0.062}$	$0.356 \pm 0.050$	$0.568{\scriptstyle\pm0.021}$
Graph-level GNN	$0.816{\scriptstyle \pm 0.068}$	$0.377{\scriptstyle \pm 0.058}$	0.419±0.070	$0.526{\scriptstyle \pm 0.081}$

# Real World microF1

Method	PPI-BP	HPO-NEURO	HPO-METAB	EM-USER
SUB-GNN (Ours)	0.324±0.013	0.632±0.010	0.537±0.023	0.751±0.021
Node Averaging	$0.289{\scriptstyle \pm 0.043}$	$0.490 \pm 0.059$	$0.443 \pm 0.063$	$0.744 \pm 0.086$
Meta Node (GIN)	$0.277 \pm 0.040$	$0.233{\scriptstyle \pm 0.086}$	$0.151 \pm 0.073$	$0.550 \pm 0.025$
Meta Node (GAT)	$0.308{\scriptstyle\pm0.032}$	$0.259 \pm 0.063$	$0.138{\scriptstyle \pm 0.034}$	$0.536 \pm 0.047$
Sub2Vec Neighborhood	$0.309{\scriptstyle\pm0.023}$	$0.211 \pm 0.068$	$0.132 \pm 0.047$	$0.503 \pm 0.035$
Sub2Vec Structure	$0.307{\scriptstyle\pm0.013}$	$0.223 {\pm} 0.065$	$0.124 \pm 0.025$	$0.742 \pm 0.023$
Sub2Vec N & S Concat	$0.295{\scriptstyle\pm0.011}$	$0.206 \pm 0.073$	$0.114 \pm 0.021$	$0.536 \pm 0.047$
Graph-level GNN	$0.291 \pm 0.026$	$0.577 \pm 0.015$	$0.480 \pm 0.026$	$0.505 \pm 0.04$

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# Channel Ablation Analysis

#### aligns with their inductive biases

SUB-GNN Channel	DENSITY	CUT RATIO	CORENESS	COMPONENT
Position (P)	0.758±0.046	0.516±0.083	0.581±0.044 🗸	0.958±0.098 🗸
Neighborhood (N)	$0.777 \pm 0.057$	0.313±0.087	$0.485{\scriptstyle \pm 0.075}$	$0.823 \pm 0.089$
Structure (S)	0.919±0.016 🗸	0.629±0.039 🗸	0.663±0.058 🗸	0.600±0.170
All (P+N+S)	0.894±0.025	$0.458 \pm 0.101$	$0.659{\scriptstyle\pm0.092}$	$0.726 \pm 0.120$