

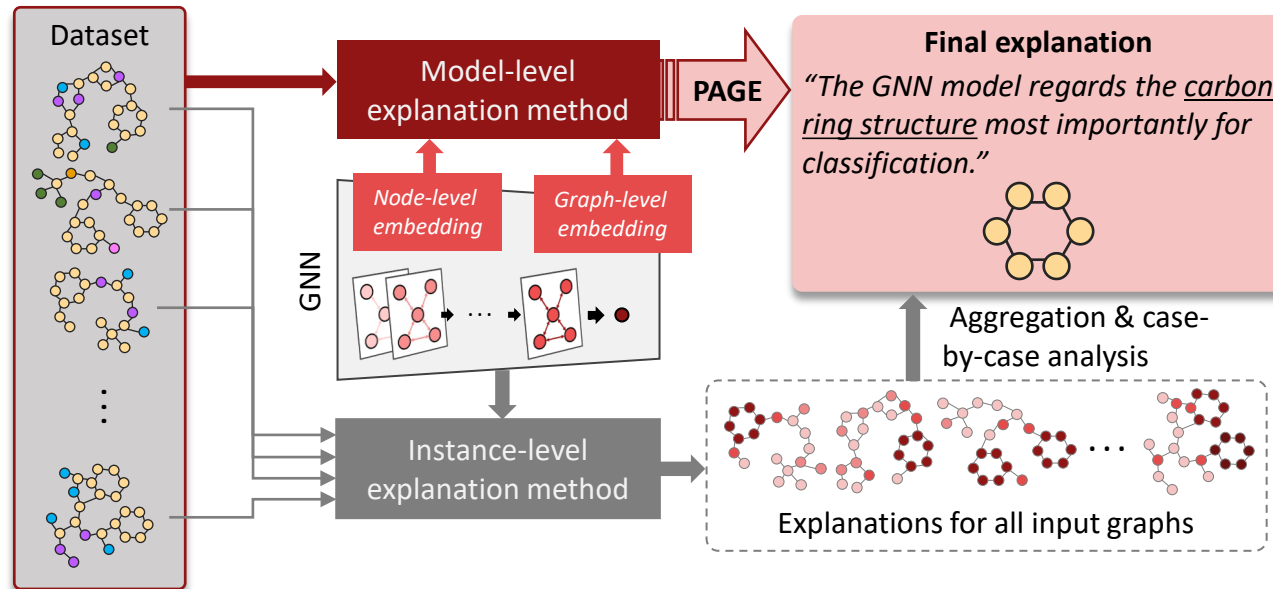
PAGE: Prototype-Based Model-Level Explanations for Graph Neural Networks

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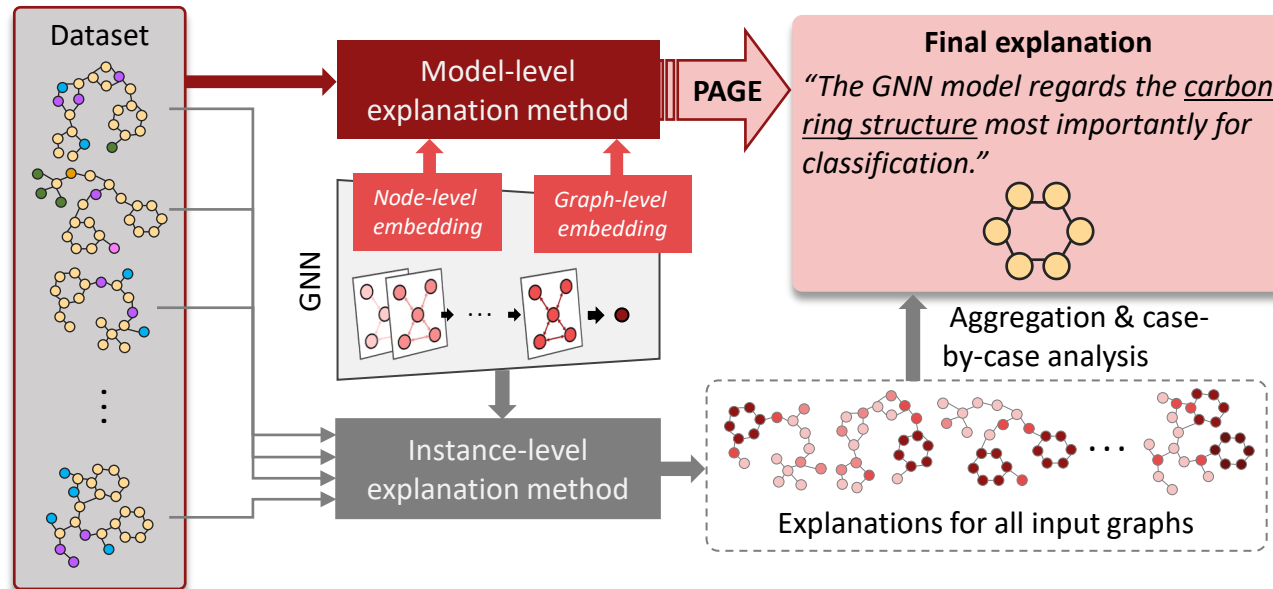


Model-level explanation



- **Model-level explanations** explain the decisions of a deep neural network model by finding the pattern that leads to a specific decision.
- Previous instance-level explanations are not fit for such type of explanations, often requiring aggregating explanations or large-scale case-by-base analysis.

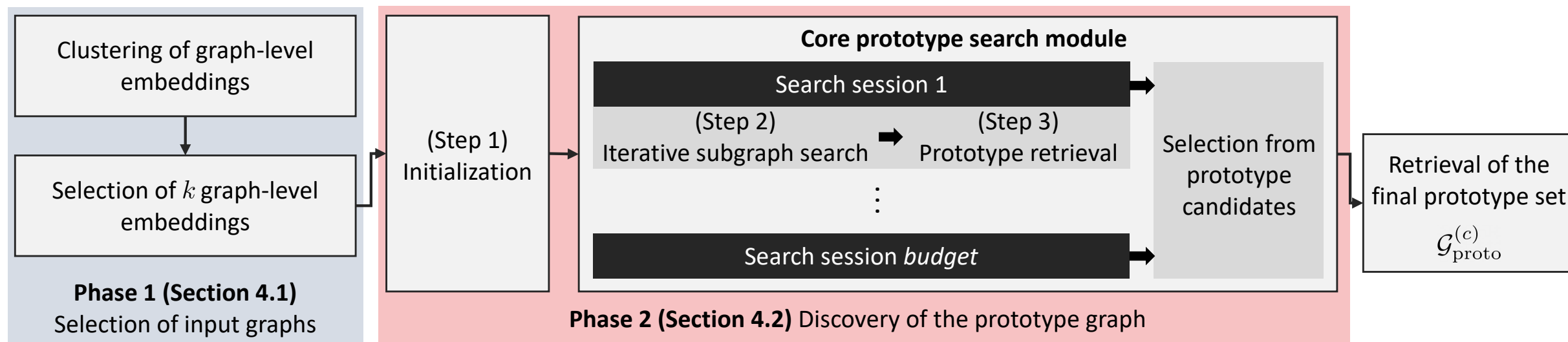
Model-level explanations of graph neural networks (GNNs)



Explanations for GNNs have mostly focused on instance-level explanation



We propose PAGE, a **model-level** explanation method for graph classification GNN models.



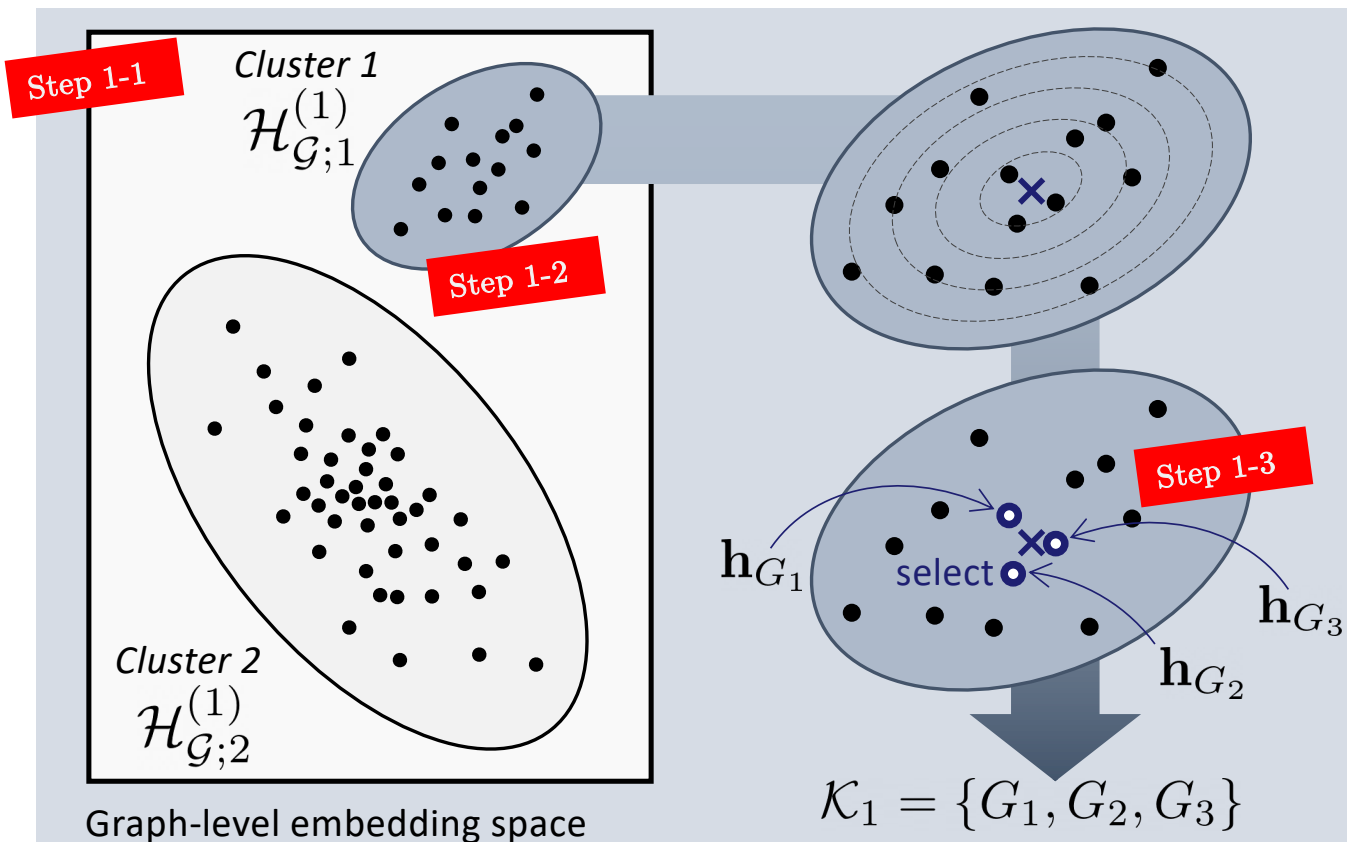
→ Main objective of Phase 1: **Reduce the search space** of the prototype by selecting a small subset of input graphs to work on.

→ In Phase 2, we find the prototype graph from the graphs from Phase 1.

→ Main approach: Search common subgraph **guided by** the embedding / representation vectors from the underlying GNN

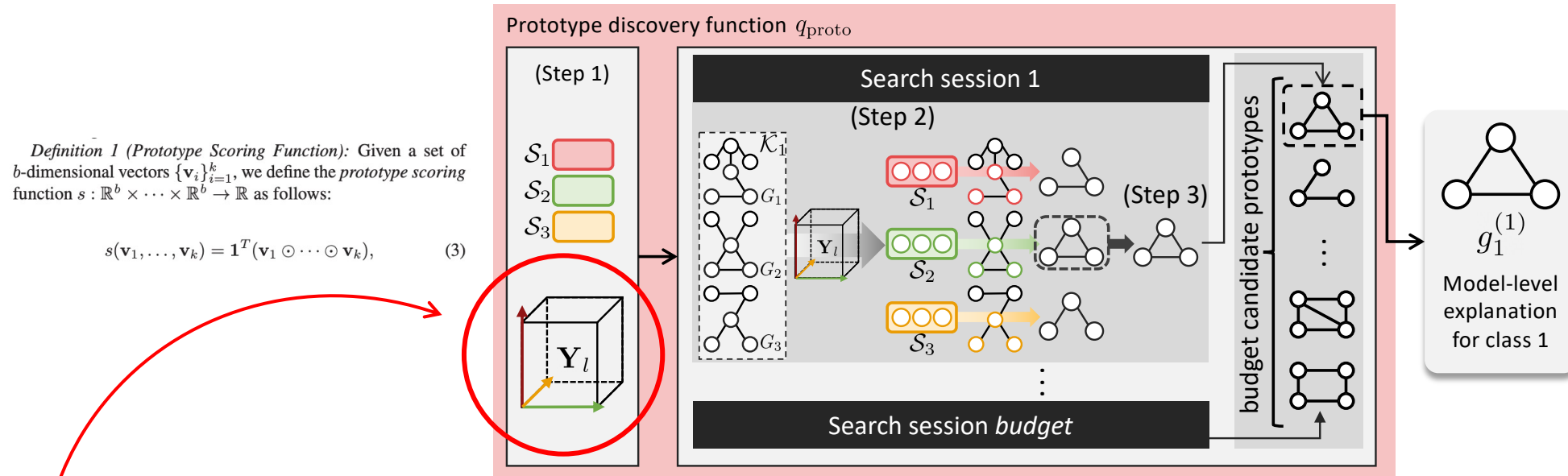
1. Step 1: Pre-compute all node-tuple-wise similarity scores
2. Step 2 (**Core prototype search module**): Extract common subgraph pattern via searching for graph patterns with highest overall similarity score

Phase 1: Selection from **graph-level** embeddings



- (Step 1-1) Use the graph dataset to acquire **graph-level** embeddings for all input, where it contains semantic information learnt from the GNN.
- (Step 1-2) Use clustering to capture different sub-pattern within each class.
- (Step 1-3) Only select nodes that are **near the centroid** for each cluster.

Phase 2: Prototype discovery with Prototype Scoring Function

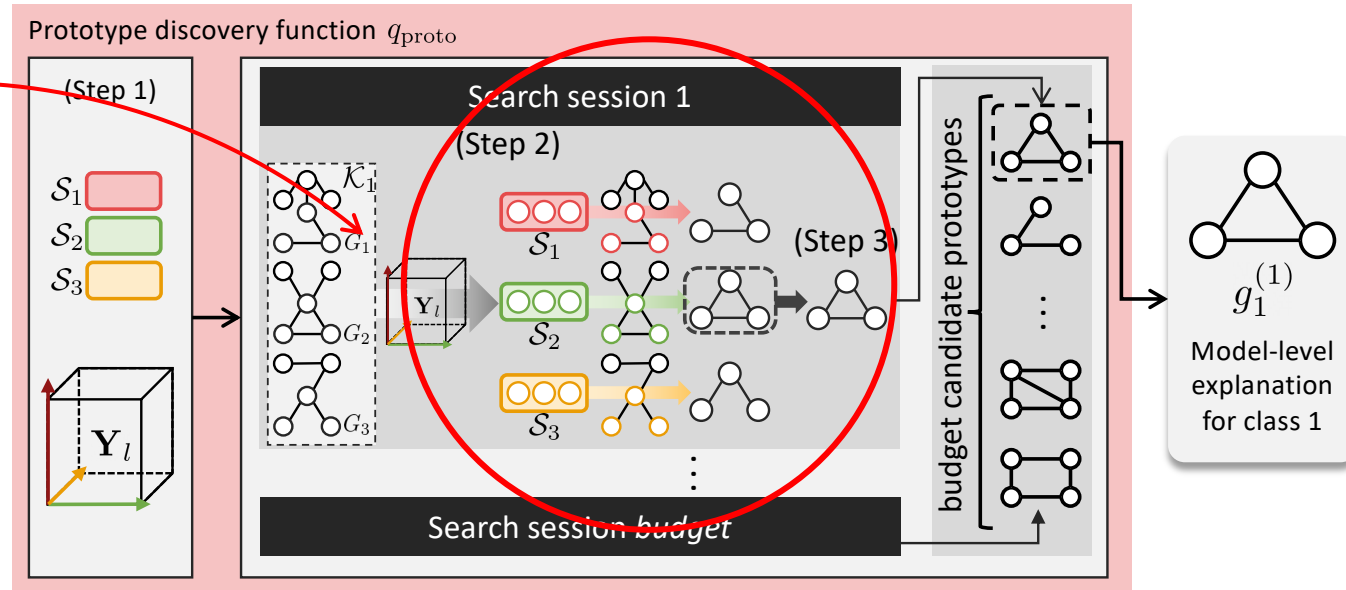


Objective: We want the nodes of the **final prototype graph** to have a **strong alignment** with the nodes of the k selected graphs.

Question: How can we efficiently capture node-wise similarity between all k selected graphs, which is to be used during the discovery process?

Solution: Define a new Prototype Scoring Function, and pre-compute a **similarity tensor** that captures all node-wise alignments from **node-level** embeddings from each k graphs

Phase 2: Prototype discovery with Prototype Scoring Function



Objective: We want the nodes of the **final prototype graph** to have a **strong alignment** with the nodes of the k selected graphs.

Question: Given the node-level similarity scores, how do we extract the common graph pattern?

Solution: Perform an **iterative walk** for each k graph **simultaneously**, selecting the nodes that has the **highest similarity scores as the next node** during the process.

Datasets for explainable AI on graphs

*TP: Test performance of GNN

Dataset	n	$\sum_i \mathcal{V}_i$ (Avg.)	$\sum_i \mathcal{E}_i$ (Avg.)	TP	Ground-truth explanations
BA-house	2,000	21,029 (10.51)	62,870 (31.44)	1.000	Ground-truth explanations: Small subgraph motifs (house-shaped, grid-shaped)
BA-grid	2,000	29,115 (14.56)	91,224 (45.61)	0.9583	
Benzene	12,000	246,993 (20.58)	523,842 (43.65)	0.9444	Chemical molecule groups (Real-world knowledge-based explanations)
MUTAG	4,337	131,488 (30.32)	266,894 (61.54)	0.7247	
Solubility	708	9,445 (13.34)	9,735 (13.75)	0.8717	
MNIST-sp	70,000	5,250,000 (75)	41,798,306 (696.63)	0.7595	Visual semantics

Qualitative evaluation: Comparison against ground-truth explanations

	BA-house	BA-grid	Benzene	MUTAG	Solubility (Class 0)	Solubility (Class 1)
Ground truth prototype						
XGNN						
PAGE						
● Head ● Body ● None ● Carbon ● Oxygen ● Nitrogen ● Chlorine ● Iodine ● Phosphorus ● Bromine ● Hydrogen ● Fluorine ● Sulfur						

Compared to XGNN [1], PAGE is able to extract explanations that much more resembles the ground-truth explanations (prototypes) for each dataset (synthetic and real-world).

Quantitative evaluation: Accuracy, Density, Consistency, Faithfulness

Method	BA-house	BA-grid	Solubility	MUTAG	Benzene	MNIST-sp
PAGE	0.5238	0.8571	0.3290	0.9090	0.6667	N/A
XGNN	0.2500	0.3200	0.2341	0.6875	0.2500	N/A

(a) Accuracy ($\uparrow$).

Method	BA-house	BA-grid	Solubility	MUTAG	Benzene	MNIST-sp
PAGE	0.0308	0.0615	0.0846	0.1216	0.0639	0.1025
XGNN	0.2152	0.2705	0.3213	0.1269	0.2227	0.0242

(c) Consistency ($\downarrow$).

Method	BA-house	BA-grid	Solubility	MUTAG	Benzene	MNIST-sp
PAGE	0.1481	0.1563	0.0462	0.1389	0.1111	0.2195
XGNN	0.1235	0.1667	0.1195	0.1875	0.1094	0.3593

(b) Density ($\downarrow$).

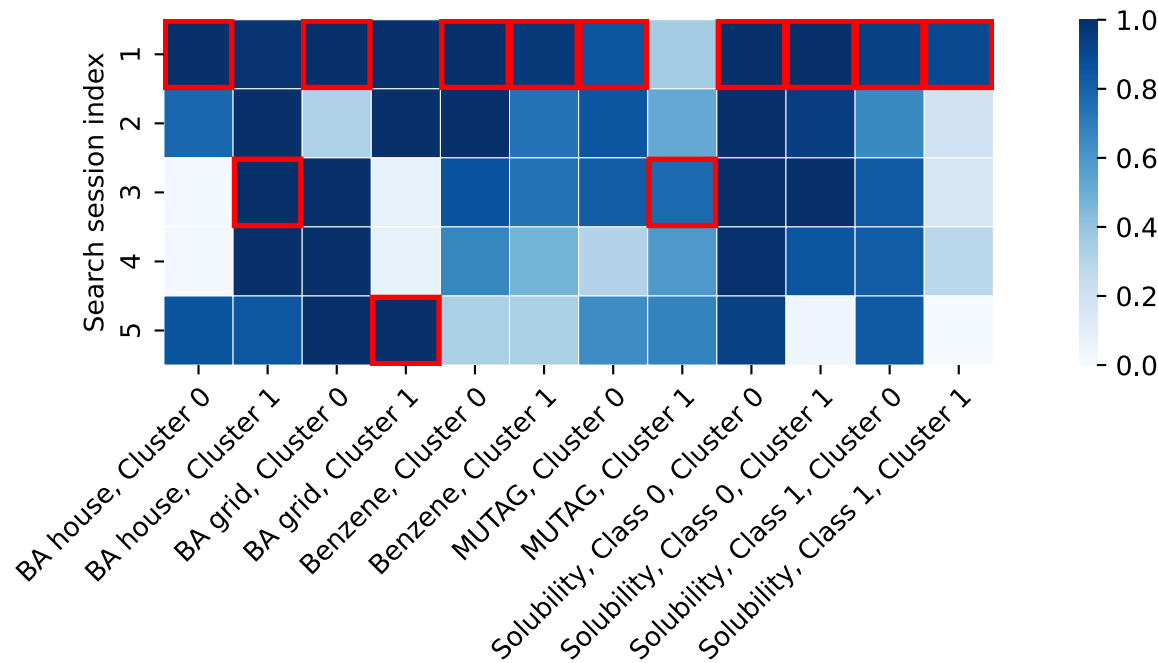
Method	BA-house	BA-grid	Solubility	MUTAG	Benzene	MNIST-sp
PAGE	0.7340	0.5636	0.2164	0.4430	0.2364	0.8182
XGNN	-0.4037	-0.1636	0.0983	0.2504	-0.3091	0.1273

(d) Faithfulness ($\uparrow$).

For 4 different metrics, PAGE shows clear superiority against XGNN for all 6 datasets.

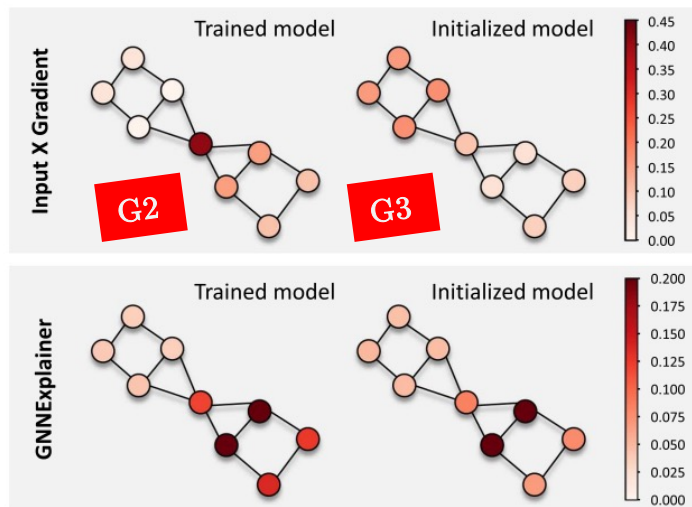
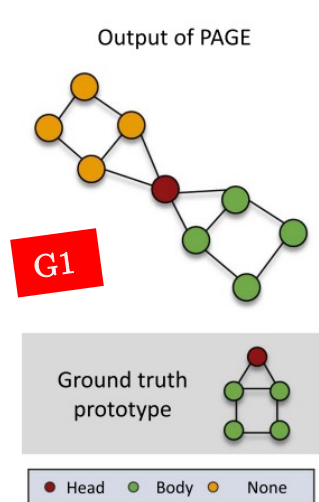
- **Accuracy:** How much the prototype overlaps with the ground-truth explanation?
- **Density:** How much each method produces explanations with lower graph density? (i.e., sparsity)
- **Consistency:** How much robust is each method for different GNN settings (e.g., # of hidden dimensions)?
- **Faithfulness:** How much the quality of the explanation correlate with the actual performance of the GNN?

How much search trials are required?



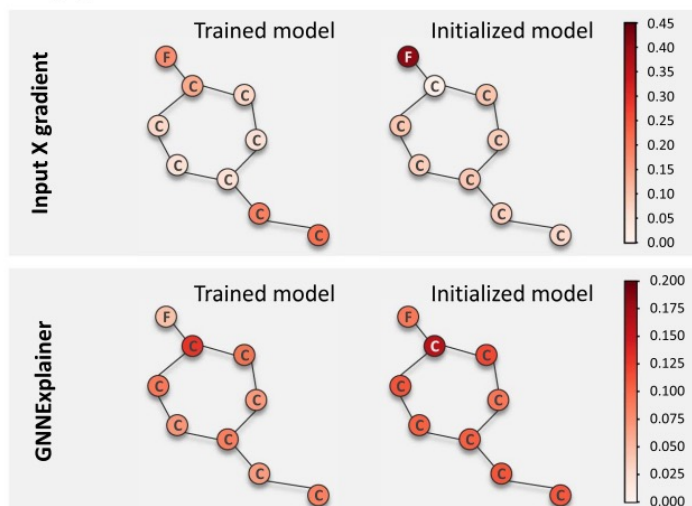
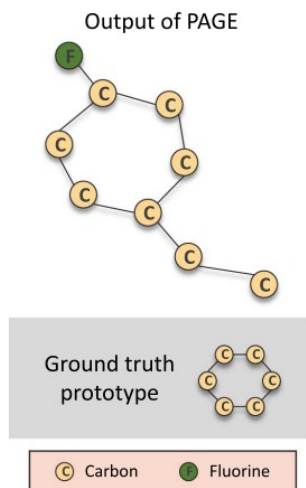
- During phase 2, we implement a strategy that involves performing multiple attempts of extraction (with different starting points).
- Empirically, we find that PAGE can find the **best result** in the **first try** for most of the time.

How much does instance-level methods agree with our results?



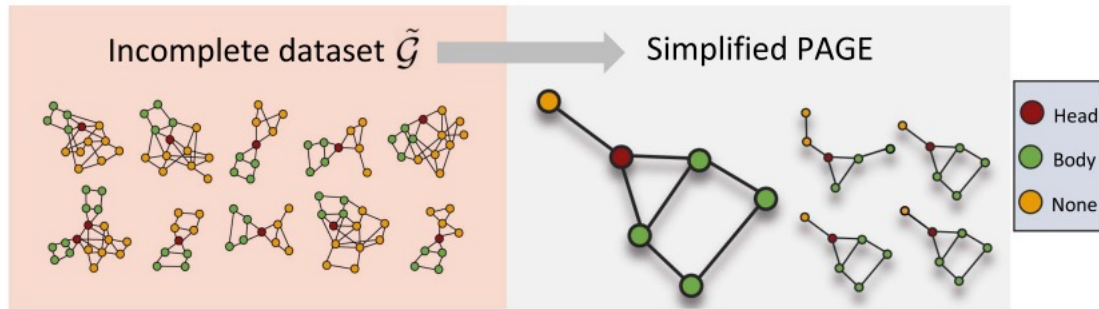
(a) BA-house.

- We also explore the alignment between PAGE and other well-known instance-level explanation methods (i.e., Input X Gradient, GNExplainer)
- We ask each instance-level explanation: How much do you think the explanation of PAGE includes the important parts in terms of explanation?
- Procedure:
 - Get **G1** from PAGE.
 - Select an instance-level explanation method.
 - Ask to explain **G1** as a heatmap (**G2**).
 - Also ask the same when the GNN is not trained (**G3**), which is considered as a 'baseline'.
- Results show that instance-level methods generally **agree** that the prototypes of **PAGE** does include **essential subgraph patterns** learned by the GNN.

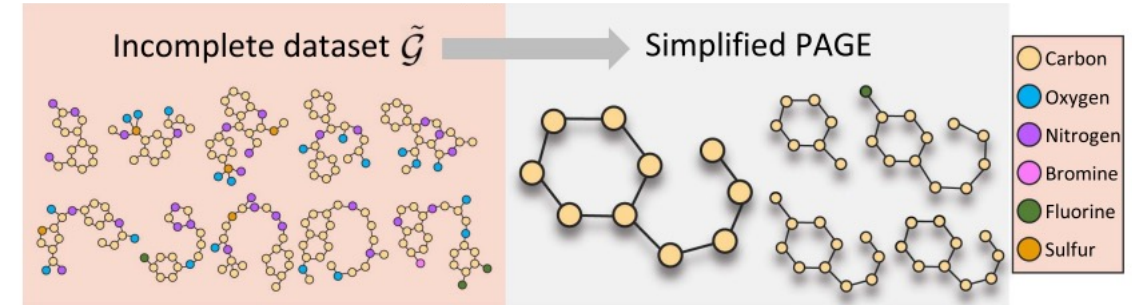


(b) Benzene.

How does PAGE perform when only a subset of dataset is available?



(a) BA-house.



(b) Benzene.

Empirical results clearly indicate PAGE can reliably run when the dataset is incomplete.

- We generate an incomplete dataset with only 10 graphs.
- We then run a simplified version of PAGE, where it skips Phase 1 (selection of k graphs via clustering) and replaces by random selection.
- We then run PAGE multiple times and observe the output.

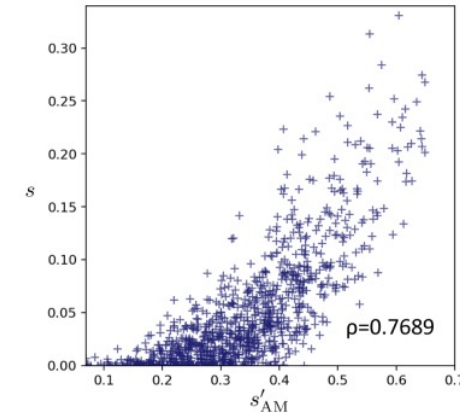
On the efficiency and quality of the Prototype Scoring Function

We compare our Prototype Scoring Function with naïve baselines, including a pairwise average of the arithmetic (AM) & geometric mean (GM).

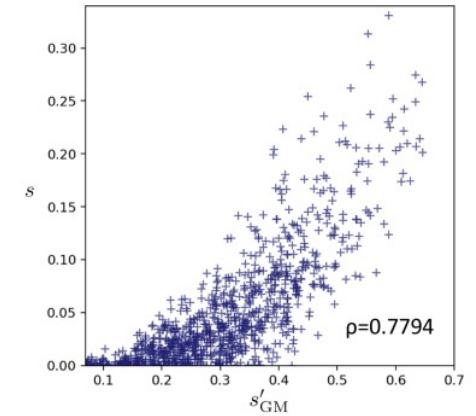
TABLE IV
COMPARISON OF DIFFERENT PROTOTYPE SCORING FUNCTIONS IN TERMS OF
THE RUNTIME COMPLEXITY IN MICROSECONDS (MEAN $\pm$ STANDARD
DEVIATION)

Prototype scoring function	s	s'_{AM}	s'_{GM}
Time (μs)	14.42 ± 12.65	38.34 ± 17.81	31.39 ± 17.81

The runtime comparison shows that our proposed Prototype Scoring Function is faster than alternatives.



(a) s'_{AM} versus s



(b) s'_{GM} versus s

We also show that there are a high correlation between our Prototype Scoring Function and the alternatives, providing an intuitive understanding of what it calculates.

- In our work, we propose PAGE, a novel model-level explanation of a GNN model that performs graph classification.
- In contrast to XGNN, which relies on reinforcement learning that requires carefully designed reward functions along with domain knowledge, our method discovers explanations within the dataset.
- Our future avenues include expanding our method to GNNs with node-level or edge-level tasks.

Thank you for your attention!

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