



FAITH: Few-Shot Graph Classification with Hierarchical Task Graphs

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Code:<https://github.com/SongW-SW/FAITH>

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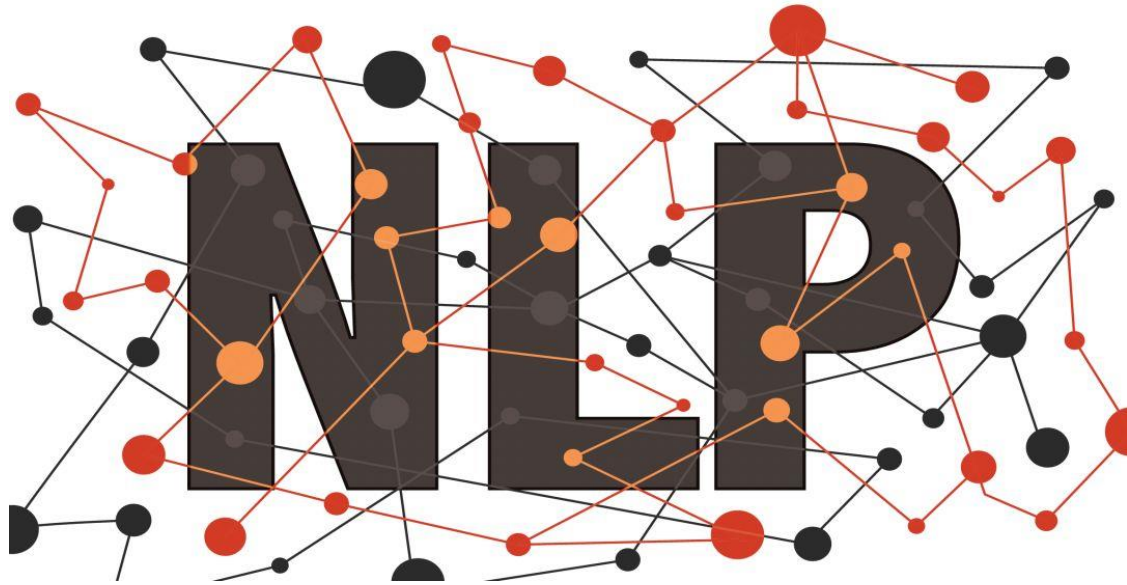
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NATURAL LANGUAGE PROCESSING



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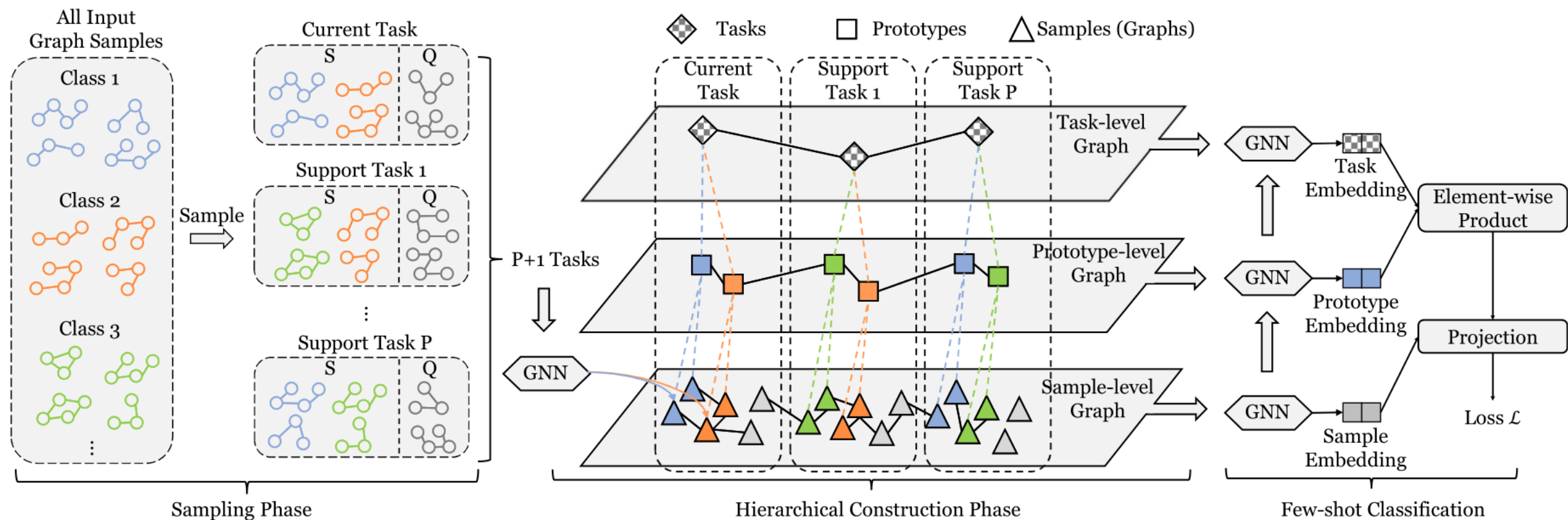


Introduction

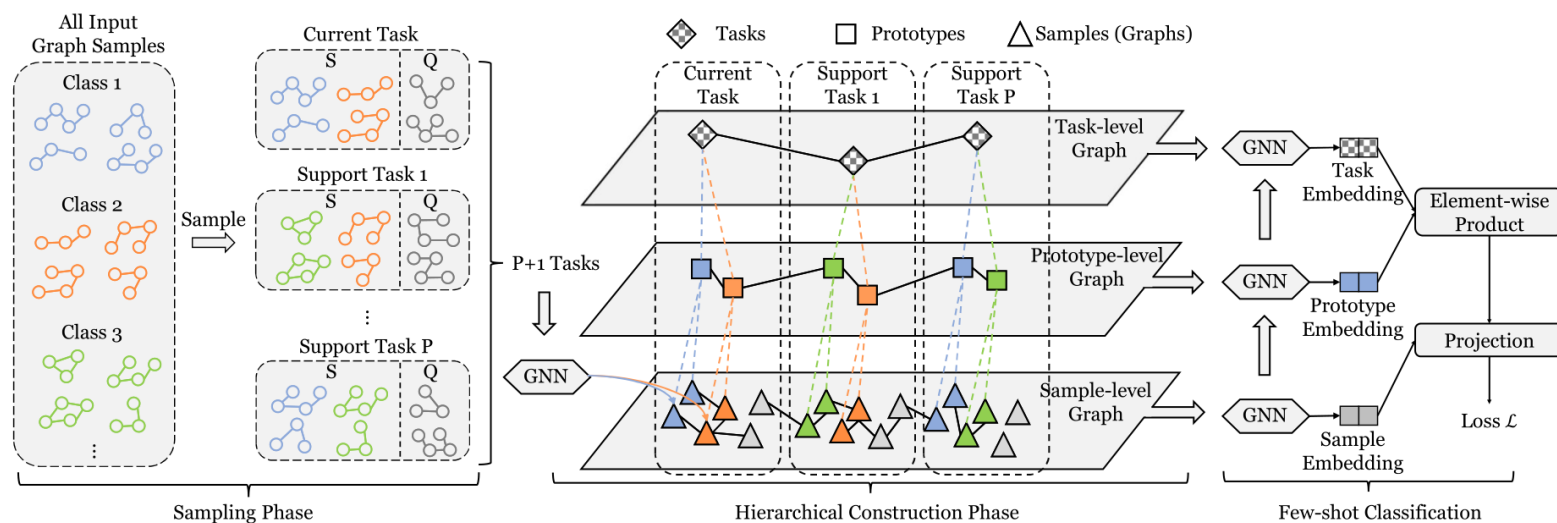
existing few-shot graph classification methods treat different meta-training tasks independently without considering task correlations.

For example, the target task of the **toxicity property prediction** with the meta-training task of the **chemical activity prediction** has stronger task correlations.

Method



Method



Loss-based Sampling for Support Tasks

$$\mathbf{p}_i = \text{softmax}(\text{MLP}(\frac{1}{K} \sum_{j=1}^K \mathbf{z}_i^j)), \quad (1)$$

$$\mathcal{L}_{\text{sample}} = -\frac{1}{N} \sum_{i=1}^N \sum_{j=1}^C y_{i,j} \log p_{i,j}, \quad (2)$$

Constructing Hierarchical Task Graphs

$$\mathbf{A}_s = \mathbf{A}'_s + \mathbf{A}''_s,$$

$$\mathbf{A}'_s(i, j) = \cos(\mathbf{Z}_s(i), \mathbf{Z}_s(j))$$

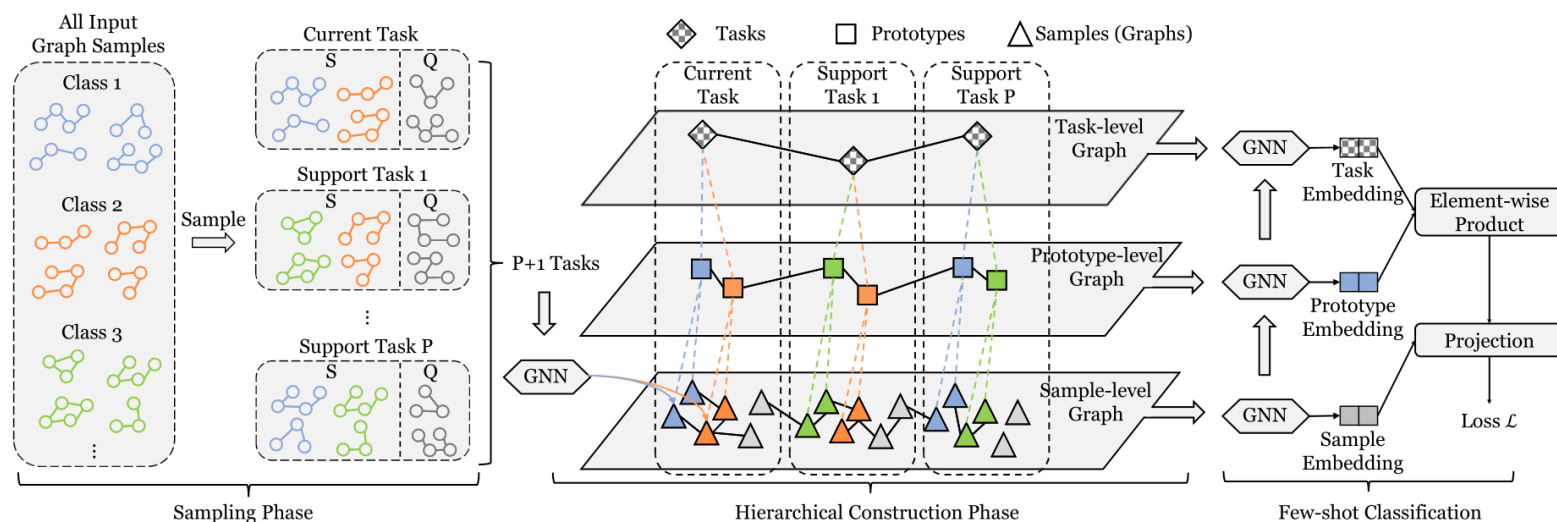
$$\mathbf{A}''_s(i, j) = \begin{cases} 1, & \text{if } y_i = y_j \\ 0, & \text{otherwise} \end{cases}, \quad (3)$$

$$\mathbf{H}_s = \text{GNN}_h^{(s)}(\mathbf{Z}_s, \mathbf{A}_s), \quad (4)$$

$$\mathbf{G}_s = \text{GNN}_g^{(s)}(\mathbf{Z}_s, \mathbf{A}_s), \quad (5)$$

$$\mathbf{Z}_p(i) = \text{softmax}(\mathbf{G}_s^i)^\top \mathbf{H}_s^i, \quad (6)$$

Method



Task-specific Few-shot Classification

$$z_{i,j}^k = (\mathbf{s}_i^k)^\top \mathbf{W}(\mathbf{p}_j^k \circ \mathbf{t}^k), \quad (7)$$

$$\bar{z}_{i,j}^k = \exp(z_{i,j}^k) / (\sum_{j=1}^N \exp(z_{i,j}^k))$$

$$\mathcal{L}_{class} = -\frac{1}{(P+1)Q} \sum_{k=1}^{(P+1)} \sum_{i=1}^Q \sum_{j=1}^N y_{i,j}^k \log \bar{z}_{i,j}^k, \quad (8)$$

$$\mathcal{L} = \mathcal{L}_{class} + \alpha \mathcal{L}_{sample}, \quad (9)$$



Experiment

Table 1: Results of all methods with different few-shot settings on four benchmark datasets. The best results are shown in bold.

Methods	Letter-high		ENZYMES		TRIANGLES		Reddit-12K	
	5-shot	10-shot	5-shot	10-shot	5-shot	10-shot	5-shot	10-shot
WL	65.27 ± 7.67	68.39 ± 4.69	55.78 ± 4.72	58.37 ± 3.84	51.25 ± 4.02	53.26 ± 2.95	40.26 ± 5.17	42.57 ± 3.69
Graphlet	33.76 ± 6.94	37.59 ± 4.60	53.17 ± 5.92	55.30 ± 3.78	40.17 ± 3.18	43.76 ± 3.09	33.76 ± 6.94	37.59 ± 4.60
PN	68.48 ± 3.28	72.60 ± 3.01	53.72 ± 4.37	55.79 ± 3.95	69.56 ± 3.97	73.12 ± 3.64	42.31 ± 2.32	43.23 ± 2.01
Relation	51.14 ± 4.21	52.54 ± 4.04	41.39 ± 4.73	43.27 ± 3.49	46.09 ± 3.10	49.15 ± 3.49	34.89 ± 3.76	37.76 ± 3.09
GSM	69.91 ± 5.90	73.28 ± 3.64	55.42 ± 5.74	60.64 ± 3.84	71.40 ± 4.34	75.60 ± 3.67	41.59 ± 4.12	45.67 ± 3.68
AS-MAML	69.44 ± 0.75	75.93 ± 0.53	49.83 ± 1.12	52.30 ± 1.43	78.42 ± 0.67	80.39 ± 0.56	36.96 ± 0.74	41.47 ± 0.83
FAITH	71.55 ± 3.58	76.65 ± 3.26	57.89 ± 4.65	62.16 ± 4.11	79.59 ± 4.05	80.79 ± 3.53	42.71 ± 4.18	46.63 ± 4.01

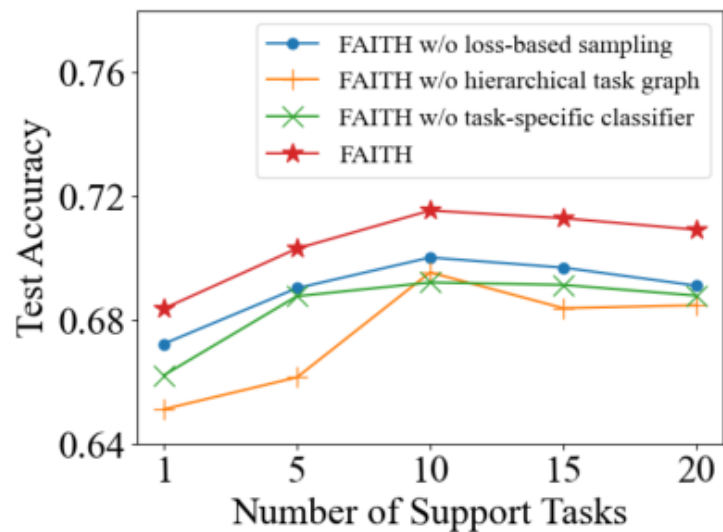


Experiment

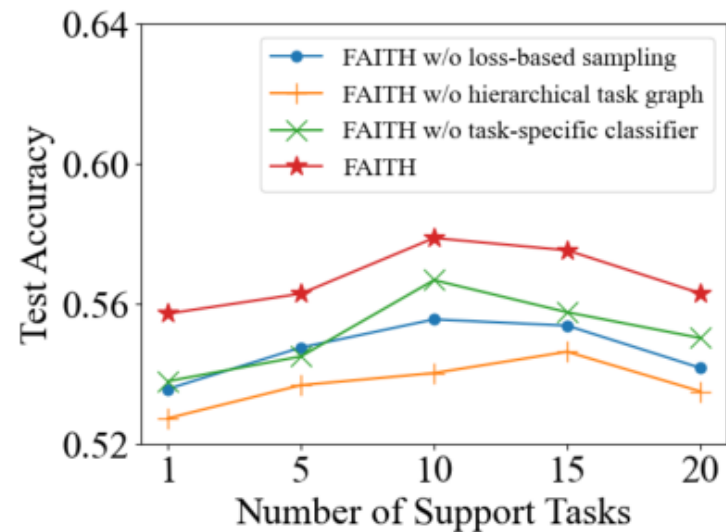
Table 2: Detailed statistics of used datasets.

Dataset	$ \mathcal{Y}_f / \mathcal{Y}_t $	# Graphs	# Nodes	# Edges
Letter-high	4/11	2,250	4.67	4.50
ENZYMES	2/4	600	32.63	62.14
TRIANGLES	3/7	2,000	20.85	35.50
Reddit-12K	4/7	1,111	391.41	456.89

Experiment



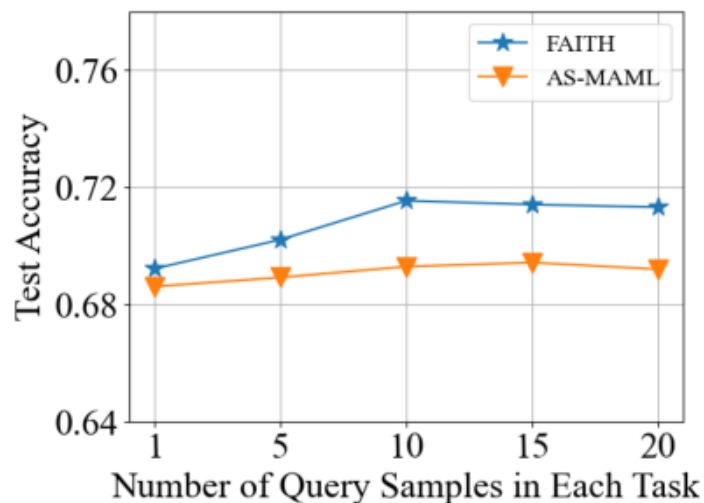
(a) Results on Letter-high



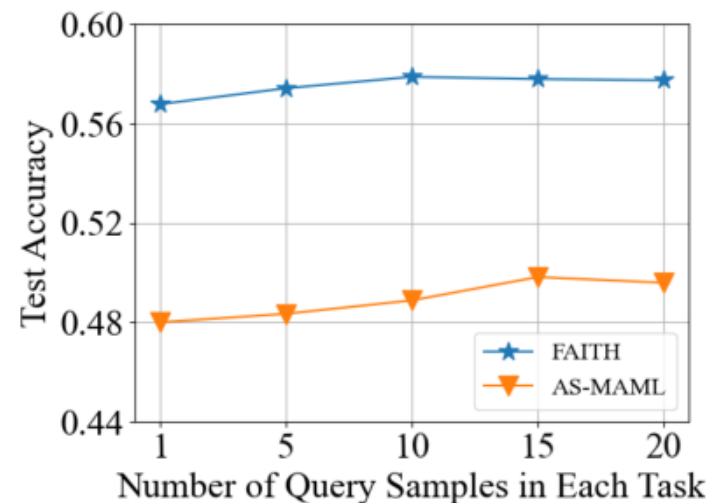
(b) Results on ENZYMES

Figure 2: Ablation study on Letter-high and ENZYMES.

Experiment



(a) Results on Letter-high



(b) Results on ENZYMES

Figure 3: Accuracy with respect to the number of query samples of FAITH and AS-MAML on two datasets.



Thank you!



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