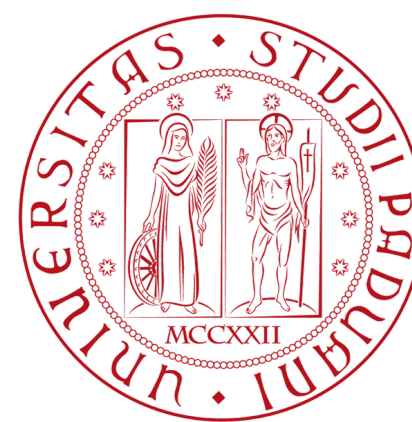


A Meta-Learning Approach for Graph Representation Learning in Multi-Task Settings

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University of Padova

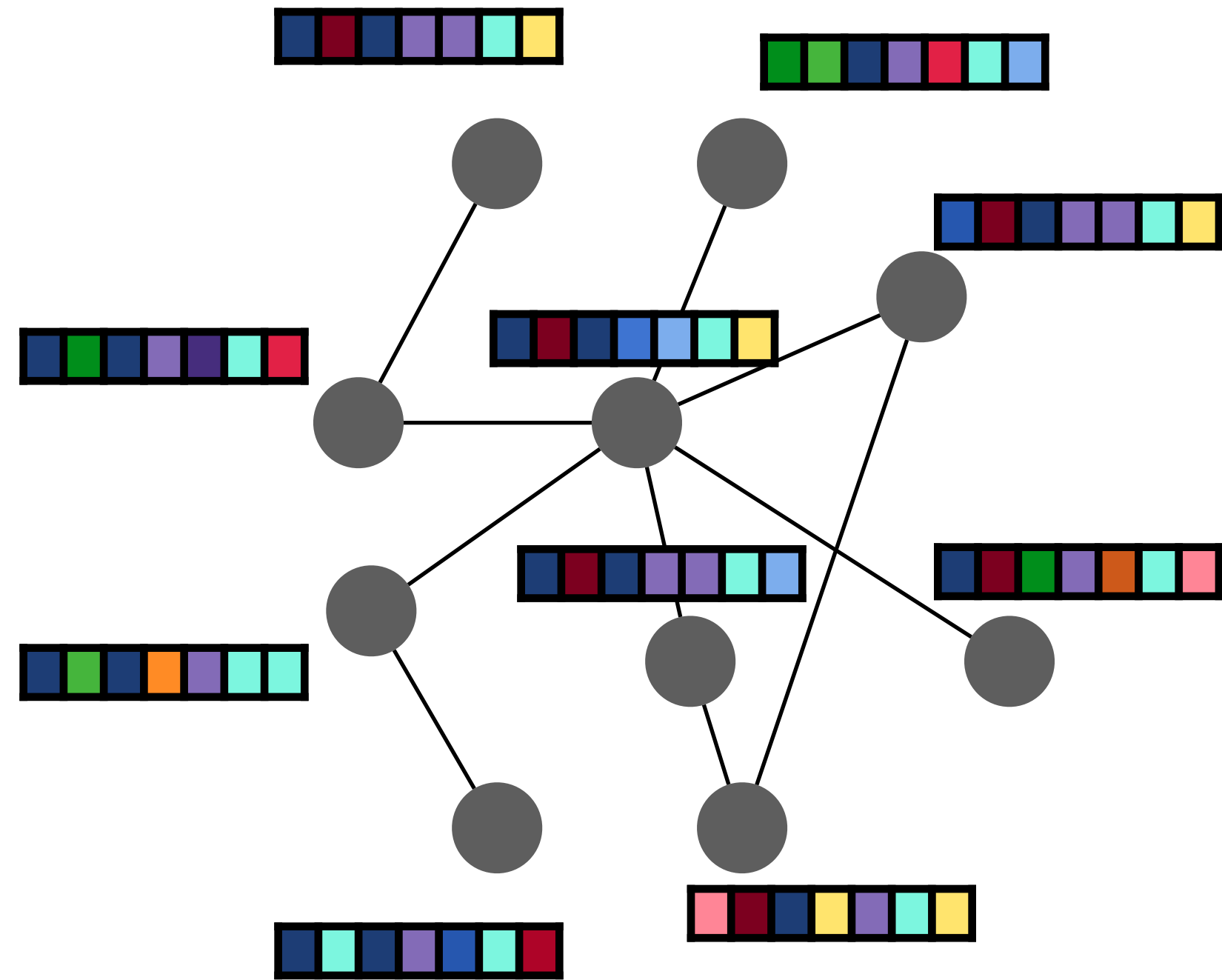
{davide.buffelli, fabio.vandin}@unipd.it



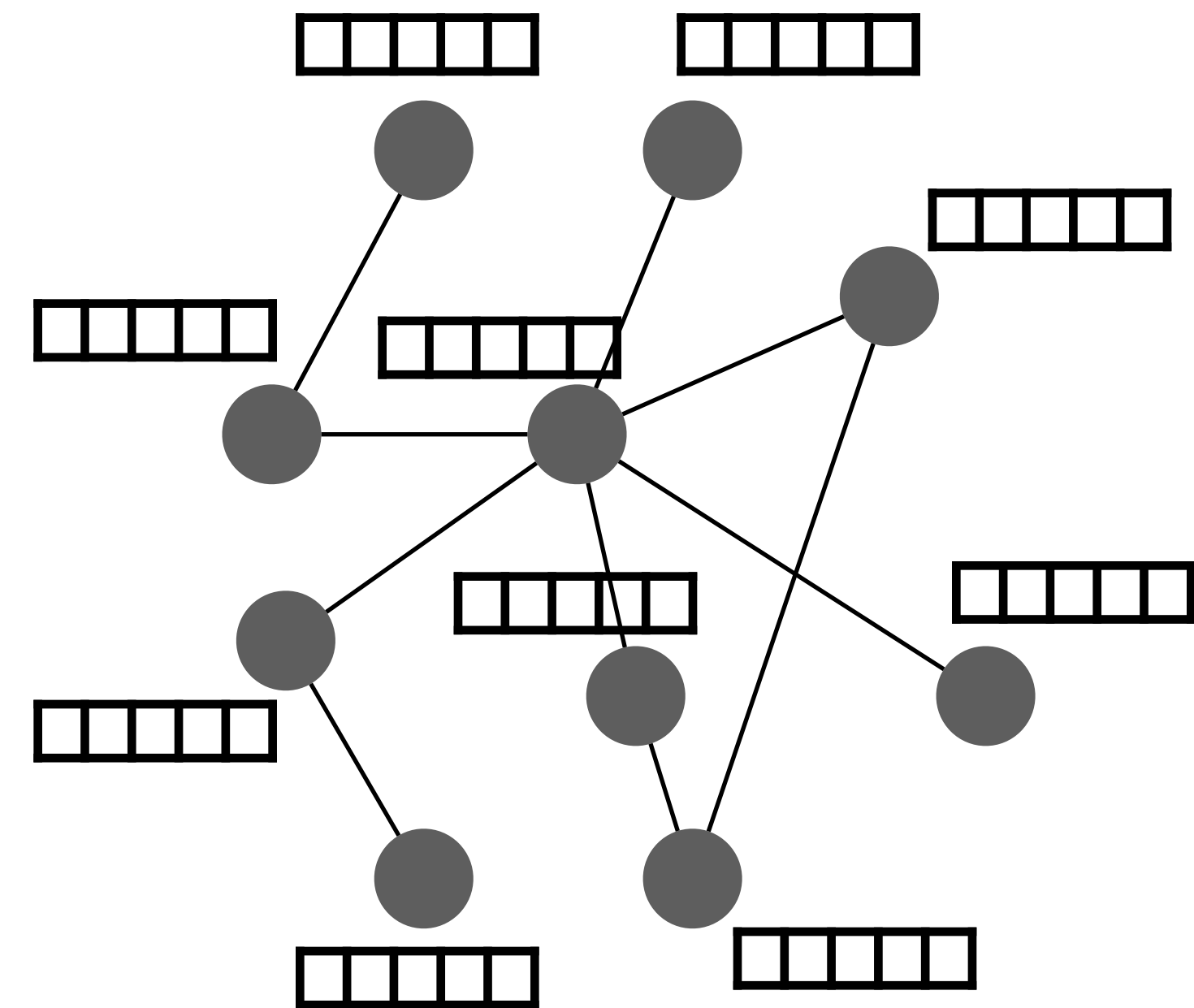
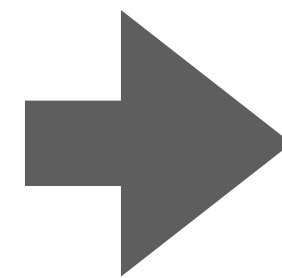
Graph Representation Learning

Node
Features  $\in \mathbb{R}^D$

Node
Embedding  $\in \mathbb{R}^d, \quad d \ll D$

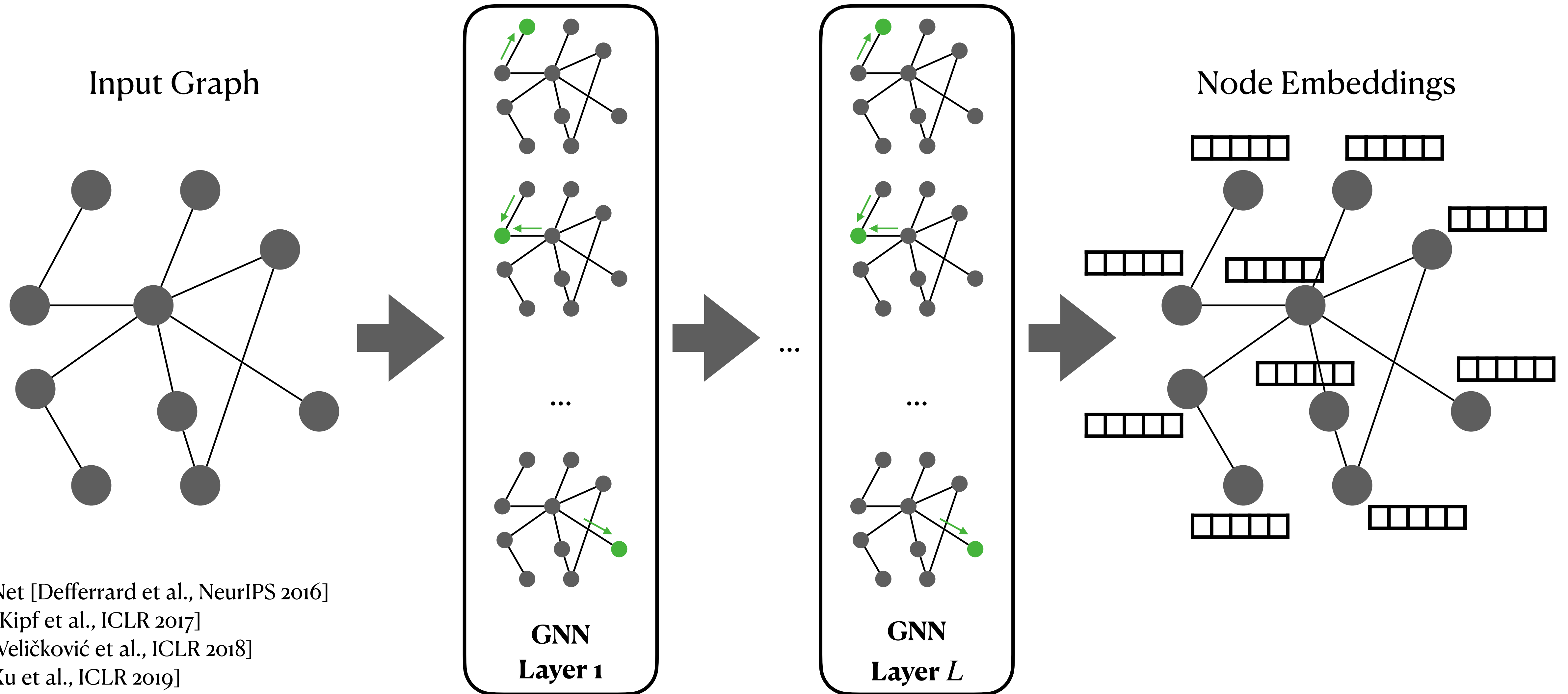


Input Graph



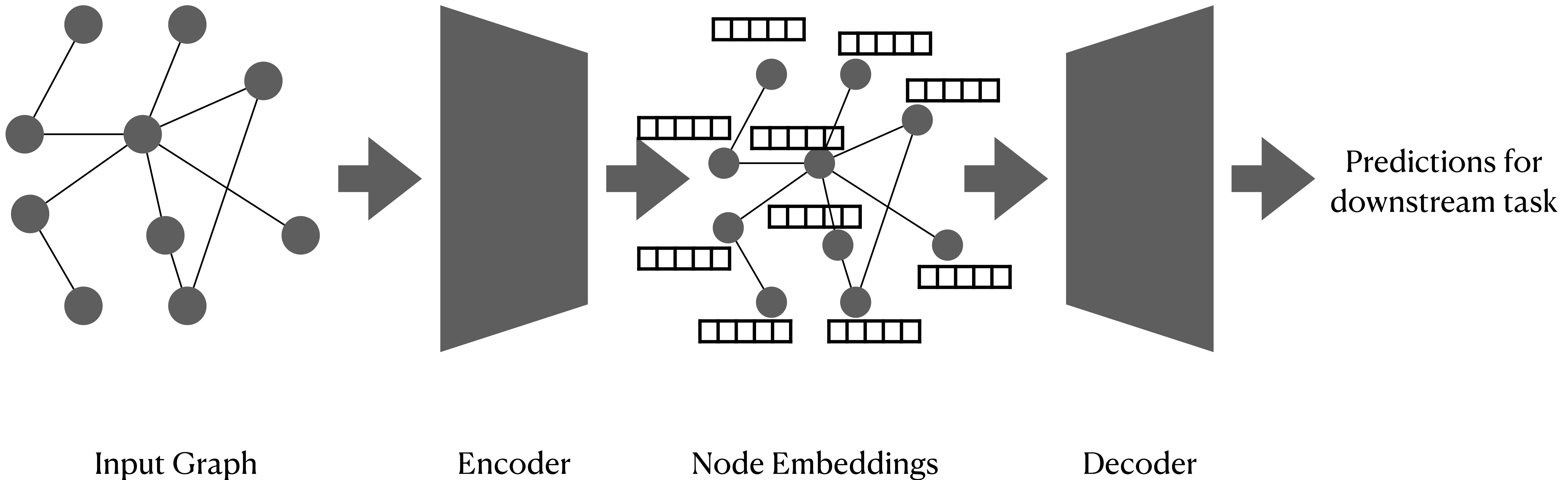
Node Embeddings

Graph Neural Networks



Graph Representation Learning

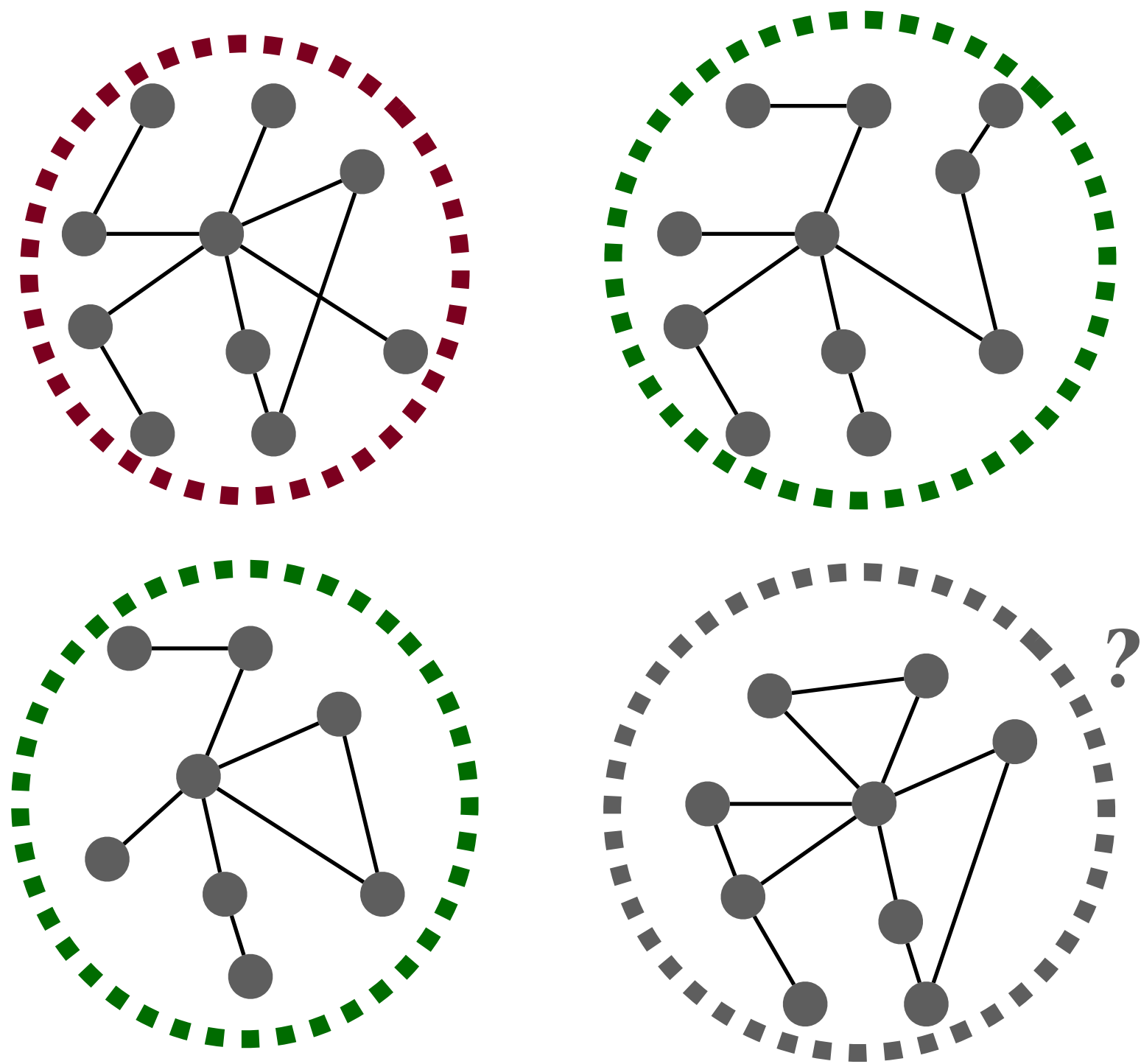
Typical Framework



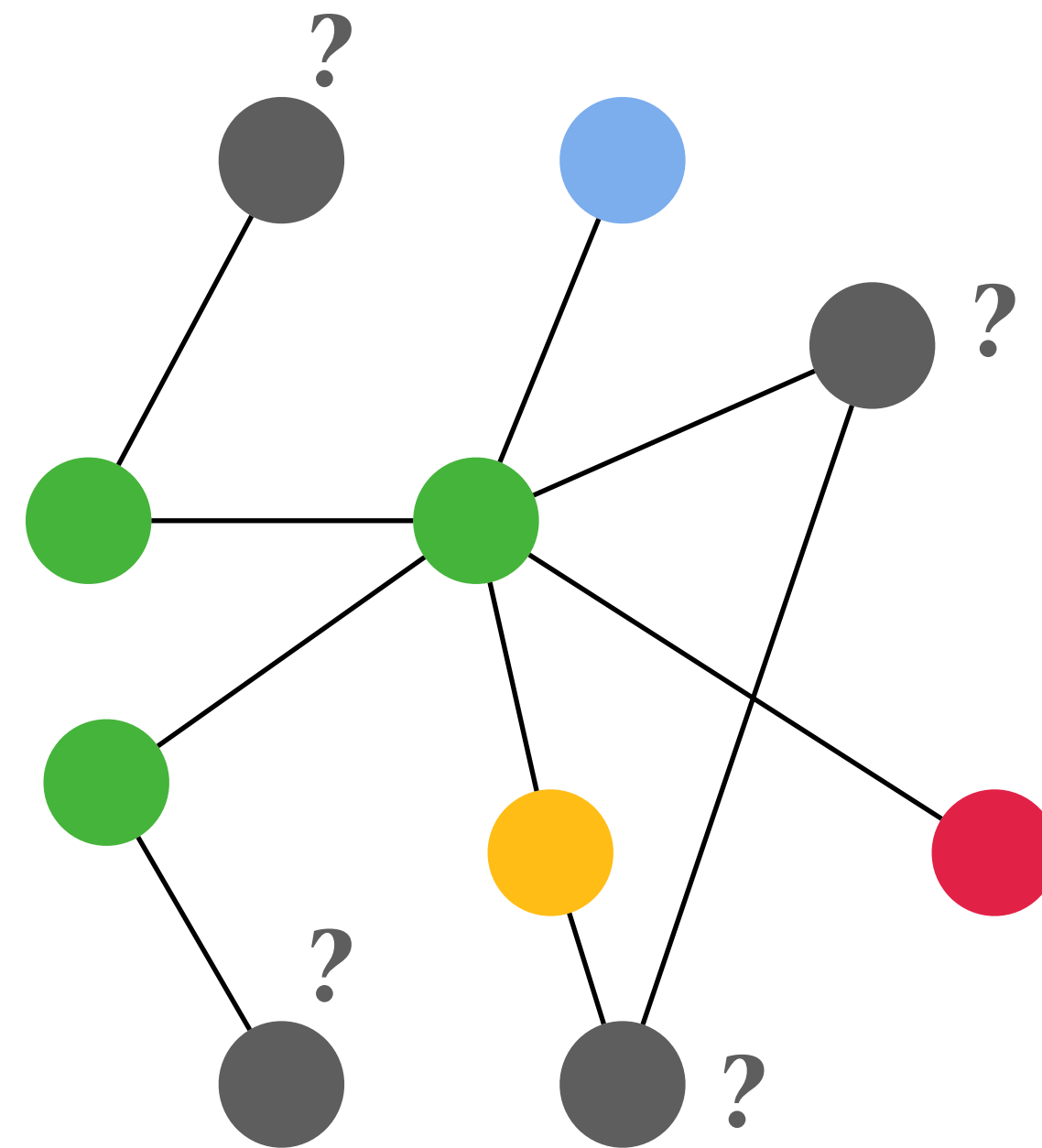
Graph Representation Learning

Applications

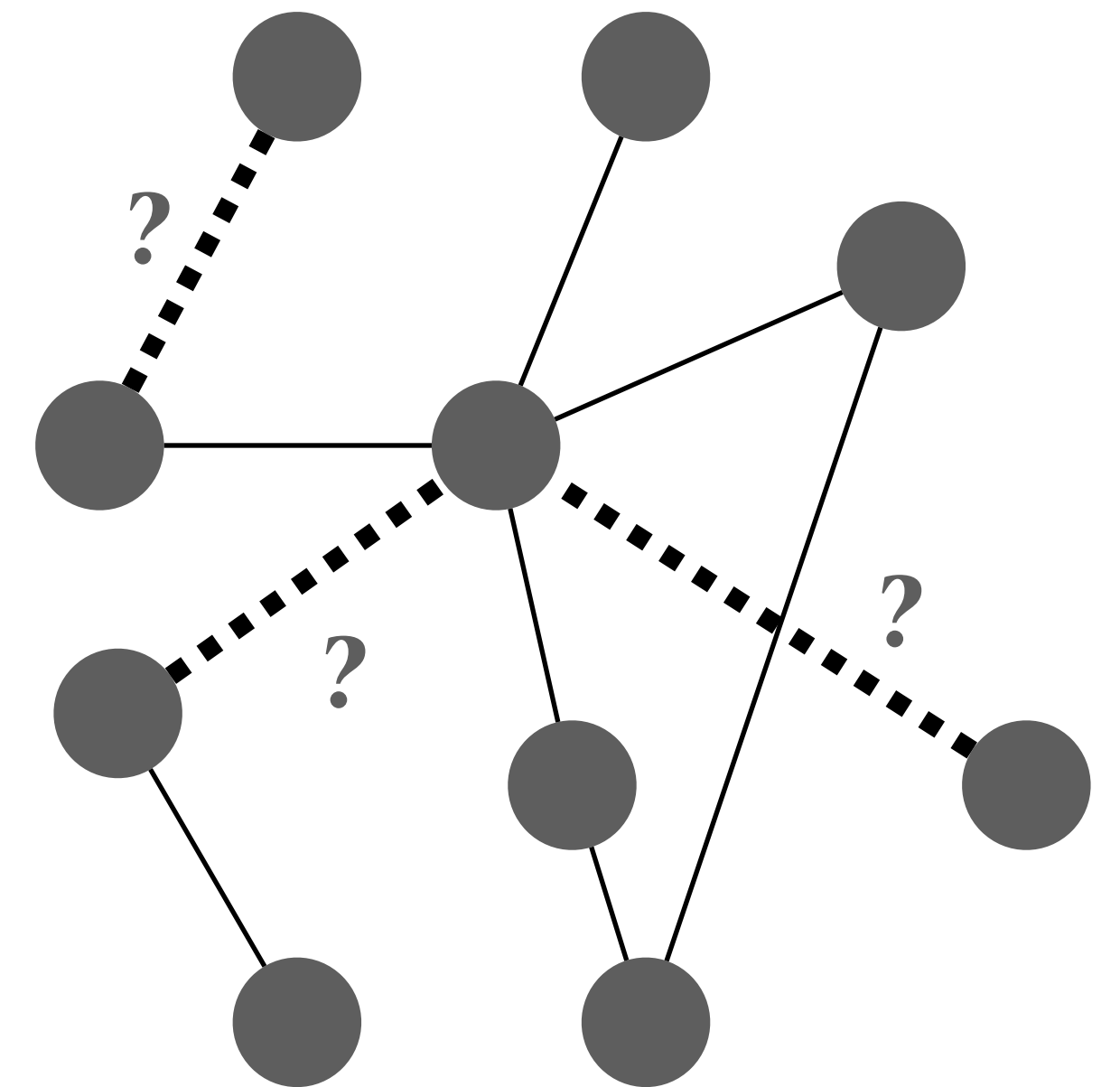
Graph Classification



Node Classification



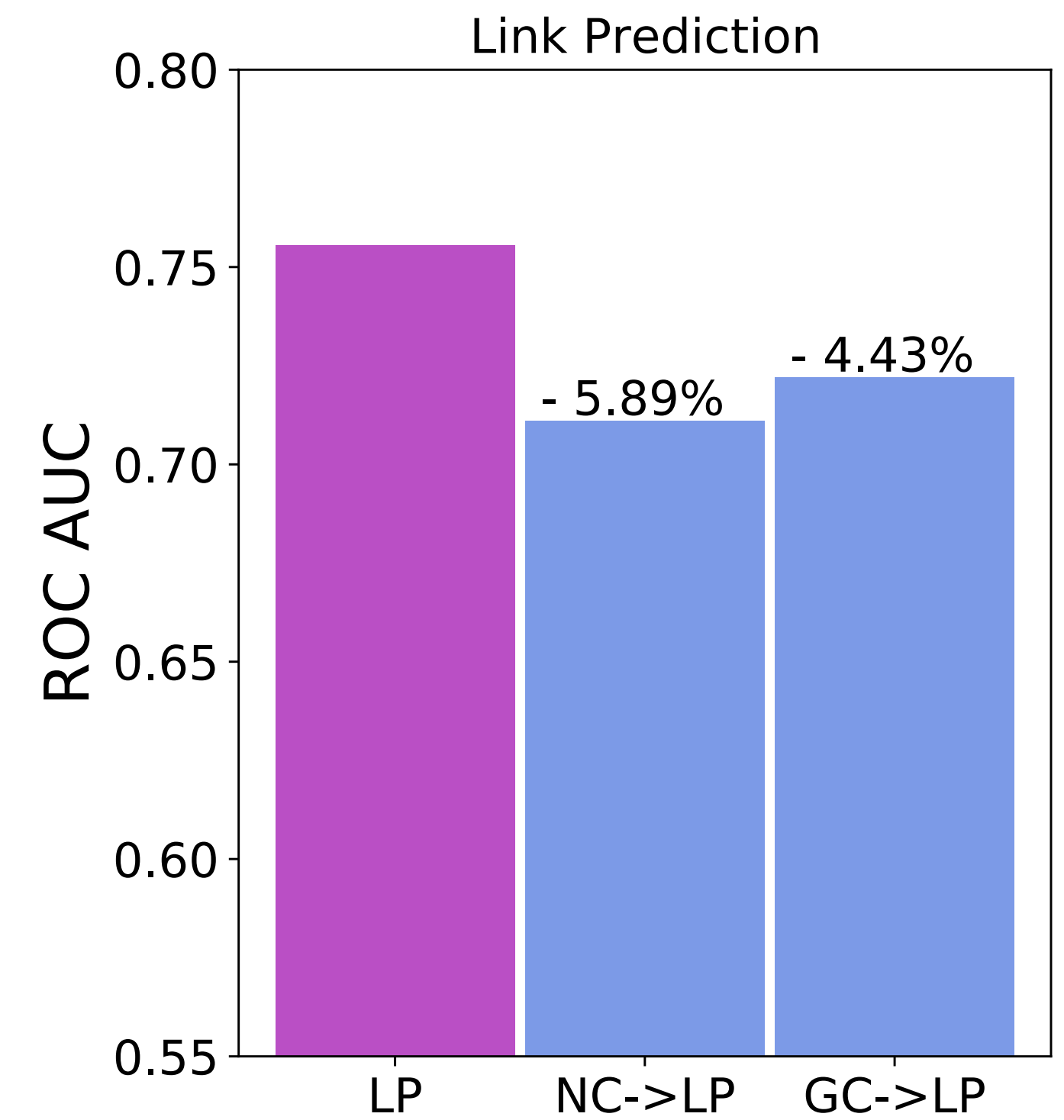
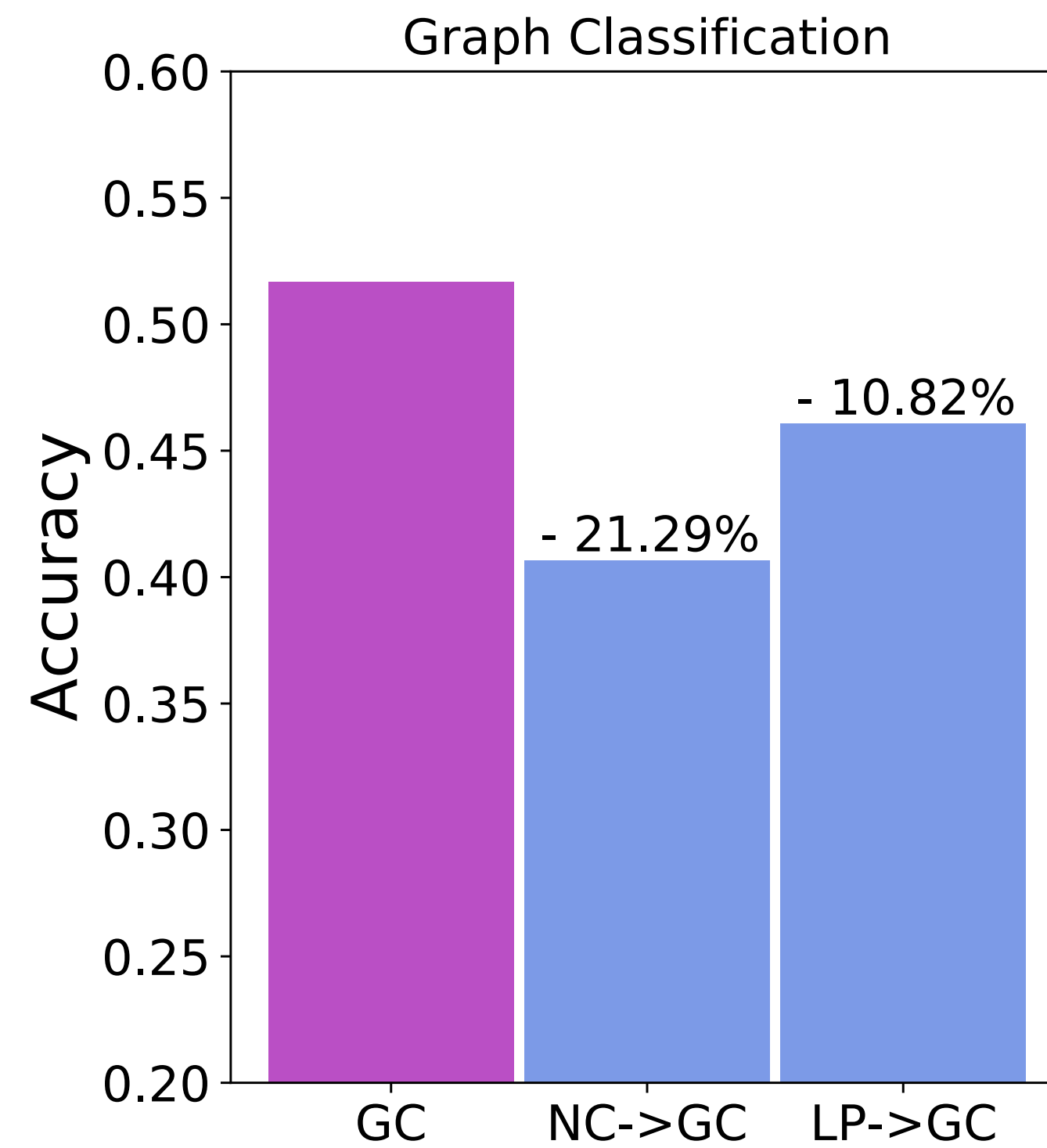
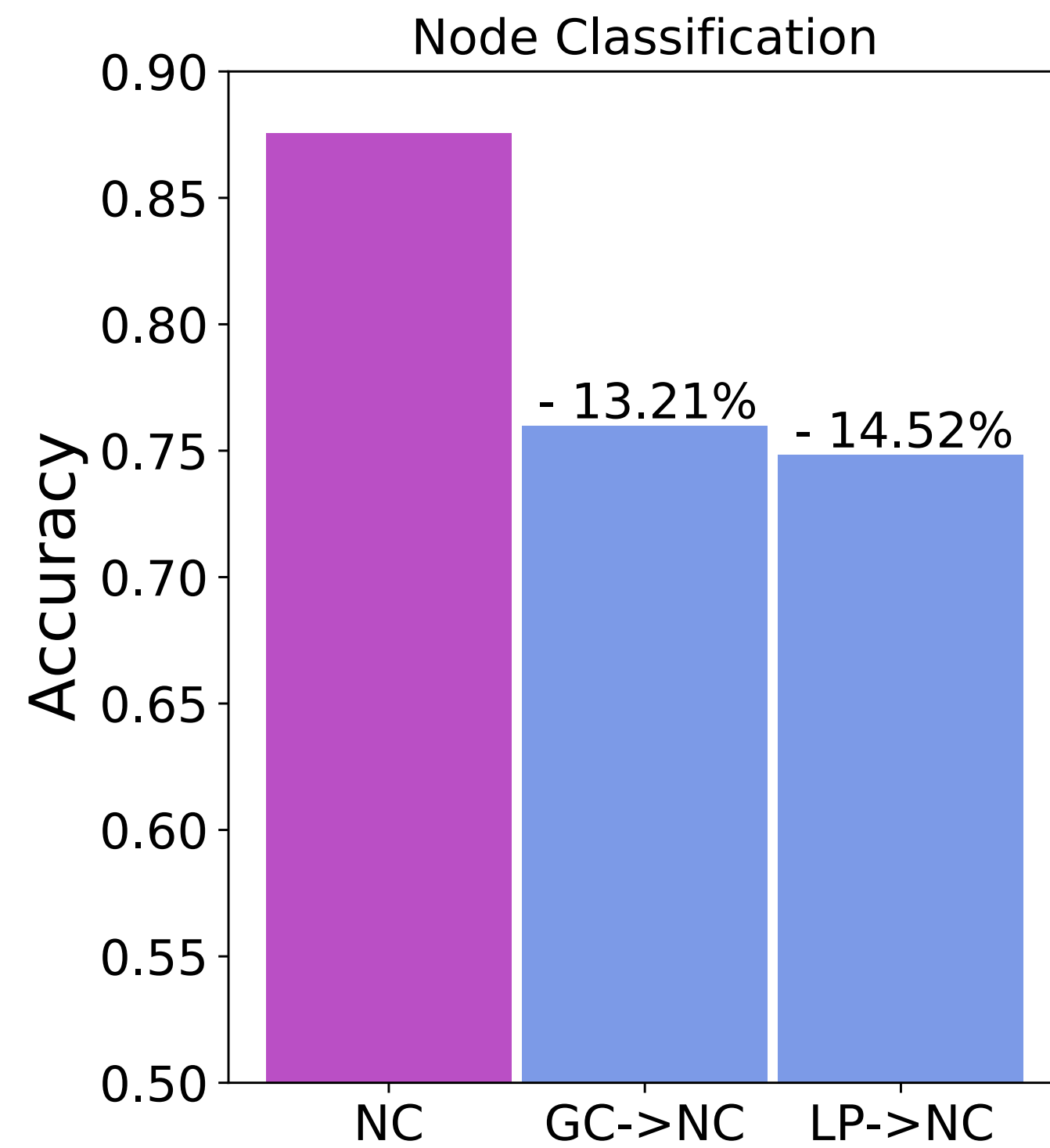
Link Prediction



Graph Representation Learning

Transferability of Embeddings

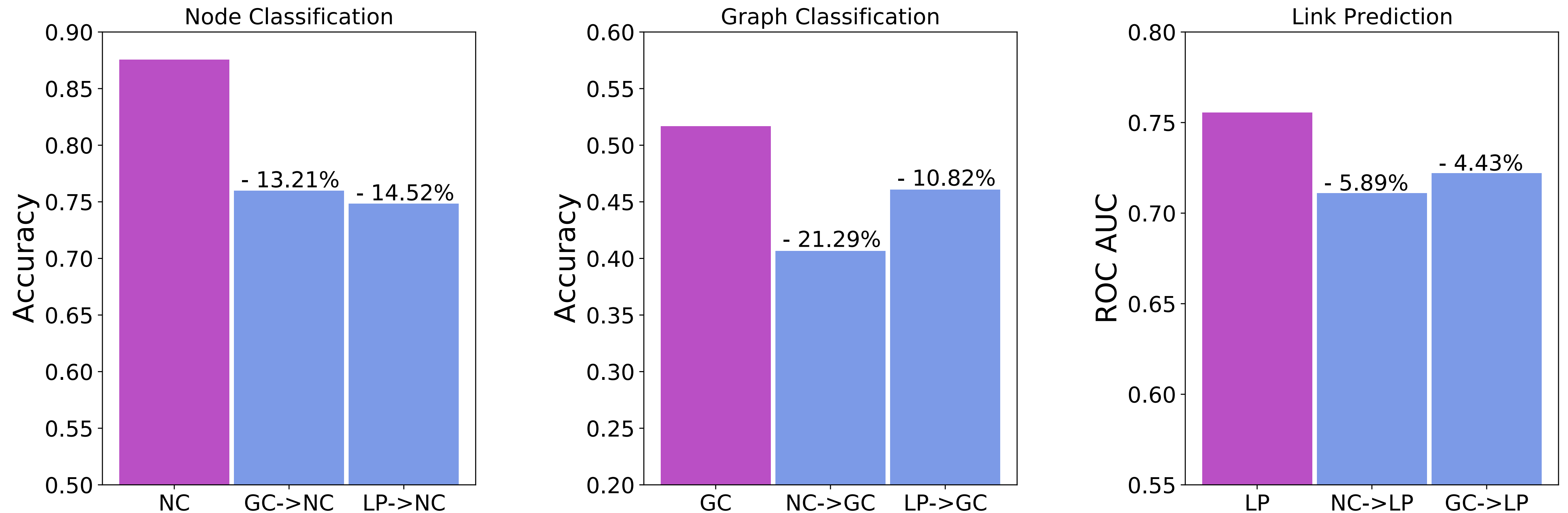
Original Embeddings **Transferred Embeddings**



Graph Representation Learning

Transferability of Embeddings

Original Embeddings Transferred Embeddings

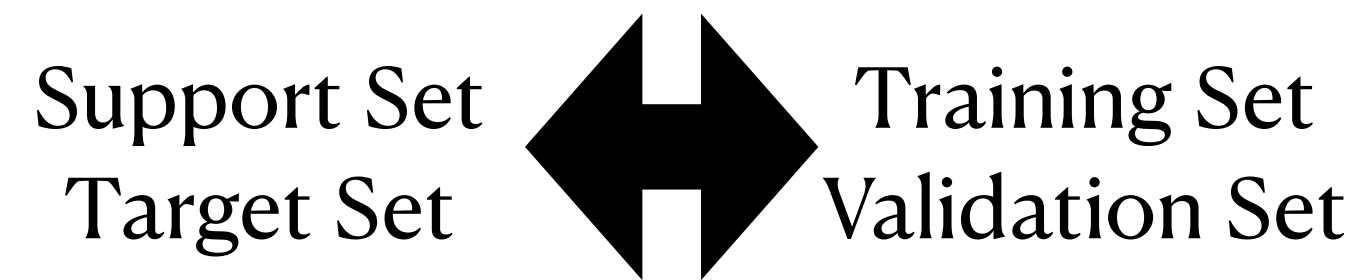


Can we generate node embeddings that generalize across tasks?

SAME: Single-Task Adaptation for Multi-Task Embeddings

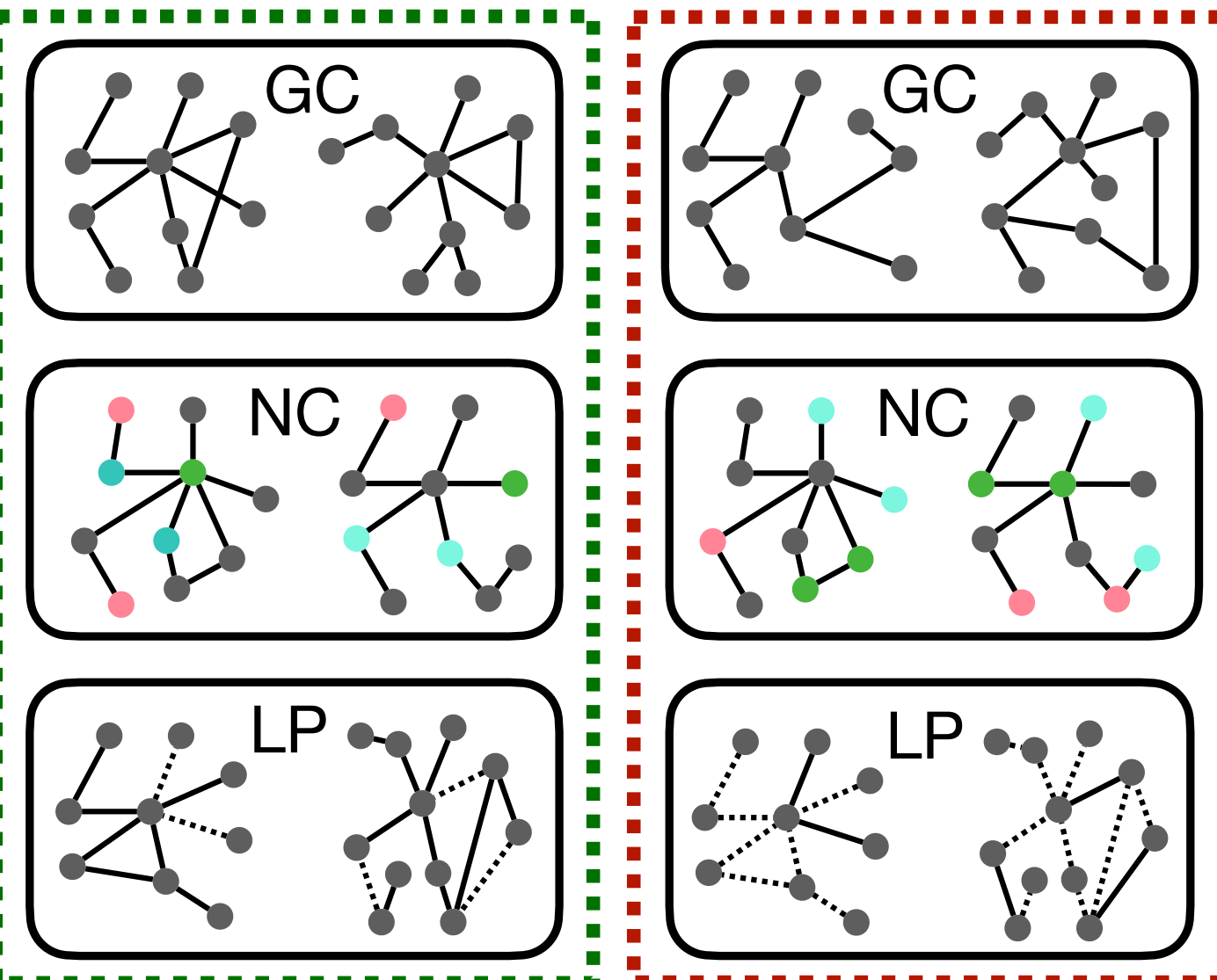
Multi-Task Episodes

- 1 task-specific support set per task
- 1 task-specific target set per task



Support Set

Target Set



Multi-Learning Procedure

- Separate adaptation for each task
- Unique outer loop update

Two variants:

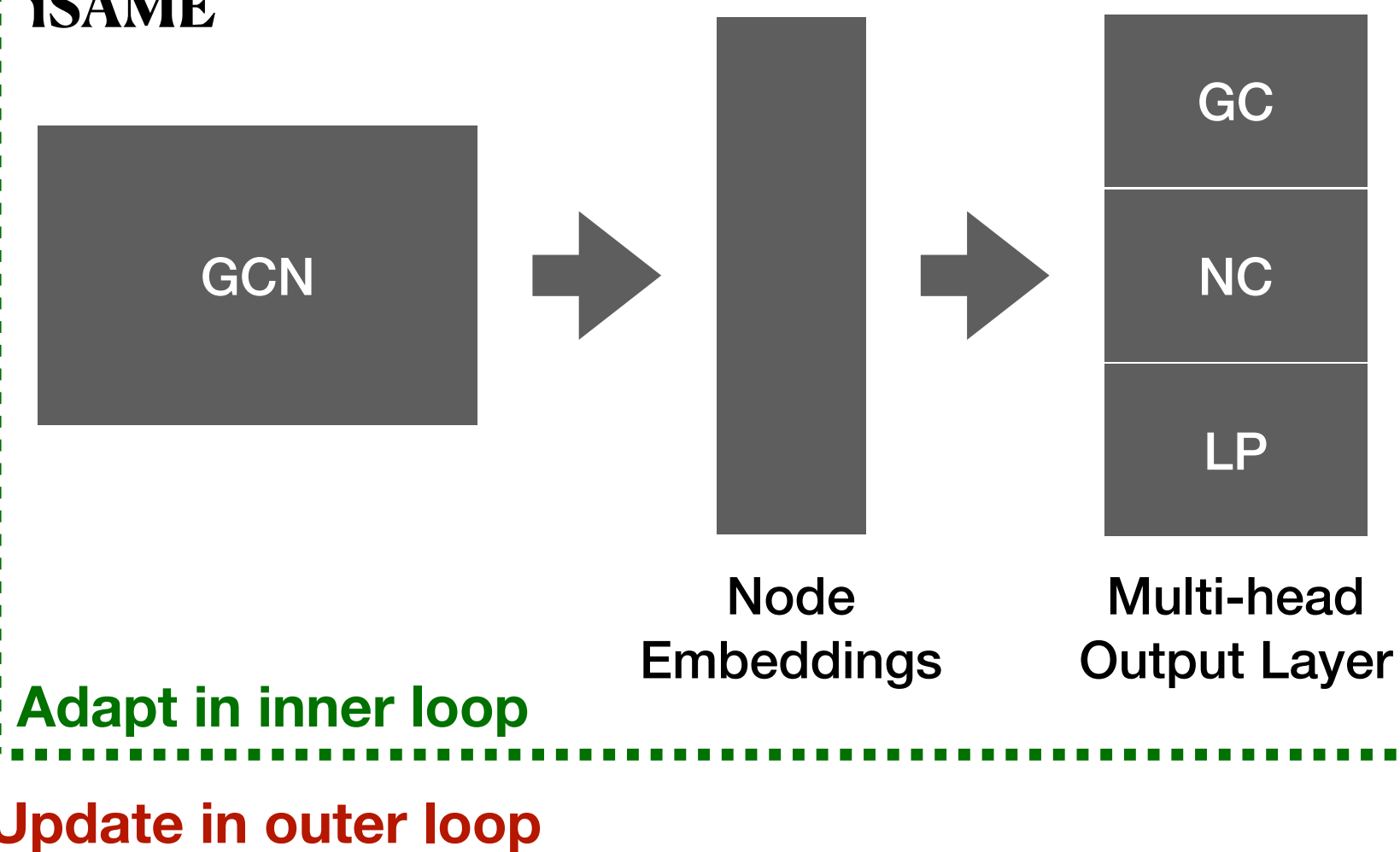
- eSAME
- iSAME

Algorithm 1: Proposed Meta-Learning Procedure

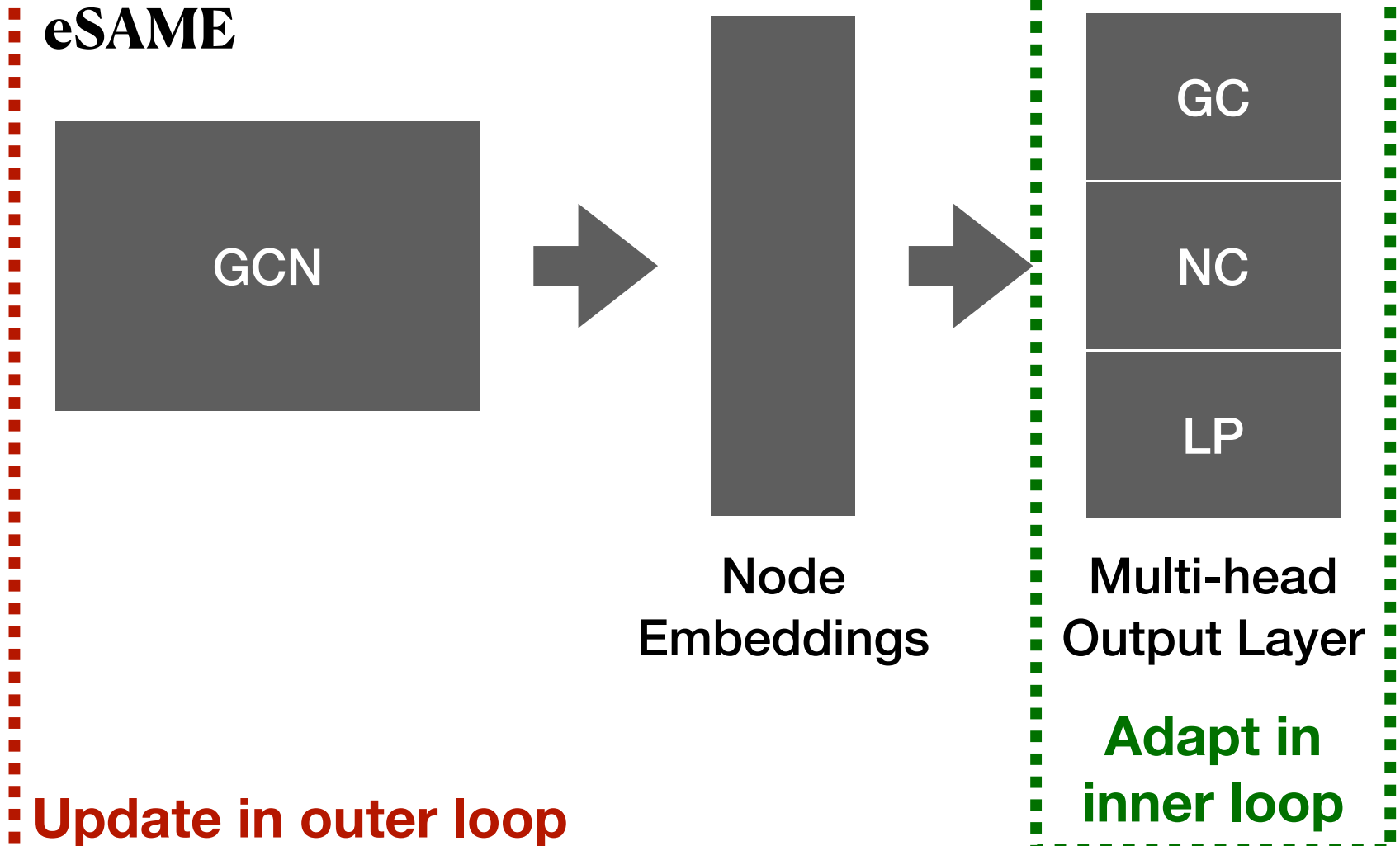
```

Input : Model  $f_\theta$ ; Episodes  $\mathcal{E} = \{\mathcal{E}_1, \dots, \mathcal{E}_n\}$ 
init( $\theta$ )
for  $\mathcal{E}_i$  in  $\mathcal{E}$  do
    o_loss  $\leftarrow$  0
    for  $t$  in (GC, NC, LP) do
         $\theta'^{(t)} \leftarrow \theta$ 
         $\theta'^{(t)} \leftarrow \text{ADAPT}(f_\theta, \mathcal{S}_{\mathcal{E}_i}^{(t)}, \mathcal{L}_{\mathcal{E}_i}^{(t)})$ 
        o_loss  $\leftarrow$  o_loss + TEST( $f_{\theta'^{(t)}}$ ,  $\mathcal{T}_{\mathcal{E}_i}^{(t)}, \mathcal{L}_{\mathcal{E}_i}^{(t)}$ )
    end
     $\theta \leftarrow \text{UPDATE}(\theta, \text{o\_loss}, \theta'^{(GC)}, \theta'^{(NC)}, \theta'^{(LP)})$ 
end
    
```

iSAME



eSAME



Experiments

- 1. Do iSAME and eSAME lead to high quality node embeddings in the single-task setting?**
- 2. Do iSAME and eSAME lead to high quality node embeddings in the multi-task setting?**
- 3. Do iSAME and eSAME extract information that is not captured by classically trained multi-task models?**
- 4. Can the node embeddings learned by iSAME and eSAME be used to perform multiple tasks with comparable or better performance than classical multi-task models?**

Experiments

Do iSAME and eSAME lead to high quality node embeddings in the single-task setting?

Table 1: Results for a single-task model trained in a classical supervised manner (Cl), and a **linear** classifier trained on the embeddings produced by our meta-learning strategies (iSAME, eSAME).

| Task | Model | Dataset | | | |
|------|-------|----------------|----------------|----------------|----------------|
| | | ENZYMES | PROTEINS | DHFR | COX2 |
| NC | Cl | 87.5 ± 1.9 | 72.3 ± 4.4 | 97.3 ± 0.2 | 96.4 ± 0.3 |
| | iSAME | 87.3 ± 0.8 | 81.8 ± 1.6 | 96.6 ± 0.3 | 96.1 ± 0.4 |
| | eSAME | 87.8 ± 0.7 | 82.4 ± 1.6 | 96.8 ± 0.2 | 96.5 ± 0.6 |
| GC | Cl | 51.6 ± 4.2 | 73.3 ± 3.6 | 71.5 ± 2.3 | 76.7 ± 4.7 |
| | iSAME | 50.8 ± 2.9 | 73.5 ± 1.2 | 73.2 ± 3.2 | 76.3 ± 4.6 |
| | eSAME | 52.1 ± 5.0 | 72.6 ± 1.6 | 71.6 ± 2.4 | 75.6 ± 4.1 |
| LP | Cl | 75.5 ± 3.0 | 85.6 ± 0.8 | 98.8 ± 0.7 | 98.3 ± 0.8 |
| | iSAME | 81.7 ± 1.7 | 84.0 ± 1.1 | 99.2 ± 0.4 | 99.1 ± 0.5 |
| | eSAME | 80.1 ± 3.4 | 84.1 ± 0.9 | 99.2 ± 0.3 | 99.2 ± 0.7 |

Experiments

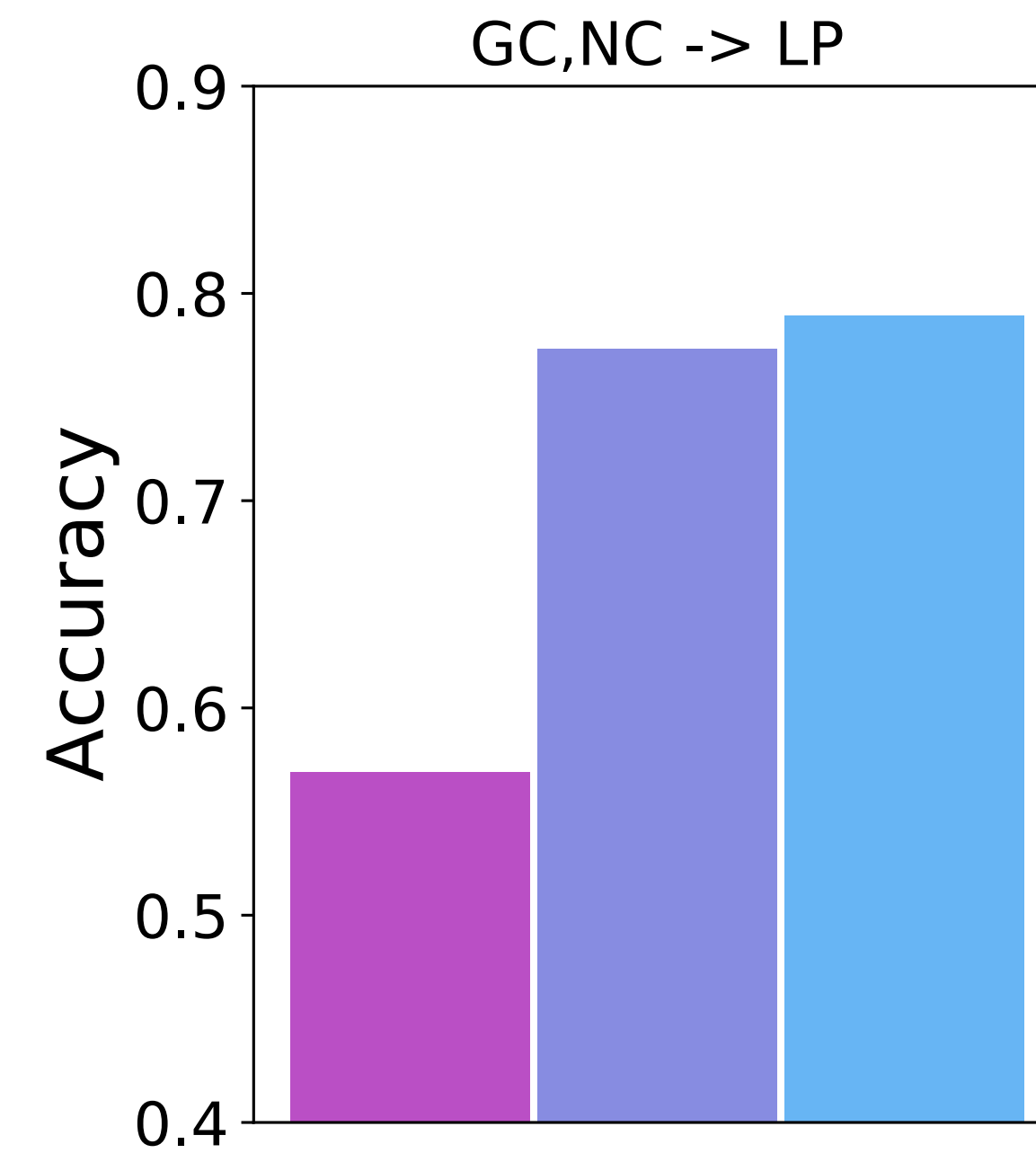
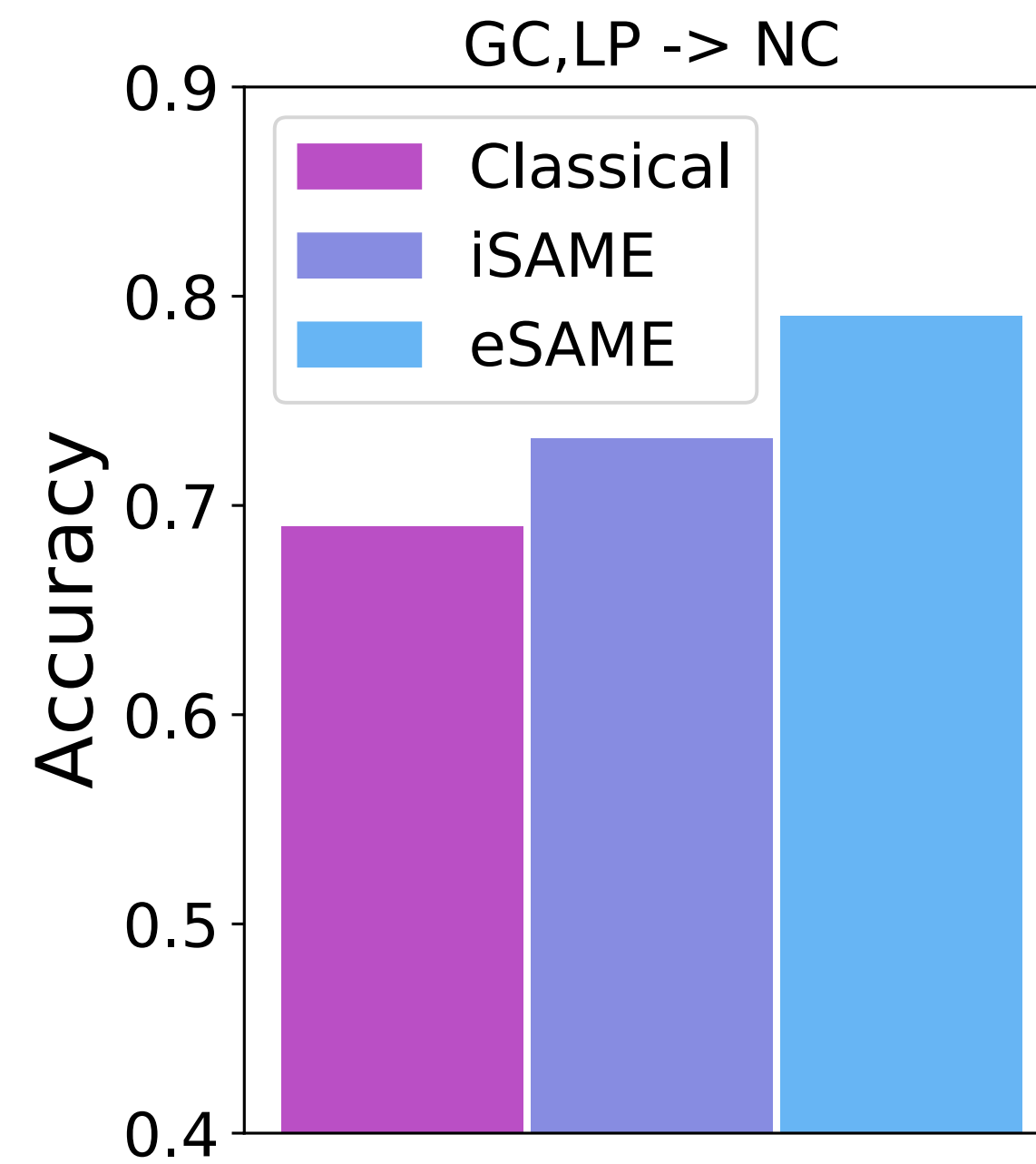
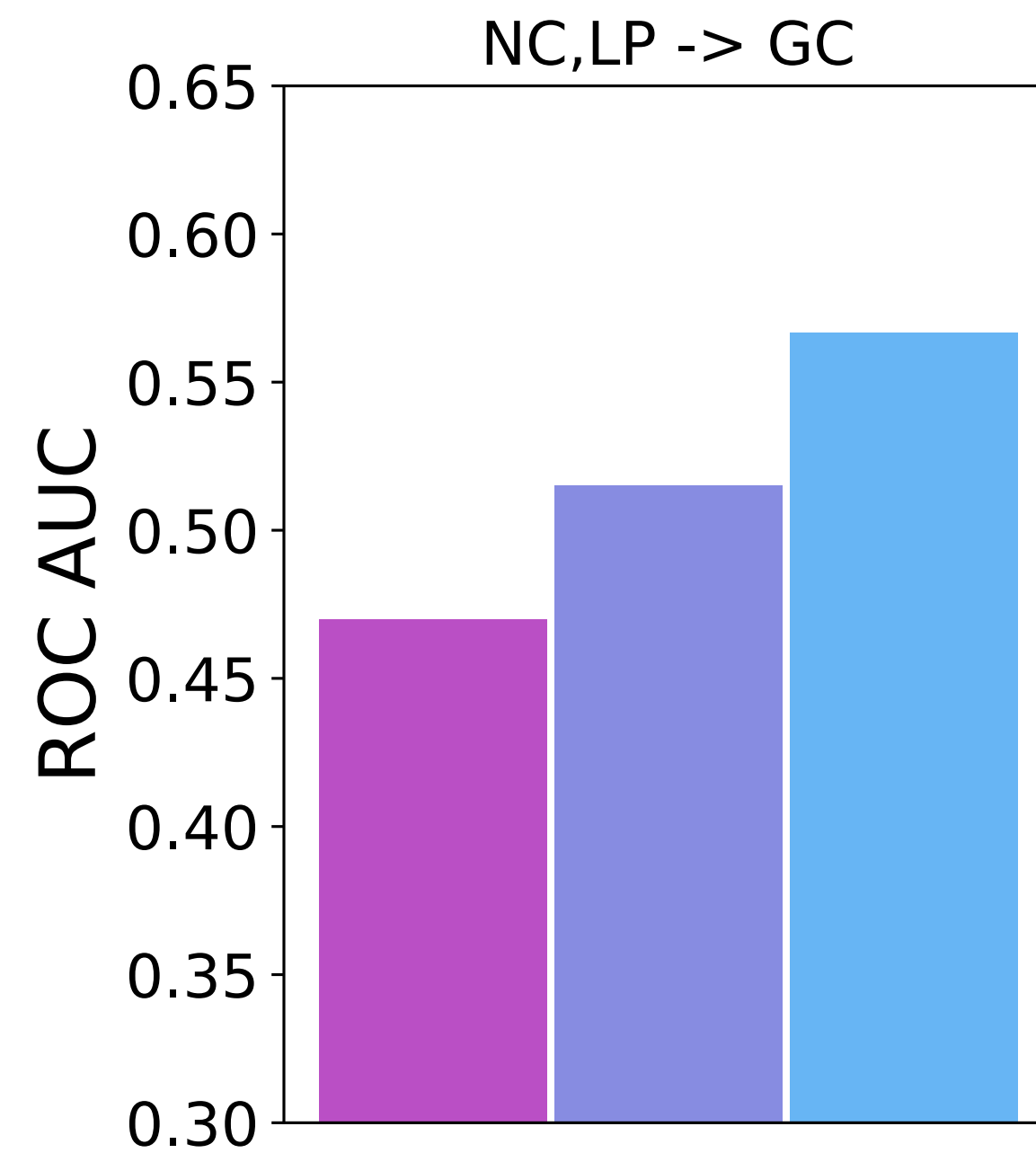
Do iSAME and eSAME lead to high quality node embeddings in the multi-task setting?

Table 2: Results for a single-task model trained in a classical supervised manner, a fine-tuned model (trained on all three tasks, and fine-tuned on the two shown tasks), and a **linear** classifier trained on node embeddings learned with our proposed strategies (iSAME, eSAME) in a multi-task setting.

| Task | | | Dataset | | | | | | | | | | | |
|-------------------------------|----|----|---------|------|------|----------|------|------|------|------|------|------|------|------|
| GC | NC | LP | ENZYMES | | | PROTEINS | | | DHFR | | | COX2 | | |
| | | | GC | NC | LP | GC | NC | LP | GC | NC | LP | GC | NC | LP |
| Classical End-to-End Training | | | | | | | | | | | | | | |
| ✓ | | | 51.6 | | | 73.3 | | | 71.5 | | | 76.7 | | |
| | ✓ | | | 87.5 | | | 72.3 | | | 97.3 | | | 96.4 | |
| | | ✓ | | | 75.5 | | | 85.6 | | | 98.8 | | | 98.3 |
| Fine-Tuning | | | | | | | | | | | | | | |
| ✓ | ✓ | | 48.3 | 85.3 | | 73.6 | 72.0 | | 66.4 | 92.4 | | 80.0 | 92.3 | |
| ✓ | | ✓ | 49.3 | | 71.6 | 69.6 | | 80.7 | 65.3 | | 58.9 | 80.2 | | 50.9 |
| | ✓ | ✓ | | 87.7 | 73.9 | | 80.4 | 81.5 | | 80.7 | 56.6 | | 87.4 | 52.3 |
| iSAME (ours) | | | | | | | | | | | | | | |
| ✓ | ✓ | | 50.1 | 86.1 | | 73.1 | 76.6 | | 71.6 | 94.8 | | 75.2 | 95.4 | |
| ✓ | | ✓ | 50.7 | | 83.1 | 73.4 | | 85.2 | 71.6 | | 99.2 | 77.5 | | 98.9 |
| | ✓ | ✓ | | 86.3 | 83.4 | | 79.4 | 87.7 | | 96.5 | 99.3 | | 95.5 | 99.0 |
| ✓ | ✓ | ✓ | 50.0 | 86.5 | 82.3 | 71.4 | 76.6 | 87.3 | 71.2 | 95.5 | 99.5 | 75.4 | 95.2 | 99.2 |
| eSAME (ours) | | | | | | | | | | | | | | |
| ✓ | ✓ | | 51.7 | 86.1 | | 71.5 | 79.2 | | 70.1 | 95.7 | | 75.6 | 95.5 | |
| ✓ | | ✓ | 51.9 | | 80.1 | 71.7 | | 85.4 | 70.1 | | 99.1 | 77.5 | | 98.8 |
| | ✓ | ✓ | | 86.7 | 82.2 | | 80.7 | 86.3 | | 96.6 | 99.4 | | 95.6 | 99.1 |
| ✓ | ✓ | ✓ | 51.5 | 86.3 | 81.1 | 71.3 | 79.6 | 86.8 | 70.2 | 95.3 | 99.5 | 77.7 | 95.7 | 98.8 |

Experiments

Do iSAME and eSAME extract information that is not captured by classically trained multi-task models?



Experiments

Can the node embeddings learned by iSAME and eSAME be used to perform multiple tasks with comparable or better performance than classical multi-task models?

Table 3: Δ_m (%) results for a classical multi-task model (CI), a fine-tuned model (FT; trained on all three tasks and fine-tuned on two) and a **linear** classifier trained on the node embeddings learned using our meta-learning strategies (iSAME, eSAME) in a multi-task setting.

| Task | | | Model | Dataset | | | |
|------|----|----|-------|-----------------|----------------|-----------------|-----------------|
| GC | NC | LP | | ENZYMES | PROTEINS | DHFR | COX2 |
| ✓ | ✓ | | CI | -0.1 ± 0.5 | 4.0 ± 1.0 | -0.3 ± 0.2 | 0.5 ± 0.1 |
| | | | FT | -4.5 ± 1.2 | 0.1 ± 0.5 | -7.4 ± 1.4 | 0.1 ± 0.4 |
| | | | iSAME | -2.3 ± 0.9 | 2.7 ± 1.5 | -1.2 ± 0.4 | -1.6 ± 0.2 |
| | | | eSAME | -0.8 ± 0.8 | 3.2 ± 1.4 | -1.8 ± 0.3 | -1.2 ± 0.3 |
| ✓ | | ✓ | CI | -25.3 ± 3.2 | -5.3 ± 1.2 | -28.3 ± 4.3 | -21.4 ± 3.4 |
| | | | FT | -5.1 ± 1.9 | -5.4 ± 1.5 | -24.5 ± 3.7 | -22.6 ± 3.8 |
| | | | iSAME | 4.1 ± 0.5 | -0.2 ± 0.9 | 0.2 ± 3.2 | 0.2 ± 0.5 |
| | | | eSAME | 3.2 ± 0.4 | -1.2 ± 1.1 | -0.7 ± 3.4 | -0.8 ± 0.7 |
| | ✓ | ✓ | CI | 7.2 ± 2.7 | 6.8 ± 0.9 | -29.1 ± 7.7 | -28.2 ± 4.5 |
| | | | FT | -1.0 ± 0.3 | 3.1 ± 1.2 | -28.9 ± 6.4 | -28.3 ± 4.2 |
| | | | iSAME | 4.4 ± 1.1 | 6.1 ± 1.0 | -0.1 ± 6.2 | -0.6 ± 2.5 |
| | | | eSAME | 3.9 ± 1.3 | 6.1 ± 1.1 | 0.1 ± 6.4 | -0.6 ± 2.6 |
| ✓ | ✓ | ✓ | CI | 1.6 ± 1.3 | 2.9 ± 0.3 | -18.9 ± 2.3 | -16.9 ± 3.1 |
| | | | iSAME | 1.5 ± 1.0 | 2.2 ± 0.2 | -0.5 ± 1.4 | -0.9 ± 1.3 |
| | | | eSAME | 1.8 ± 0.9 | 2.8 ± 0.2 | -1.0 ± 1.7 | -0.4 ± 1.2 |

Thank you for watching!

Don't hesitate to come to our virtual booth and have a chat.

**You can also contact us at:
{davide.buffelli, fabio.vandin}@unipd.it**