





Towards Sparse Hierarchical Graph Classifiers

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Graph classification

- Typical machine learning task aimed at graph-structured data (e.g. want to classify molecular structures, citation networks, social circles)
- Graphs represent the generalization of images (grid-like graphs) → aim to extend readily established *image classification* techniques
- Typical image classification pipeline: alternate between convolution (feature detection) and pooling (down-sampling)—want this for graphs!



Existing work

- Graph convolutional layers: plenty!
 - Spectral: Bruna et al.(2014), Defferrard et al.(2016), Kipf et al.(2017)
 - *Message-passing*: Gilmer et al.(2017)
 - o Attention-based: Monti et al.(2016), Veličković et al.(2017)
- ► Pooling mechanisms: comparatively few...
 - Globally pool after each/the final message-passing step
 - Progressively coarsen the graph—most approaches assume a pre-defined assignment of nodes to clusters
 - Learn a differentiable pooling mechanism based on the *structure of the data*: Ying et al.(2018), Anon.(2018)



Hierarchical (pooling) strategies

- DiffPool: compute soft clustering assignments of nodes from the original graph to nodes from the resulting graph
- End-to-end trainable architecture with SOTA results
- Main issue: soft clustering assignments require storing the assignment matrix and therefore O(kV²) memory!
- Graph U-Net: does not suffer from the issue above, but the architecture was not evaluated on classification task benchmarks



Our contributions

- An intuitive architecture for graph classification that closely resembles the computation model for image classification
- Results comparable to state-of-the-art on classification benchmark tasks
- A drastic reduction in the GPU memory requirement (from O(V²) to O(V + E))





- Input graph → a matrix of node features, X ∈ ℝ^{N×F}, and an adjacency matrix, A ∈ ℝ^{N×N}
- A is binary and symmetric
- If the graph is featureless, use 1-hot encoding of node degree information to construct X
- A CNN-inspired network for graph classification should contain the following layers: *convolutional*, *pooling*, *readout* (i.e. flattening layer in an image CNN used for the final prediction)



Model

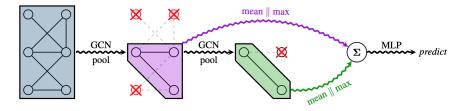


Figure 1: The full pipeline of our model (for k = 0.5), leveraging several stacks of interleaved convolutional/pooling layers (that, unlike DiffPool, *drop* rather than aggregate nodes), as well as a JK-net-style summary, combining information at different scales.



Convolutional layer

 Apply the *mean-pooling* propagation rule (as in GCN or Const-GAT):

$$MP(\mathbf{X}, \mathbf{A}) = \sigma(\hat{\mathbf{D}}^{-1}\hat{\mathbf{A}}\mathbf{X}\Theta + \mathbf{X}\Theta')$$

- $\hat{\mathbf{A}} = \mathbf{A} + \mathbf{I}_N$ is the adjacency matrix with self-loops
- $\hat{\mathbf{D}}$ is the degree matrix: $\hat{D}_{ii} = \sum_{j} \hat{A}_{ij}$



- ► Use Graph U-Net architecture to reduce the number of nodes N via pooling ratio k ∈ (0, 1]
- ► Layer outputs [*kN*] nodes
- ► Learn projection score p used as gating values → ensure lower scoring nodes retain comparatively less features



Pooling layer—cont'd

Obtain pooled graph (X', A') as:

$$\vec{y} = \frac{\mathbf{X}\vec{p}}{\|\vec{p}\|},$$
$$\vec{i} = \operatorname{top-}k(\vec{y}, k),$$
$$\mathbf{X}' = (\mathbf{X} \cdot \operatorname{tanh}(\vec{y}))_{\vec{i}},$$
$$\mathbf{A}' = \mathbf{A}_{\vec{i},\vec{i}}$$

This only requires a pointwise projection operation and slicing into the original matrices!



Readout layer

- ► Want "flattening" operation analogous to image CNNs
- Take the average and max of all learnt node embeddings for output graph (X^k, A^k) of the k-th block:

$$ec{s}^k = rac{1}{N^k} \sum_{i=1}^{N^k} ec{x}^k_i \| \max_{i=1}^{N^k} ec{x}^k_i \|$$

Inspired by JK-net, summarise the graph by summing:

$$ec{s} = \sum_{k=1}^{K} ec{s}^k$$

• An MLP predicts the class for \vec{s} at the "tail" of the model



Experiments

- Benchmark datasets: Enzymes, Proteins, DD, Collab
- ▶ 10-fold cross-validation, compare to Ying et al.(2018)

- ► Three {GCN layer, pooling layer} blocks
- Preserve k = 80% of the nodes on each pool



Results

	Datasets			
Model	Enzymes	D&D	Collab	Proteins
Graphlet	41.03	74.85	64.66	72.91
Shortest-path	42.32	78.86	59.10	76.43
1-WL	53.43	74.02	78.61	73.76
WL-QA	60.13	79.04	80.74	75.26
PatchySAN	_	76.27	72.60	75.00
GraphSAGE	54.25	75.42	68.25	70.48
ECC	53.50	74.10	67.79	72.65
Set2Set	60.15	78.12	71.75	74.29
SortPool	57.12	79.37	73.76	75.54
DiffPool-Det	58.33	75.47	82.13	75.62
DiffPool-NoLP	62.67	79.98	75.63	77.42
DiffPool	64.23	81.15	75.50	78.10
Ours	64.17	78.59	74.54	75.46



Memory usage

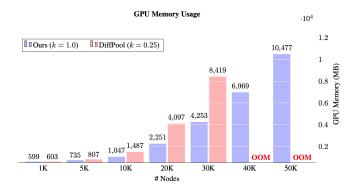
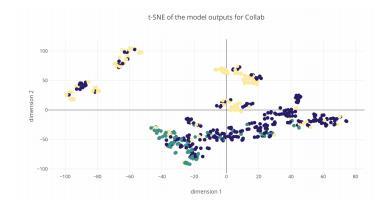


Figure 2: GPU memory usage of our method (with no pooling; k = 1.0) and DiffPool (k = 0.25) during training on Erdős-Rényi graphs [9] of varying node sizes (and |E| = 2|V|). Both methods ran with 128 input and hidden features, and three Conv-Pool layers. "OOM" denotes out-of-memory.



Qualitative analysis





Future directions

- Apply the computation model to large datasets (particularly relevant for the neuroscience domain, where brain meshes are encoded by graphs with +100K nodes).
- Improve the pooling mechanism (e.g. by making the pooling ratio k learnable).
- Make the "mirroring" operation (unpooling) independent of the pooling results.





Questions?

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