Scaling Graph Transformers with Expander Graphs

Danica Sutherland UBC + Amii (she/her)

based on:

Exphormer: Scaling Graph Transformers with Expander Graphs (ICML 2023; arXiv:2303.06147) **Even Sparser Graph Transformers** (Spexphormer; not quite on arXiv yet, but soon!)

with:



Hamed Shirzad UBC



Ameya Velingker Google



Balaji venkatachalam Google \rightarrow Meta

SFU VCR/AI seminar, June 2024



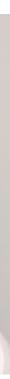
Ali Sinop Google (Exphormer only)



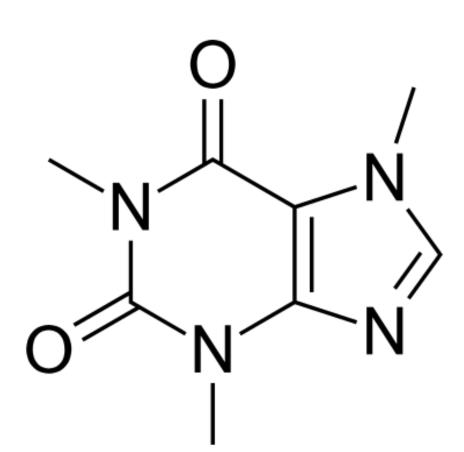
Honghao Lin CMU (Spexphormer only)



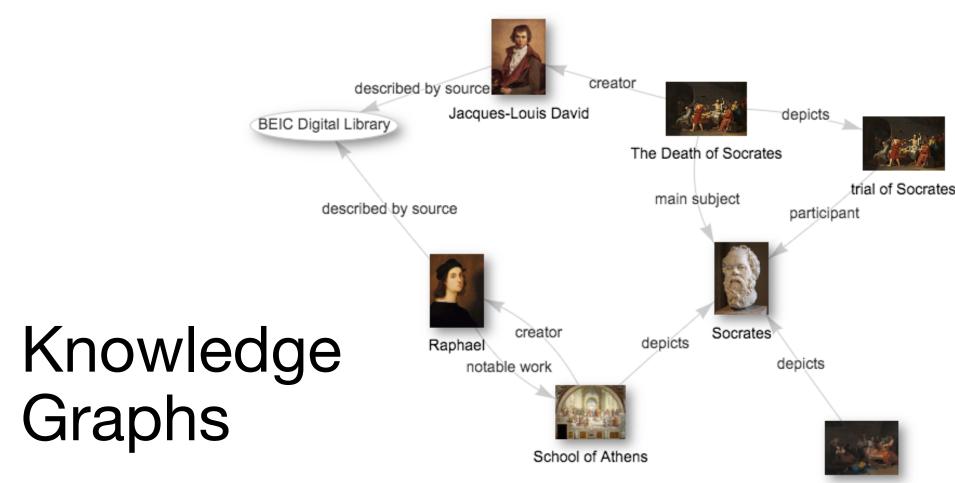
David woodrum CMU + Google (Spexphormer only)



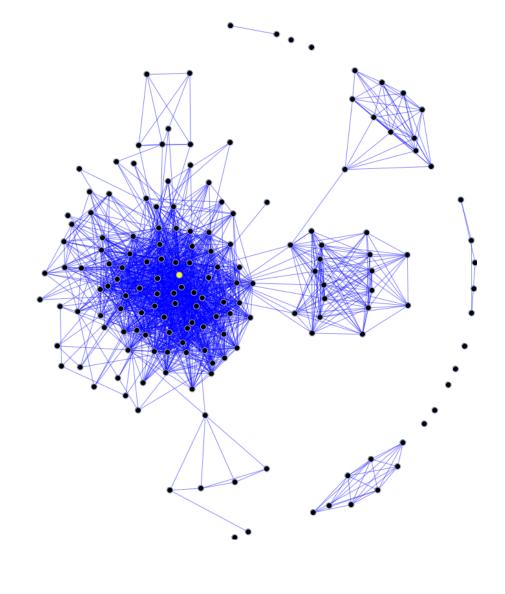
Learning on graphs



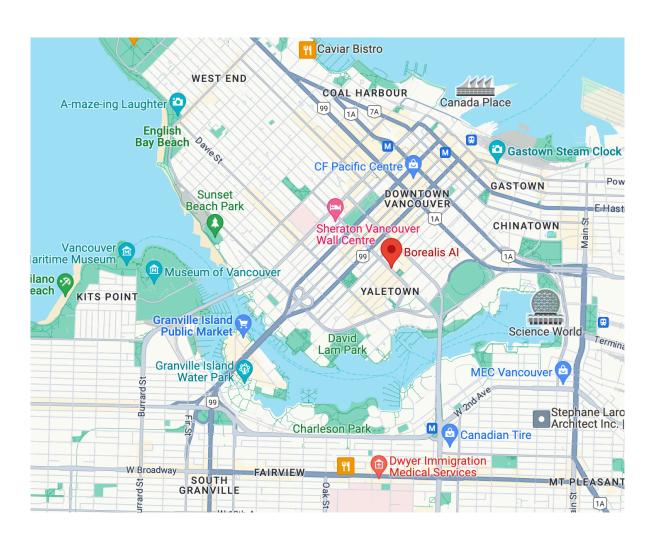
Molecular Graphs



The Death of Socrates



Social Network Graphs

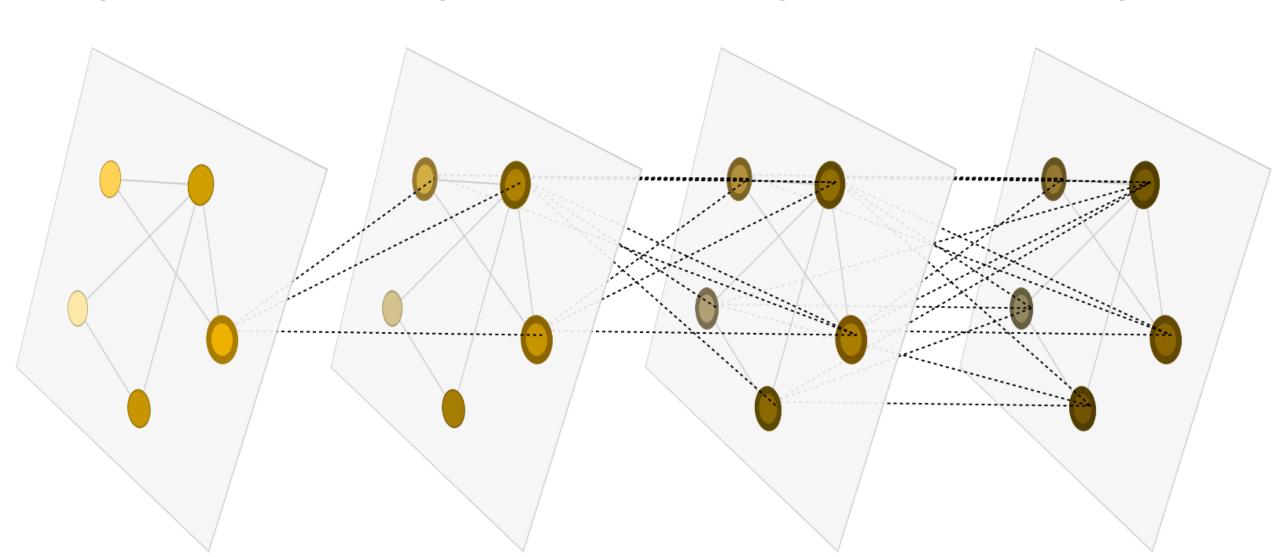


Road Network Graphs

Message-passing neural networks

J

Layer 1



• Features at each node x_i , maybe also each edge $e_{i \rightarrow j}$

Layer 0

New features
$$x_i^+ = f_v(x_i, \sum_i m_{j \to i})$$

Layer 2

Layer 3

• Message from node j to node i: $m_{i \to i} = f_e(x_i, x_i, e_{i \to i})$ for some NN f_e

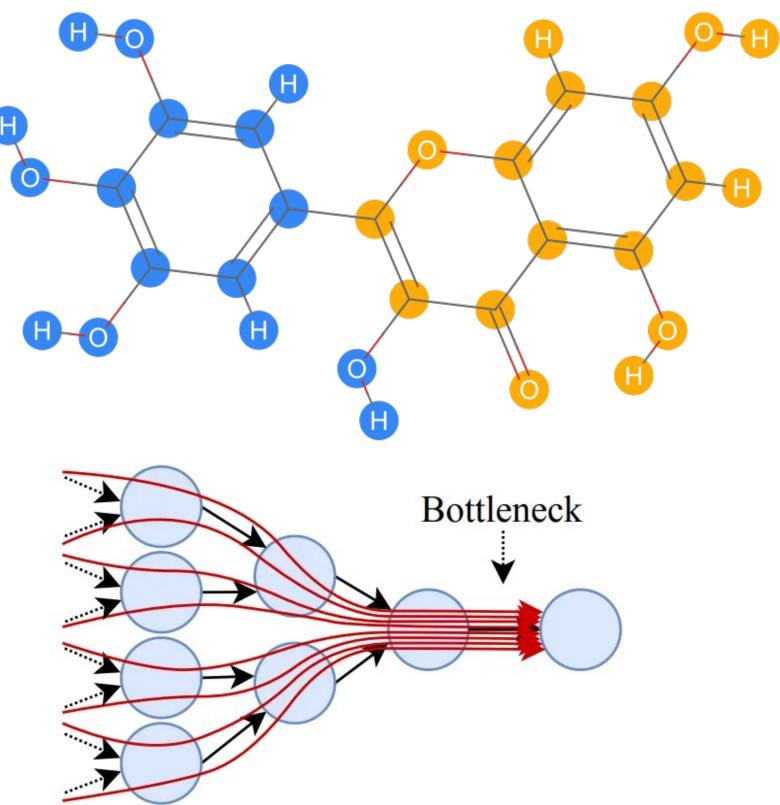
 f_i) for some NN f_v

Problems with message-passing neural nets

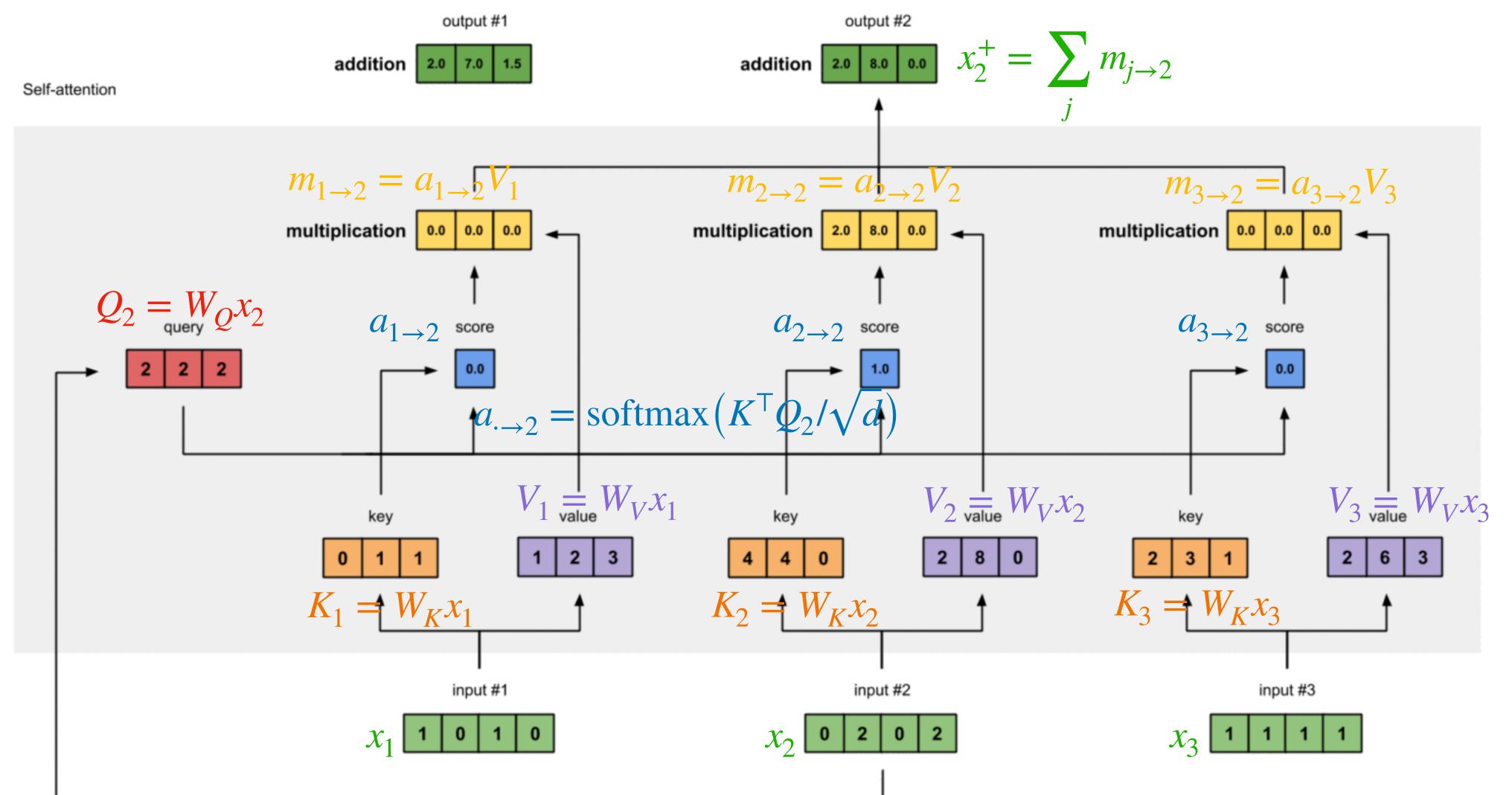
- Information only propagates along the graph edges
 - What if there are long-range dependencies?

- Over-smoothing
 - In deeper layers, all node features often end up basically the same as each other
- Over-squashing

Images from https://www.wolfram.com/language/12/molecular-structure-and-computation/molecule-graphs.html.en and Long-Range Graph Benchmark



Self-attention layers



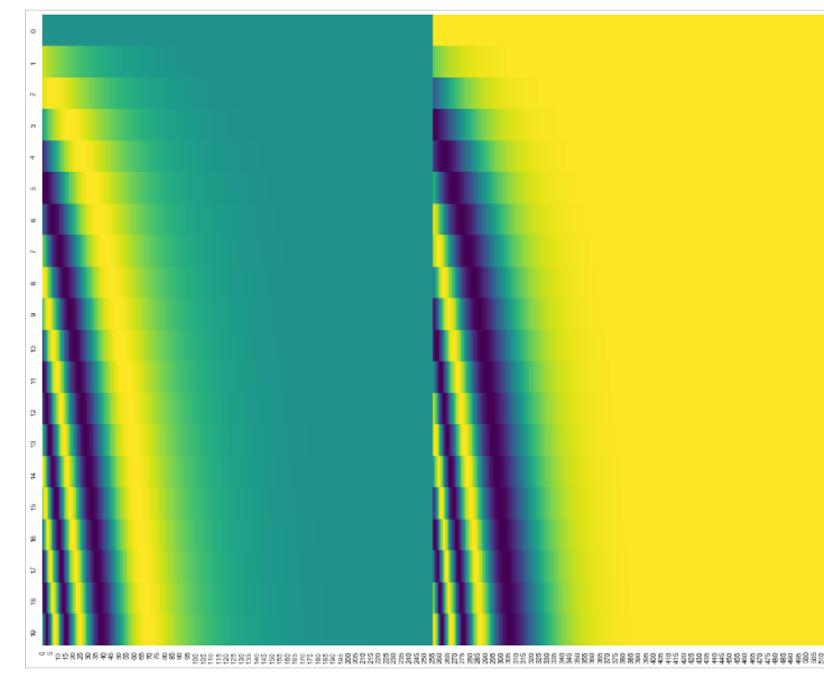
Looks like message-passing over the full graph



Positional encodings for sequences

- Problem: the order of the sequence is lost!
- Solution: add features for positional encodings that tell you "where you are"
- "Default" version: trigonometric features

$$PE_{(pos,2i)} = sin(pos_{(pos,2i+1)}) = cos(pos_{(pos,2i+1)}) = cos(pos_{(pos_{(pos,2i+1)})}) = cos(pos_{(pos_{(pos_{(pos,2i+1)})}) = cos(pos_{(pos_$$







Positional encodings

Eigenvalues and eigenfunctions of the Laplacian

Andrew Hassell

The setting

In this talk I will consider the Laplace operator, Δ , on various geometric spaces M. On a Euclidean domain,

$$\Delta f = -\sum_{i=1}^{n} \frac{\partial^2 f}{\partial x_i^2}.$$

Trigonometric functions: eigenfunctions of Laplacian on Euclidean space

Examples

• The interval [0, a]. Eigenfunctions and eigenvalues are

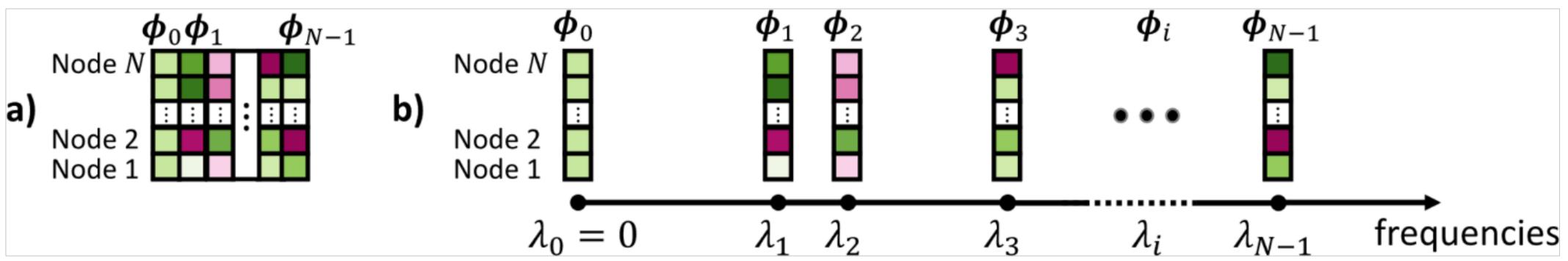
$$u_n = \sqrt{\frac{2}{a}} \sin \frac{\pi n x}{a}, \quad \lambda_n = \frac{\pi n}{a}.$$

• The torus T_{π}^2 . Eigenfunctions and eigenvalues are

$$u = \frac{1}{\pi} e^{ilx} e^{imy}, \quad \lambda = \sqrt{l^2 + m^2}.$$

Positional encodings

- Trigonometric functions: eigenfunctions of Laplacian on Euclidean space What should we do so that a self-attention layer "knows" about the
- structure of a graph?
 - "Default" choice: eigenvectors of the graph Laplacian L = D A



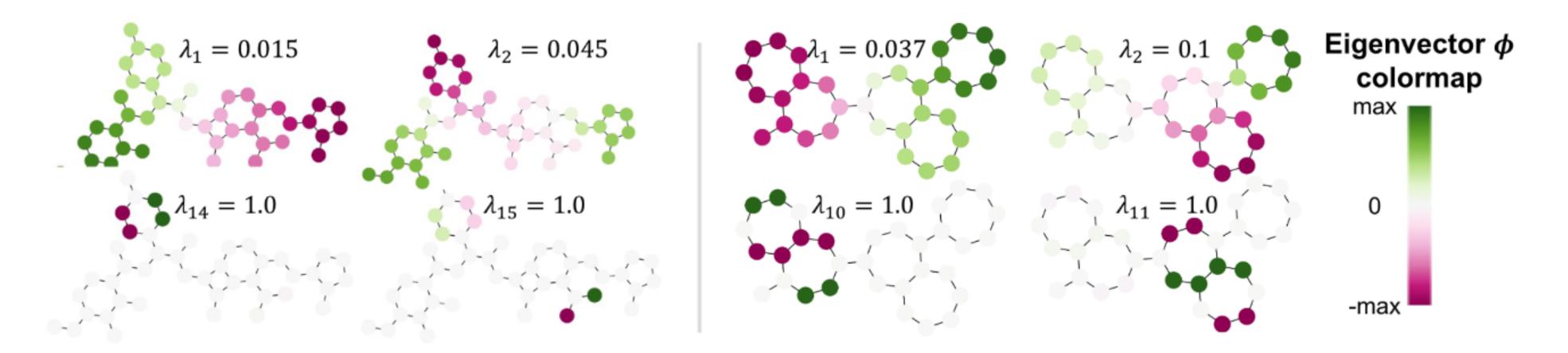
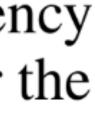
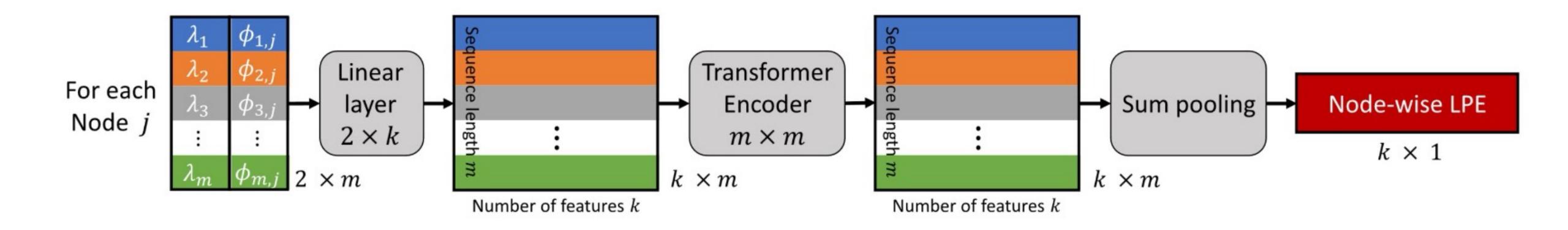


Figure 3: Examples of eigenvalues λ_i and eigenvectors ϕ_i for molecular graphs. The low-frequency eigenvectors ϕ_1, ϕ_2 are spread accross the graph, while higher frequencies, such as ϕ_{14}, ϕ_{15} for the left molecule or ϕ_{10} , ϕ_{11} for the right molecule, often resonate in local structures.



Learned positional encodings (Kreuzer et al. 2021)

Can be helpful to post-process the positional encodings further

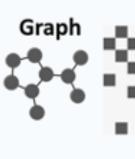


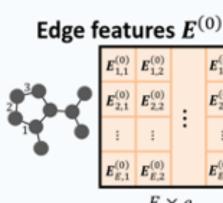
Spectral Attention Networks (SAN) (Kreuzer et al. 2021)

- Self-attention layers with learned positional encodings
- Intersperse with MLPs processing each node feature independently, as in Transformer encoders

Input graph

A: Adjacency matrix L: Laplacian matrix N: number of nodes E: number of edges. n_0 : Number of input node features e_0 : Number of input edge features 0: Computation complexity





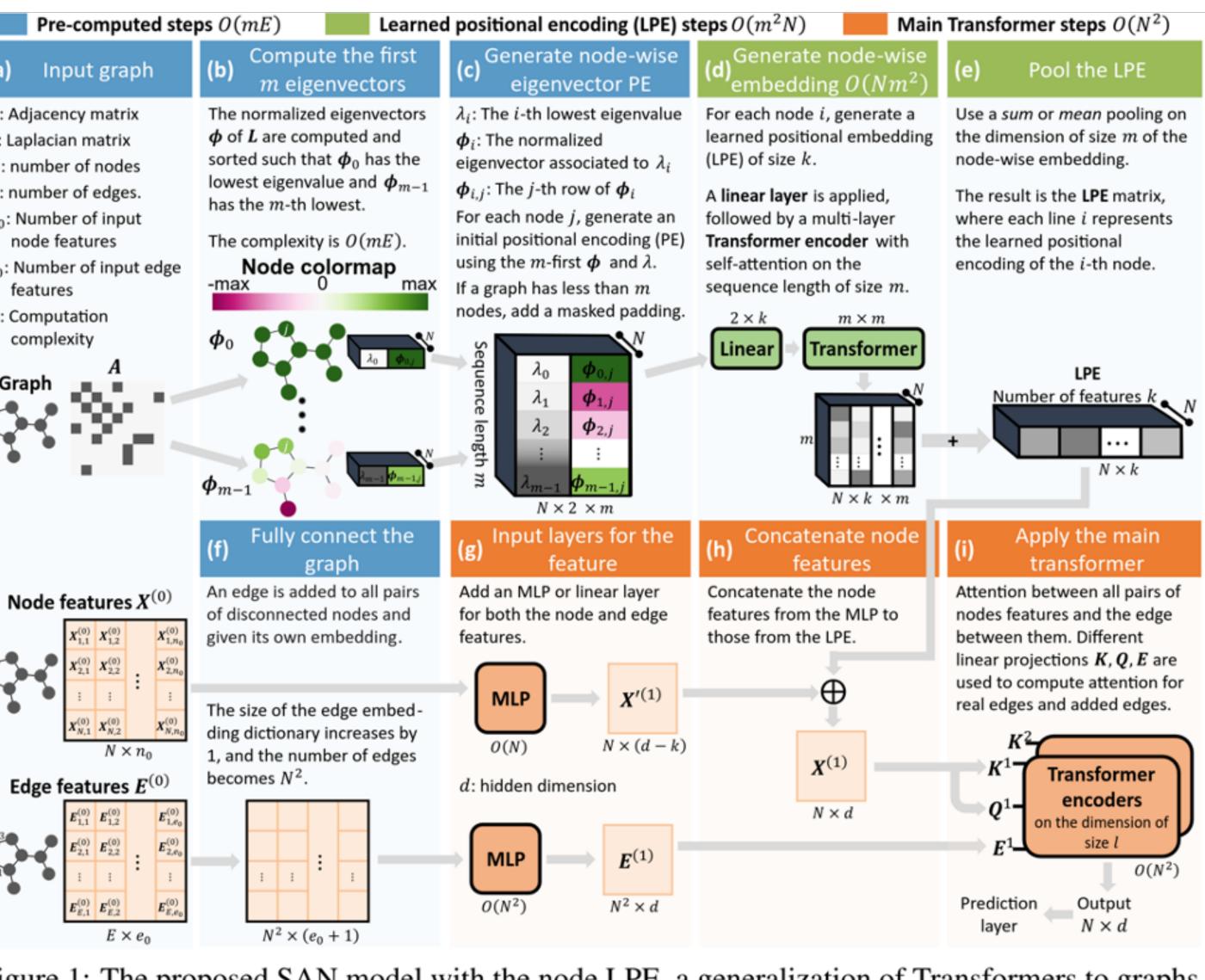
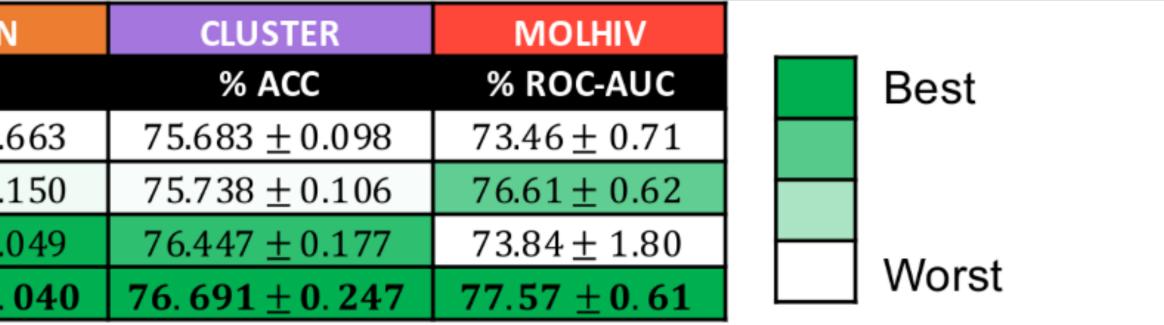


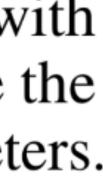
Figure 1: The proposed SAN model with the node LPE, a generalization of Transformers to graphs.

Full attention and position encodings help

Model det	ails	ZINC	PATTERN	
Attention	LPE	MAE	% ACC	
Sparse	-	0.267 ± 0.032	83.613 ± 0.0	
Sparse	Node	0.198 <u>+</u> 0.004	81.329 ± 2.1	
Full	-	0.392 ± 0.055	86.322 ± 0.0	
Full	Node	0.157 ± 0.006	86 . 441 ± 0 .	

Figure 6: Ablation study on datasets from [15, 21] for the node LPE and full graph attention, with no hyperparameter tuning other than γ taken from Figure 5. For a given dataset, all models use the same hyperparameters, but the hidden dimensions are adjusted to have $\sim 500k$ learnable parameters. Means and uncertainties are derived from four runs, with different seeds (except MolHIV).





Message passing vs graph transformers

Message passing networks

Update across edges of input graph

Capture inductive bias from input graph topology

Solution: Efficient computation: O(N + M)

X Difficulty with long-range dependencies

X Oversmoothing, oversquashing

Positional and structural encodings

<u>Graph transformers</u> Use global attention



from input graph

Long-range modeling

V Universal Approximation Theorem







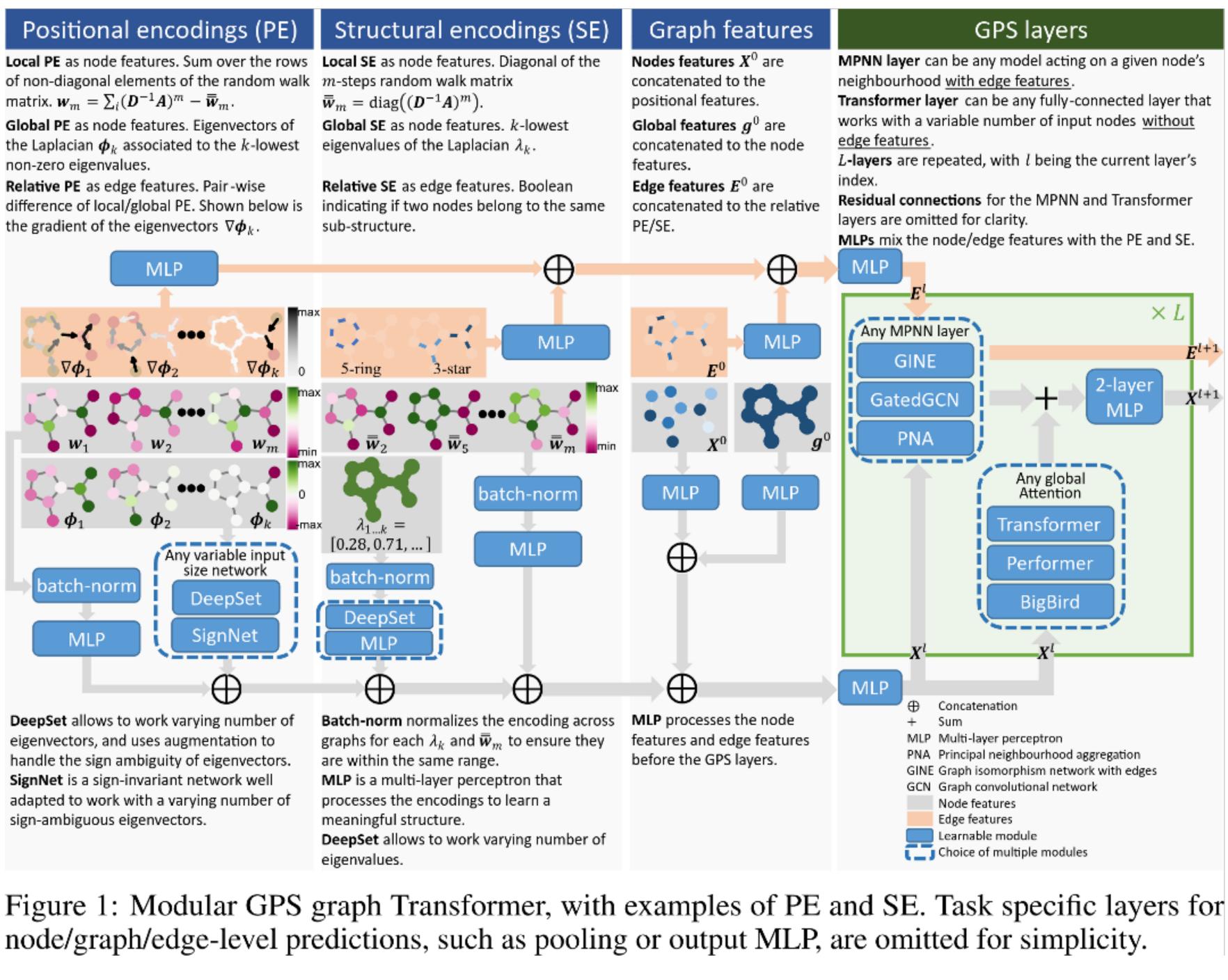
Inefficient computation: O(N²)



matrix. $\boldsymbol{w}_m = \sum_i (\boldsymbol{D}^{-1} \boldsymbol{A})^m - \overline{\boldsymbol{w}}_m$.

Global PE as node features. Eigenvectors of the Laplacian $\boldsymbol{\phi}_k$ associated to the k-lowest non-zero eigenvalues.

the gradient of the eigenvectors $\nabla \phi_k$.



GraphGPS

(Rampášek et al., 2022)

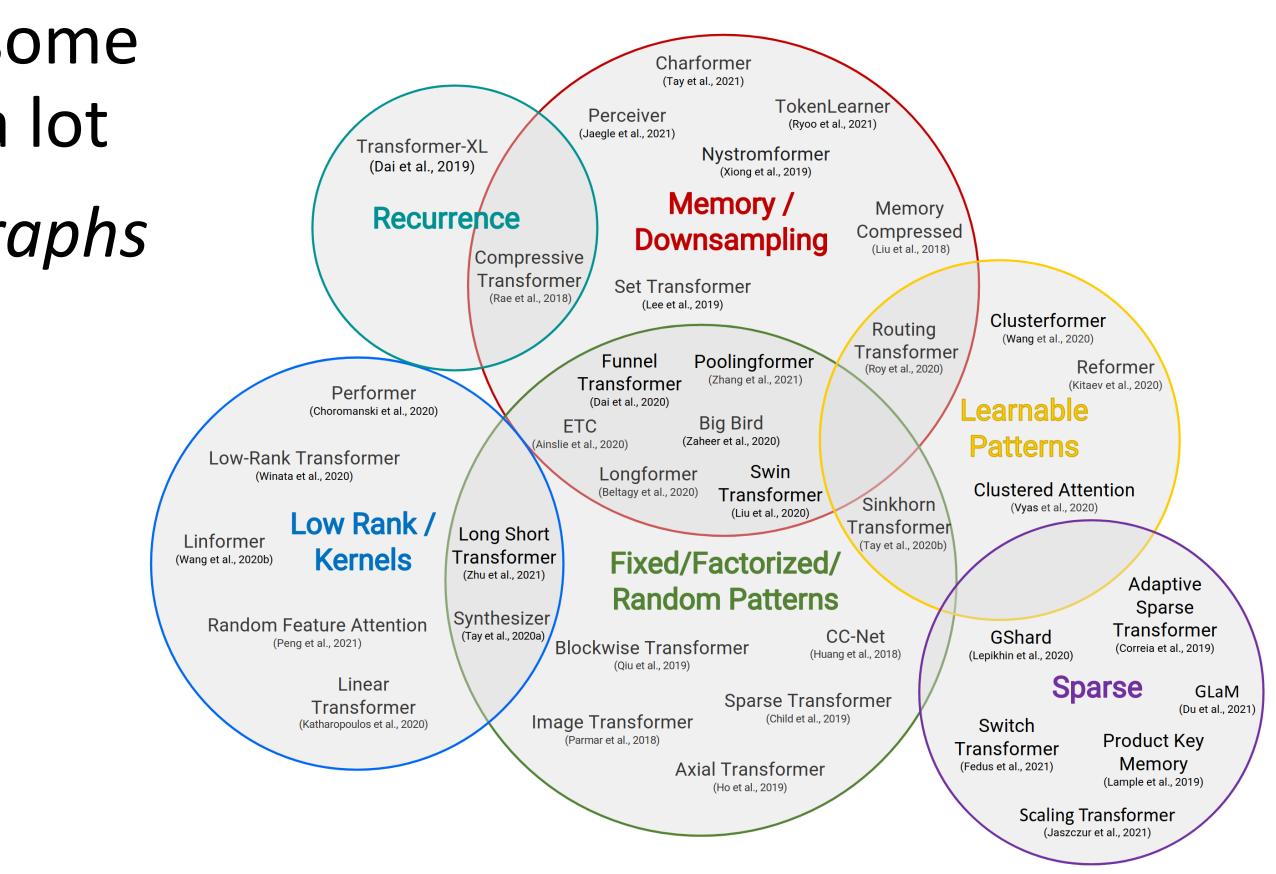
- Framework to combine transformer layers with messagepassing layers
- MPNN layers better at preserving graph structure
- Transformer layers better at certain kinds of dependence

Sparse attention and Exphormer

Reducing the N^2 attention cost

- GraphGPS paper tried to apply some to graphs...but it hurt accuracy a lot
- Need something *designed for graphs*

Lots of work on more-efficient attention mechanisms for sequences

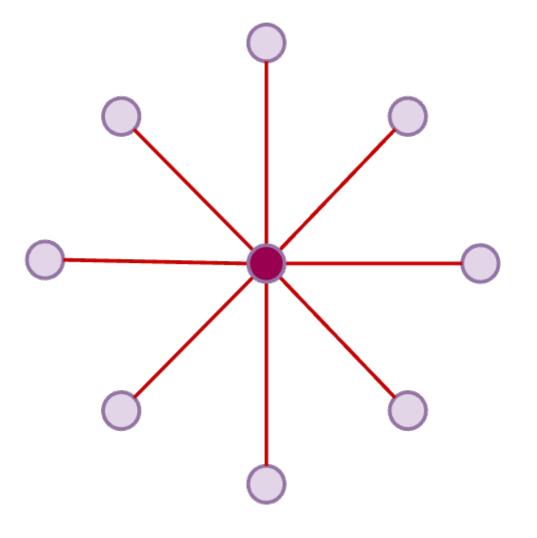


Attention graph

- We'll take an approach of "sparse attention"
- Start by "running" attention along the graph edges (both ways)
 Adjacent nodes are reasonably likely to "matter" to each other
- Full transformers augment this attention graph with the complete graph
 - Every node connected to every other
 - Allows easy long-range communication
 - Adds too many edges! Makes it slow + memory-hungry

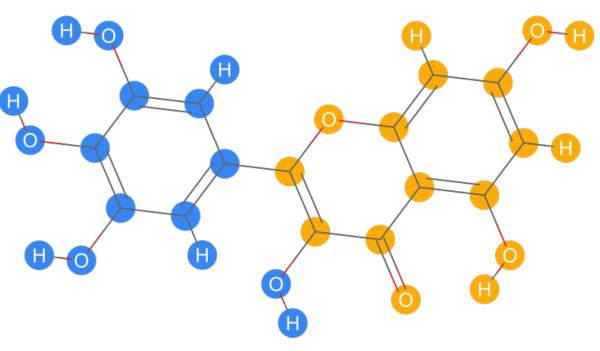
Augmenting attention with virtual nodes

- Add a "virtual" node (or four) that's connected to every graph node
- Now any node can "talk to" any other node in at most 2 hops
- Also common in MPNNs
- Problem: oversquashing



Expander graphs

- How can we connect far-away parts of the graph without severe oversquashing?
- We want a sparse approximation to the complete graph
 - No too many edges
 - Can get from anywhere to anywhere quickly
 - Have many routes from place to place, to avoid oversquashing
- Mathematical concept called expander graphs
 - Approximate the complete graph (in many senses), with low degree



Expander graphs

Theorem 4.1. A d-regular ϵ -expander G on n vertices spectrally approximates the complete graph K_n on n vertices:²

$$(1-\epsilon)\frac{1}{n}L_K \preceq \frac{1}{d}L_G \preceq (1+\epsilon)\frac{1}{n}L_K.$$

Spectral approximation is known to preserve the cut structure in graphs. As a result, a sparse attention mechanism based on expander edges retains spectral properties of the full attention mechanism: cuts, vertex expansion, and so on.

Lemma 4.2. (*Alon*, 1986) *Let* G = (V, E) *be a d-regular* ϵ -expander graph on n = |V| nodes. For any initial distribution $\pi^{(0)}: V \to \mathbb{R}^+$ and any $\delta > 0$, $\pi^{(t)}$ satisfies

$$\|\pi^{(t)} - \frac{1}{n}\|_1 \le \delta$$

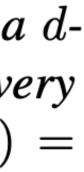
as long as $t = \Omega\left(\frac{1}{\epsilon}\log(n/\delta)\right)$.

Theorem 4.3. (Alon, 1986) Suppose G = (V, E) is a dregular ϵ -expander graph on n vertices. Then, for every vertex v and $k \geq 0$, the k-hop neighborhood B(v,r) = $\{w \in V : d(v, w) \leq k\}$ has

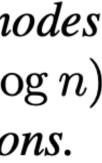
 $|B(v,r)| \ge \min\{(1+c)^k, n\}$

for some constant c > 0 depending on d, ϵ . In particular, we have that $diam(G) = O_{d,\epsilon}(\log n)$.

Corollary 4.4. If a sparse attention mechanism on n nodes is a d-regular ϵ -expander graph, then stacking $O_{d,\epsilon}(\log n)$ transformer layers models all pairwise node interactions.







Expander graphs

CS366: Graph Partitioning and Expanders

[general info] [lecture notes] [exams and projects]

what's new

• 2/14 midterm

general information

Instructor: Luca Trevisan, Gates 474, Tel. 650 723-8879, email trevisan at stanford dot edu

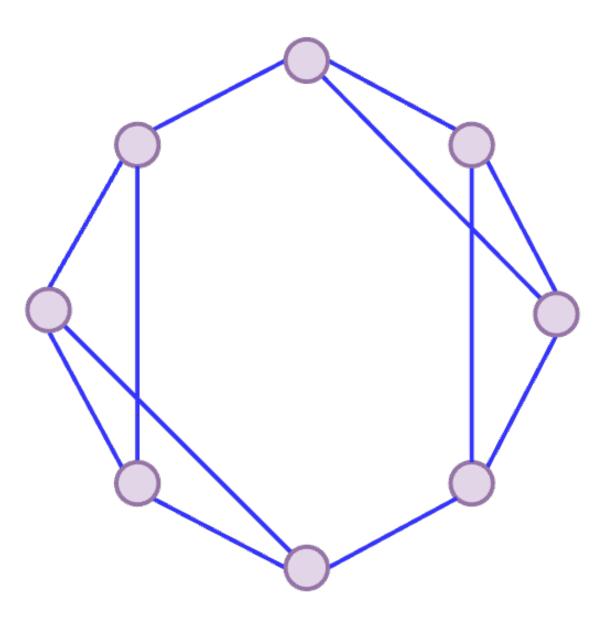
Expander graphs turn out to have a slew of applications not only in math but also in computer science and physics. They can be used to create error-correcting codes and to figure out when simulations based on random numbers converge to the reality they are trying to simulate. Neurons can be modeled in a graph that some researchers believe forms an expander, due to the limited space for connections inside the brain. The graphs are also useful to geometers who try to understand how to traverse complicated surfaces, among other problems.

https://www.quantamagazine.org/new-proof-shows-that-expander-graphs-synchronize-20230724/

Constructing expander graphs

- How to make an expander graph with max degree $\leq d$:
 - Sample d/2 random Hamiltonian cycles, combine these as edges Check (from the eigenvalues of the Laplacian) if it's an expander

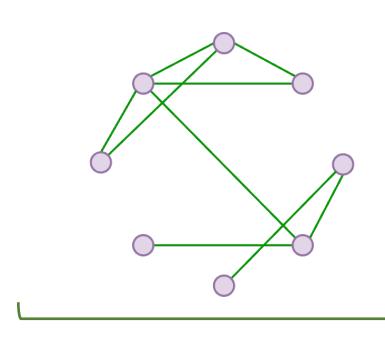
 - If not, try again (happens rarely)



Components of Exphormer

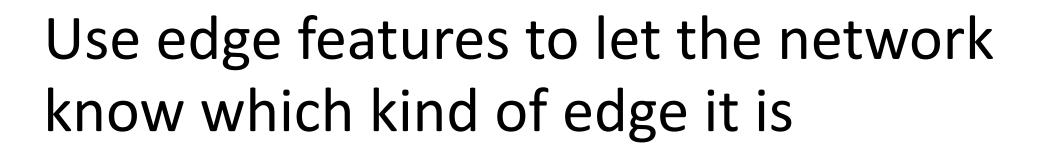
Original graph

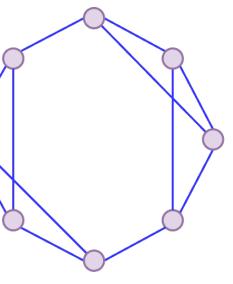
• Preserve locality from the original graph



Expander graph

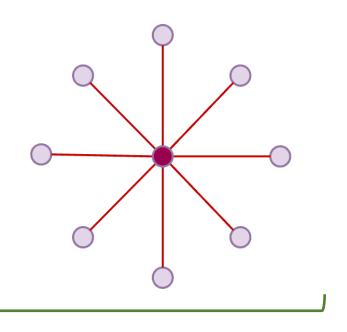
- Random walk mixing
- Constant degree
- O(N) edges





Virtual node(s)

- "Storage sink"
- Short connections between all node pairs



Exphormer: Combine all three to form the interaction graph

Universal approximation

Theorem E.3. Suppose H is the attention graph of EXPHORMER (which contains n graph nodes and potentially more virtual nodes), augmented with self loops on all nodes. Suppose H satisfies at least one of the following:

- 2. The underlying expander graph of H contains a Hamiltonian path.

Then, it follows that a sparse transformer model, with positional encodings and an attention mechanism following H, can universally approximate continuous functions $f: [0,1]^{d \times n} \to \mathbb{R}^{d \times n}$. That is, for any $1 and <math>\epsilon > 0$, there exists a sparse transformer network g, which uses the attention graph H and some positional encodings, such that $\ell^p(f,g) < \epsilon$.

1. H contains at least one node which is connected to all n graph nodes (i.e., at least one virtual node is included).



Experimental Results: Various Datasets

Table 1. Comparison of EXPHORMER with baselines on various datasets. Best results are colored in first, second, third.

Model	CIFAR10 Accuracy ↑	MalNet-Tiny Accuracy ↑	MNIST Accuracy ↑	CLUSTER Accuracy ↑	PATTERN Accuracy ↑
GCN (Kipf & Welling, 2017) GIN (Xu et al., 2018)	$55.71 {\pm} 0.381$ $55.26 {\pm} 1.527$	81.0 88.98±0.557	$90.71 {\pm} 0.218$ $96.49 {\pm} 0.252$	$\begin{array}{c} 68.50 \pm 0.976 \\ 64.72 \pm 1.553 \end{array}$	$\begin{array}{c} 71.89 \pm 0.334 \\ 85.39 \pm 0.136 \end{array}$
GAT (Veličković et al., 2018) GatedGCN (Bresson & Laurent, 2017;	64.22 ± 0.455 67.31 ± 0.311	92.1 ± 0.242 92.23 ± 0.65	$95.54{\pm}0.205$ $97.34{\pm}0.143$	$\begin{array}{r} 70.59 \pm 0.447 \\ 73.84 \pm 0.326 \end{array}$	$\begin{array}{r} 78.27 \pm 0.186 \\ 85.57 \pm 0.088 \end{array}$
Dwivedi et al., 2020) PNA (Corso et al., 2020) DGN (Beaini et al., 2021)	$70.35{\pm}0.63$ 72.84 ${\pm}0.417$	_	97.94±0.12	_	-86.68 ± 0.034
CRaWl (Toenshoff et al., 2021)	69.01±0.259	_		_	_
GIN-AK+ (Zhao et al., 2022b)	72.19±0.13	_	_	-	86.85±0.057
SAN (Kreuzer et al., 2021) K-Subgraph SAT (Chen et al., 2022a) EGT (Hussain et al., 2021) GraphGPS (Rampásek et al., 2022)	- 68.70±0.409 72.30±0.356	_ 93.50±0.41	- 98.17±0.087 98.05±0.126	76.69 ± 0.65 77.86 ± 0.104 79.23 ± 0.348 78.02 ± 0.180	$\begin{array}{r} 86.58 {\pm} 0.037 \\ \textbf{86.85 {\pm} 0.037} \\ \textbf{86.82 {\pm} 0.020} \\ 86.69 {\pm} 0.059 \end{array}$
EXPHORMER (ours)	74.69±0.125	$\textbf{94.02} \pm \textbf{0.209}$	$\textbf{98.55} \pm \textbf{0.039}$	$\textbf{78.07} \pm \textbf{0.037}$	86.74±0.015

Experimental Results: Long-Range Benchmark

Table 3. Comparison of EXPHORMER with baselines from the Long-Range Graph Benchmarks (LRGB, Dwivedi et al., 2022). Best results are colored in **first**, **second**, **third**.

Model	PascalVOC-SP F1 score ↑	COCO-SP F1 score ↑	Peptides-Func AP ↑	Peptides-Struct MAE ↓	PCQM-Contact MRR ↑
GCN GINE GatedGCN GatedGCN+RWSE	$\begin{array}{c} 0.1268 \pm 0.0060 \\ 0.1265 \pm 0.0076 \\ 0.2873 \pm 0.0219 \\ 0.2860 \pm 0.0085 \end{array}$	$\begin{array}{c} 0.0841 \pm 0.0010 \\ 0.1339 \pm 0.0044 \\ \textbf{0.2641} \pm \textbf{0.0045} \\ 0.2574 \pm 0.0034 \end{array}$	$\begin{array}{c} 0.5930 \pm 0.0023 \\ 0.5498 \pm 0.0079 \\ 0.5864 \pm 0.0077 \\ 0.6069 \pm 0.0035 \end{array}$	$\begin{array}{c} 0.3496 \pm 0.0013 \\ 0.3547 \pm 0.0045 \\ 0.3420 \pm 0.0013 \\ 0.3357 \pm 0.0006 \end{array}$	$\begin{array}{c} 0.3234 \pm 0.0006 \\ 0.3180 \pm 0.0027 \\ 0.3218 \pm 0.0011 \\ 0.3242 \pm 0.0008 \end{array}$
Transformer+LapPE SAN+LapPE SAN+RWSE GraphGPS	$\begin{array}{c} 0.2694 \pm 0.0098 \\ \textbf{0.3230} \pm \textbf{0.0039} \\ 0.3216 \pm 0.0027 \\ \textbf{0.3748} \pm \textbf{0.0109} \end{array}$	$\begin{array}{c} 0.2618 \pm 0.0031 \\ 0.2592 \pm 0.0158 * \\ 0.2434 \pm 0.0156 * \\ \textbf{0.3412} \pm \textbf{0.0044} \end{array}$	$\begin{array}{c} 0.6326 \pm 0.0126 \\ 0.6384 \pm 0.0121 \\ \textbf{0.6439} \pm \textbf{0.0075} \\ \textbf{0.6535} \pm \textbf{0.0041} \end{array}$	$\begin{array}{l} \textbf{0.2529} \pm \textbf{0.0016} \\ 0.2683 \pm 0.0043 \\ 0.2545 \pm 0.0012 \\ \textbf{0.2500} \pm \textbf{0.0005} \end{array}$	$\begin{array}{c} 0.3174 \pm 0.0020 \\ \textbf{0.3350} \pm \textbf{0.0003} \\ \textbf{0.3341} \pm \textbf{0.0006} \\ 0.3337 \pm 0.0006 \end{array}$
Exphormer (ours)	$\textbf{0.3975} \pm \textbf{0.0037}$	$\textbf{0.3455} \pm \textbf{0.0009}$	$\textbf{0.6527} \pm \textbf{0.0043}$	$\textbf{0.2481} \pm \textbf{0.0007}$	$\textbf{0.3637} \pm \textbf{0.0020}$

Experimental Results: Larger Graphs

Table 4. Accuracy of models with different attention mechanisms on transductive graph datasets (numbers in top rows, other than arXiv, are from Chen et al., 2022b). Chen et al. did not report NAGphormer results on this dataset.

-	-	-			
Model	ogbn-arxiv	Computer	Photo	CS	Physics
SAN	OOM	89.83 ± 0.16	94.86 ± 0.10	94.51 ± 0.15	OOM
GraphGPS	OOM	OOM	95.06 ± 0.13	93.93 ± 0.12	OOM
NAGphormer	NA	91.22 ± 0.14	95.49 ± 0.11	95.75 ± 0.09	97.34 ± 0.03
Exphormer	72.44 ± 0.28	91.59 ± 0.31	95.27 ± 0.42	95.77 ± 0.15	97.16 ± 0.13



What if the original graph is too big?



Memory usage

- GPUs only have so much memory available
- Exphormer only adds $\mathcal{O}(N)$ extra edges to the original graph

• But what if the original graph already had too many edges to run on? ogbn-proteins has 132,000 nodes (fine) but almost 40,000,000 edges!

Can easily run out of GPU memory even with just original-graph attention

Which edges matter?

- Attention on graphs tends to be kind of sparse, ish If we could throw away unimportant edges beforehand,
- we could afford to do our optimization on the rest
- But how to tell which edges matter if we can't afford to train?
- Idea: train another network to tell us which edges matter

Attention score estimation

- Idea I: narrow nets are much cheaper in memory than wide nets (lower-dimensional hidden features)
- Idea II: CPU memory is much cheaper than GPU memory

Node characteristics

nodes 🔺	cores ÷	available memory	CPU ¢	storage \$	GPU \$
3	64	4000G or 4096000M	2 x AMD Rome 7502 @ 2.50 GHz 128M cache L3	1 x 960G SSD	-
33	64	2009G or 2057500M	2 x AMD Rome 7532 @ 2.40 GHz 256M cache L3	1 x 960G SSD	-
159	48	498G or 510000M	2 x AMD Milan 7413 @ 2.65 GHz 128M cache L3	1 x SSD of 3.84 TB	4 x NVidia A100SXM4 (40 GB memory), connected via NVLink

Train a narrow network on CPU to tell which edges matter

Attention scores agree across widths

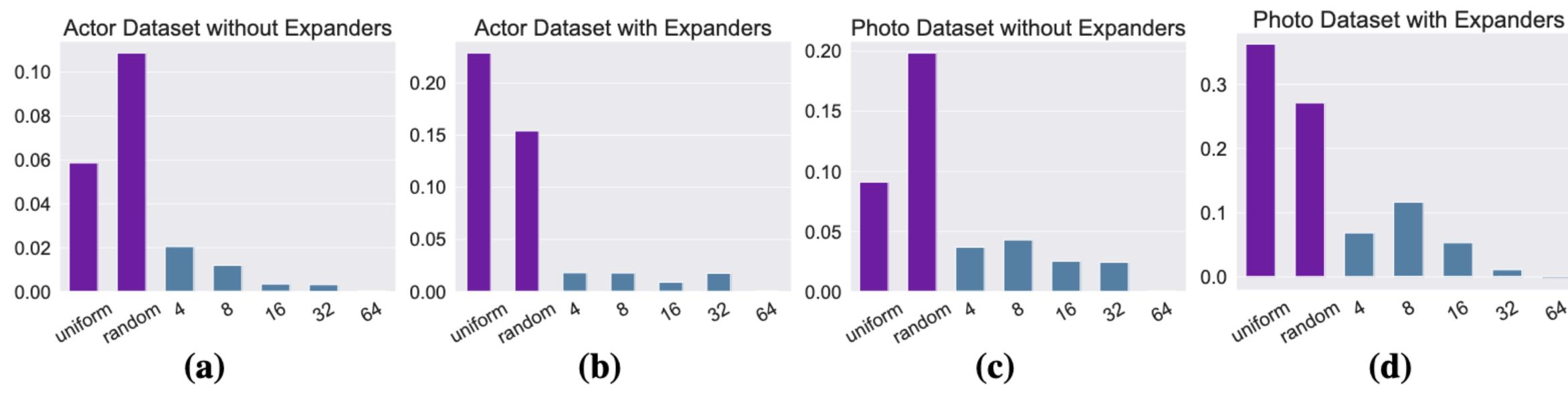
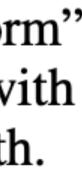


Figure 2: Energy distance between the attention scores of various networks to a network of width 64. "Uniform" refers to the baseline placing uniform scores on each neighbor, while "random" refers to the baseline with uniformly distributed logits. The remaining bars refer to networks trained on the appropriately labeled width.

Caveat I: adding layer normalization to value matrices, to make attention scores more comparable Caveat II: using an annealed temperature schedule to encourage sparser attention





Theory

 Have some partial results about existence of narrow transformers approximating wide ones

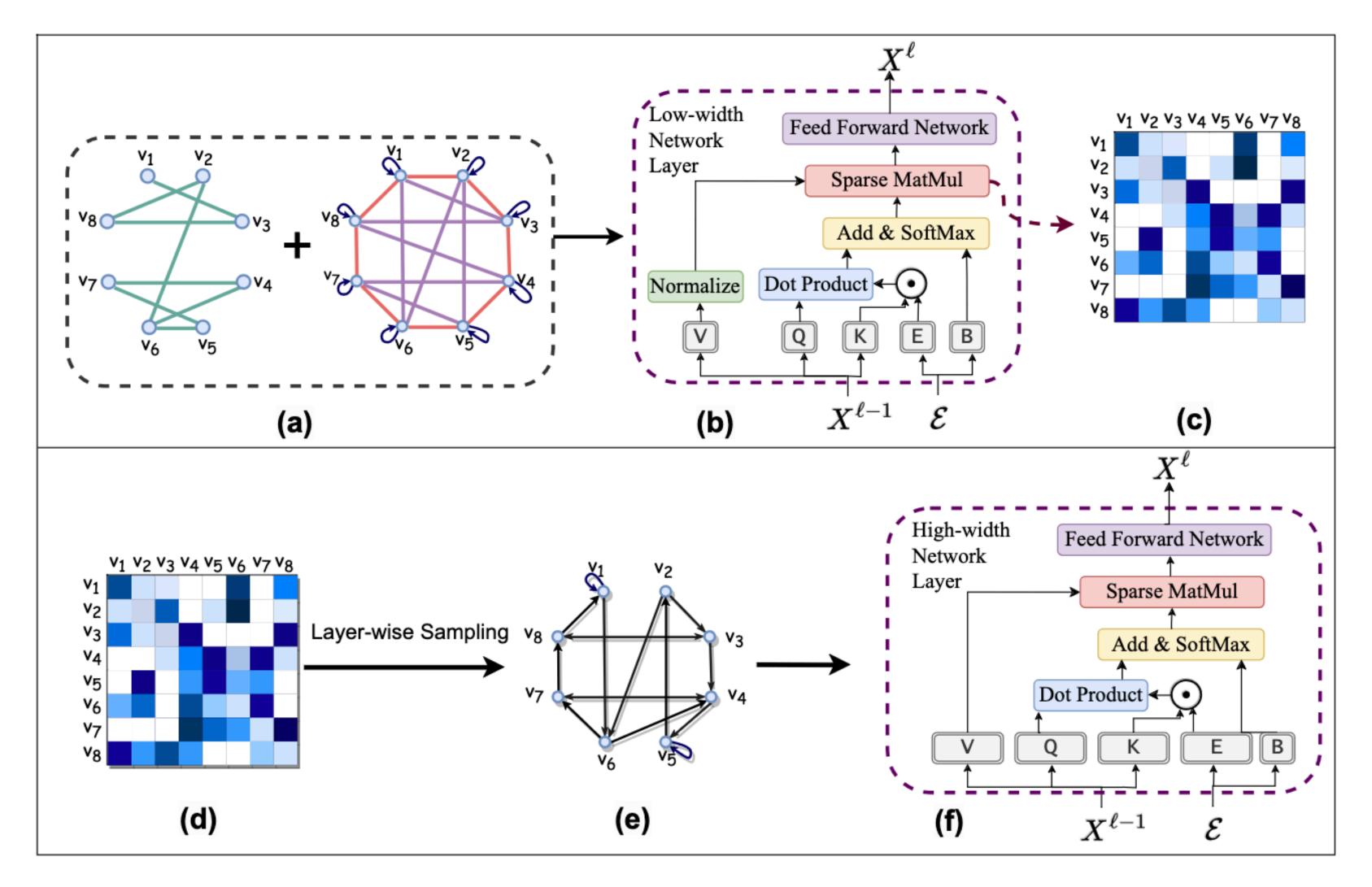
- width
- Following MLP + attention layers...not totally sure

• The first attention layer can be well-approximated with only log(N)



Choosing the important edges

- Once we know attention scores, how do we decide which to keep?
 - Sample k edges for each node proportionally to the attention score
 - Do different samples per graph, and resample in each epoch
- Theory: approximates attention matrix in spectral norm well (extending proof of Achlioptas, Karnin, Liberty, NeurIPS 2013)
- Using reservoir sampling makes it run much faster than default implementation
- Once we have the sparsified graph, run a wide transformer (on GPU) on only that sparsified graph



larger feature dimension, does not normalize V.

Figure 1: Steps of our method. (a) The attention mechanism for the attention score estimator network combines graph edges with an expander graph and self-loops. The expander graphs are constructed by combining a small number of Hamiltonian cycles – here two, in red and in purple – then confirming the spectral gap is large enough. (b) Self-attention layers in the estimator network use this sparse attention mechanism; its self-attention layers normalize V. (c, d) Attention scores are extracted from this network for each layer, and used to sample, in (e), a sparse directed graph, which becomes the attention graph for the final network (f). This network, with a much

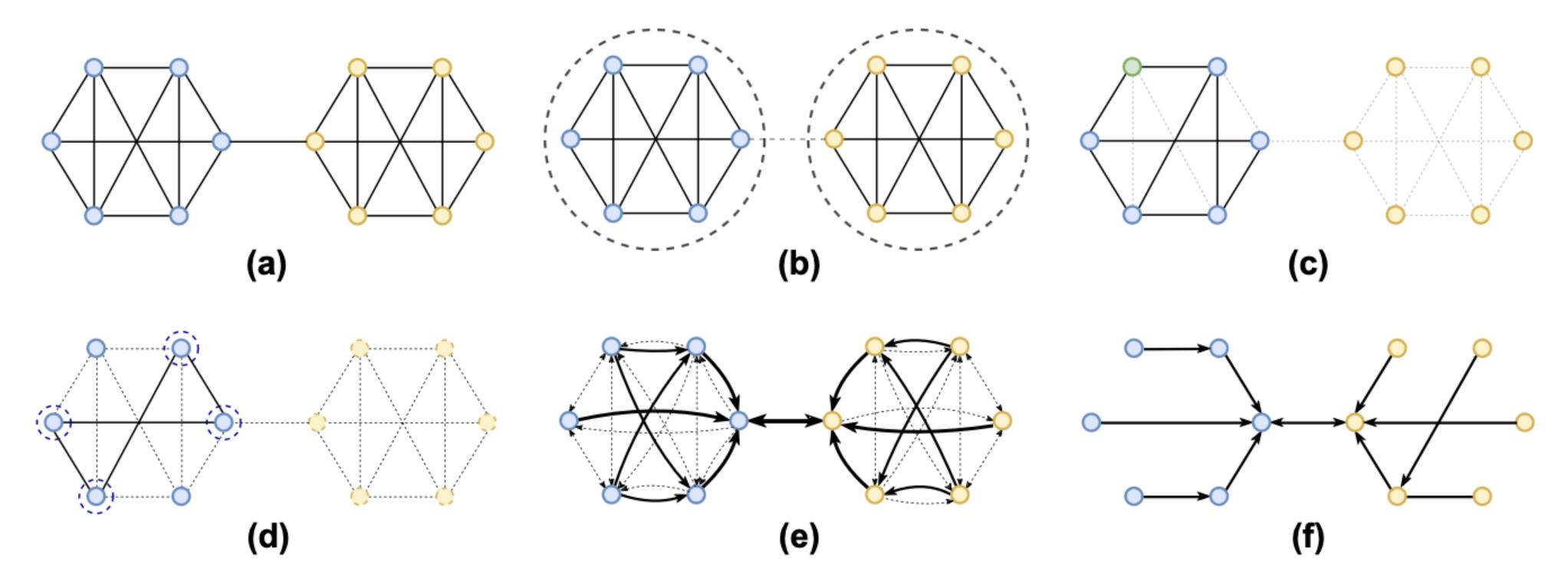
Advantage: regular graph

- Irregularly-shaped matrix multiplications are slow on GPU
- If each node has exactly k neighbours, we instead can do batched matrix multiplication with consistent shapes (much faster!)

Advantage: neighbourhood sampling

- Can do minibatching by
 - Start with some "core nodes"
- Compared to other minibatching strategies on graphs, guarantees that minibatching doesn't bias the process

Include their entire "receptive field" of other nodes that affect them



either of them leads to the correct result.

Figure 4: Figure (a) shows a very simple synthetic graph where each node has a binary classification task of determining whether there exists a node of the opposite color in the graph. This task requires learning long-range dependencies. Figure (b) shows a natural clustering of the graph, this clustering would mean no node can do its task if models are trained only on the clusters. Figure (c) shows a neighbor sampling starting from the green node, where random sampling fails to select the important edge that bridges the between the two different colored nodes. Figure (d) shows a random subset sampling strategy that a subset of nodes are likely to be selected from the same color only, making it impossible to reason about the task. (e) shows attention scores between the nodes if trained with an attention based network. Dashed lines have near zero attention scores, and thicker the lines the attention score is higher. Knowing these attention scores will mean each node with just one directional edge can do the task perfectly. The attention edges are shown in (f). In case two nodes are equally informative selecting

Table 1: Comparison of our model with other GNNs on five homophilic and three heterogeneous datasets (shown in the left and right tables, respectively). The reported metric is ROC-AUC for the Minesweeper and Tolokers datasets, and accuracy for all others.

	/	2							
Model	Computer	Photo	CS	Physics	ogbn-arxiv	Model	Actor	Minesweeper	Toloker
GCN	89.65 ± 0.52	92.70 ± 0.20	92.92 ± 0.12	96.18 ± 0.07	71.74 ± 0.29	GLOGNN	36.4 ± 1.6	51.08 ± 1.23	73.39 ± 1
GRAPHSAGE	91.20 ± 0.29	94.59 ± 0.14	93.91 ± 0.13	96.49 ± 0.06	71.49 ± 0.27	SGC	27.0 ± 0.9	-	-
GAT	90.78 ± 0.13	93.87 ± 0.11	93.61 ± 0.14	96.17 ± 0.08	72.01 ± 0.20	GCN	33.23 ± 1.16	89.75 ± 0.52	83.64 ± 0
GRAPHSAINT	90.22 ± 0.15	91.72 ± 0.13	94.41 ± 0.09	96.43 ± 0.05	68.50 ± 0.23	GRAPHGPS	-	90.63 ± 0.67	83.71 ± 0
NODEFORMER	86.98 ± 0.62	93.46 ± 0.35	95.64 ± 0.22	96.45 ± 0.28	59.90 ± 0.42	NAGPHORMER	-	84.19 ± 0.66	78.32 ± 0
GRAPHGPS	91.19 ± 0.54	95.06 ± 0.13	93.93 ± 0.12	97.12 ± 0.19	70.92 ± 0.04	NodeFormer	36.9 ± 1.0	86.71 ± 0.88	78.10 ± 1
GOAT	90.96 ± 0.90	92.96 ± 1.48	94.21 ± 0.38	96.24 ± 0.24	72.41 ± 0.40	GOAT	-	81.09 ± 1.02	83.11 ± 1
EXPHORMER+GCN	91.59 ± 0.31	95.27 ± 0.42	95.77 ± 0.15	97.16 ± 0.13	72.44 ± 0.28	EXPHORMER+GAT	38.68 ± 0.38	90.74 ± 0.53	83.77 ± 0
EXPHORMER	91.16 ± 0.26	95.36 ± 0.17	95.19 ± 0.26	96.40 ± 0.20	71.27 ± 0.27	EXPHORMER	39.01 ± 0.69	92.26 ± 0.56	83.53 ± 0
Spexphormer	90.93 ± 0.08	95.24 ± 0.12	95.00 ± 0.15	96.70 ± 0.05	70.82 ± 0.24	Spexphormer	38.44 ± 0.42	90.72 ± 0.06	83.15 ± 0

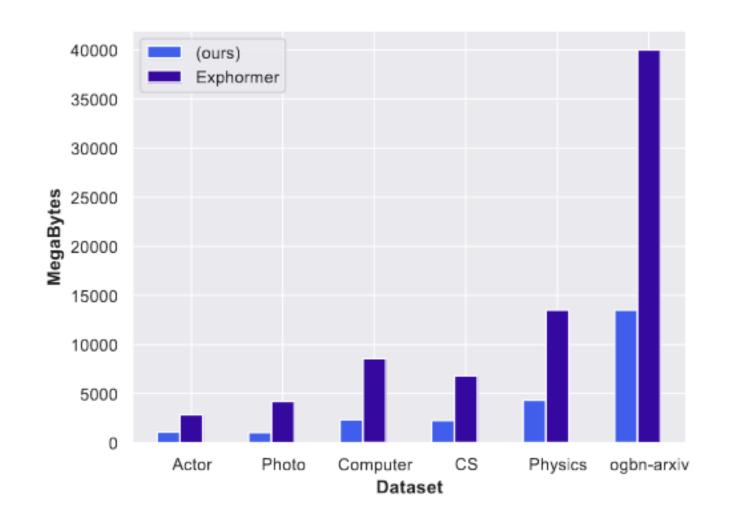
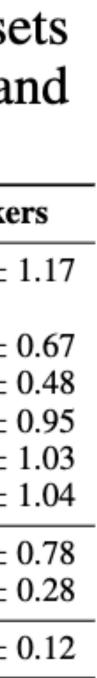


Figure 3: Comparing memory usage of our model with the Exphormer with expander degree 30.



Model	ogbn-proteins	Amazon2M	Pokec
MLP	72.04 ± 0.48	63.46 ± 0.10	60.15 ± 0.03
GCN	72.51 ± 0.35	83.90 ± 0.10	62.31 ± 1.13
SGC	70.31 ± 0.23	81.21 ± 0.12	52.03 ± 0.84
GCN-NSAMPLER	73.51 ± 1.31	83.84 ± 0.42	63.75 ± 0.77
GAT-NSAMPLER	74.63 ± 1.24	85.17 ± 0.32	62.32 ± 0.65
SIGN	71.24 ± 0.46	80.98 ± 0.31	68.01 ± 0.25
NodeFormer	77.45 ± 1.15	87.85 ± 0.24	70.32 ± 0.45
SGFORMER	79.53 ± 0.38	89.09 ± 0.10	73.76 ± 0.24
Spexphormer	$\textbf{80.66} \pm \textbf{0.21}$	$\textbf{90.32} \pm \textbf{0.01}$	$\textbf{74.73} \pm \textbf{0.04}$

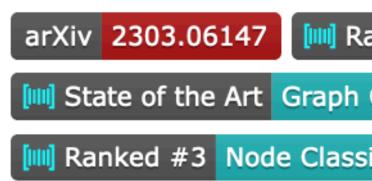
and accuracy for all others.

Table 2: Comparative results on large graph datasets, with ROC-AUC reported for the ogbn-proteins dataset

Thanks!

- Graph transformers: like regular transformers, but on graphs
- Full pairwise attention way too expensive for large graphs

Exphormers: Sparse Transformers for Graphs



- Augment attention graph with expanders + maybe virtual nodes
- Spexphormer

Exphormer

Sparsify the augmented graph with a pilot network

arXiv 2303.06147 [III] Ranked #5 Graph Classification on CIFAR10 100k [III] Ranked #3 Node Classification on COCO-SP State of the Art Graph Classification on MalNet-Tiny Manked #2 Graph Classification on MNIST Ranked #3 Node Classification on PascalVOC-SP [III] Ranked #2 Link Prediction on PCQM-Contact (MRR metric)

