

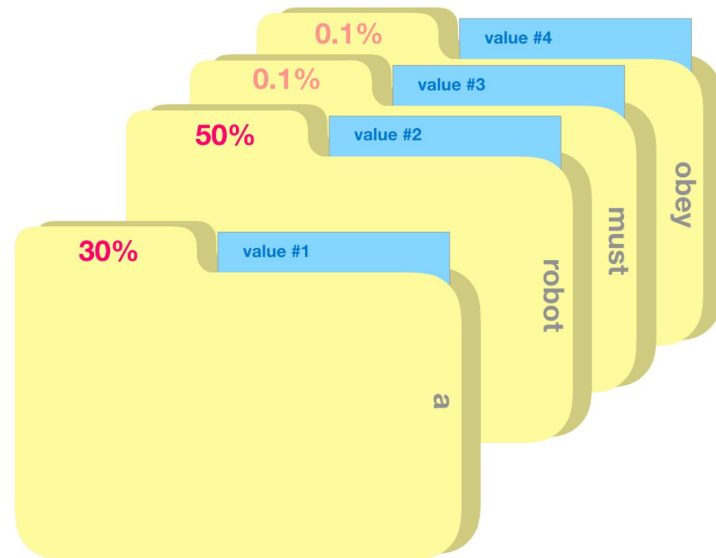
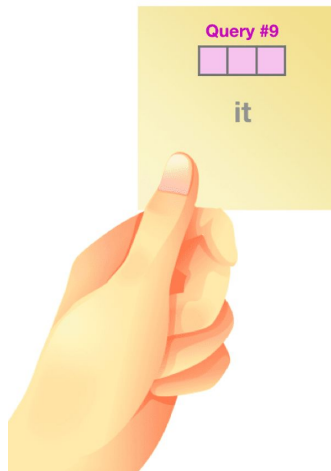
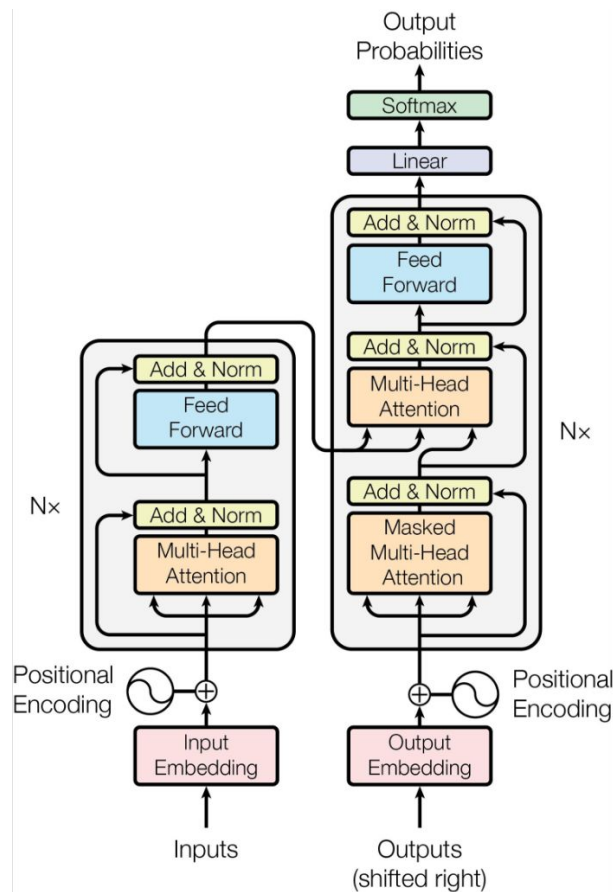
A Generalization of Transformer Networks to Graphs

Vijay Prakash Dwivedi, Xavier Bresson

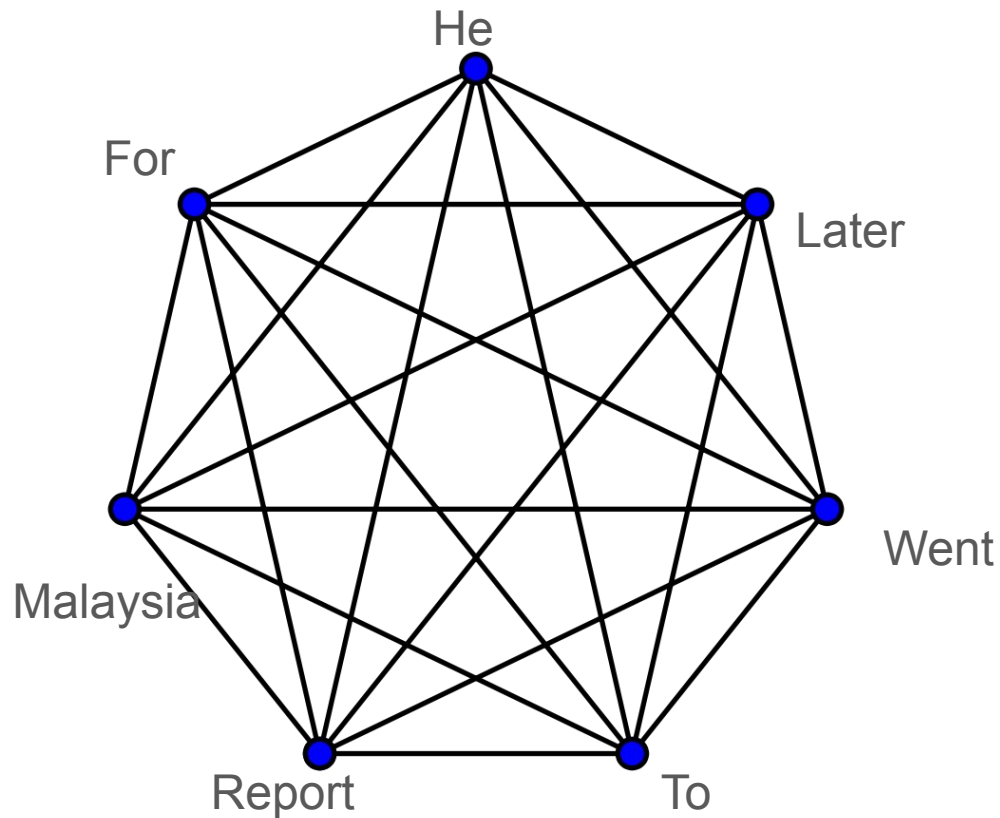
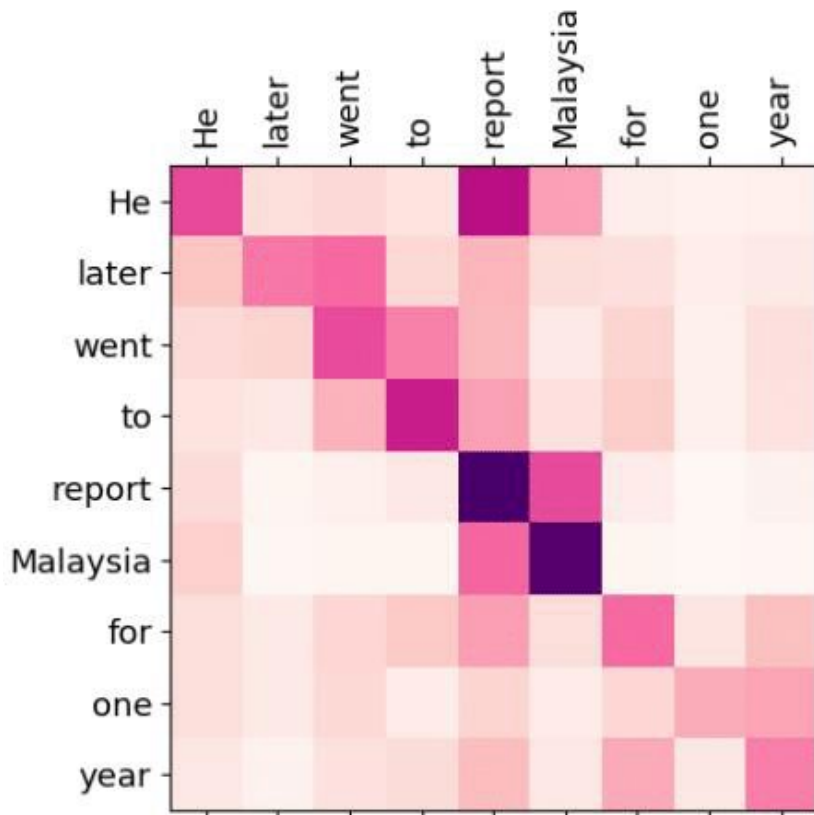
Paper report by

Aleksandr Kariakin, Lev Leontev, Nikita Ivlev

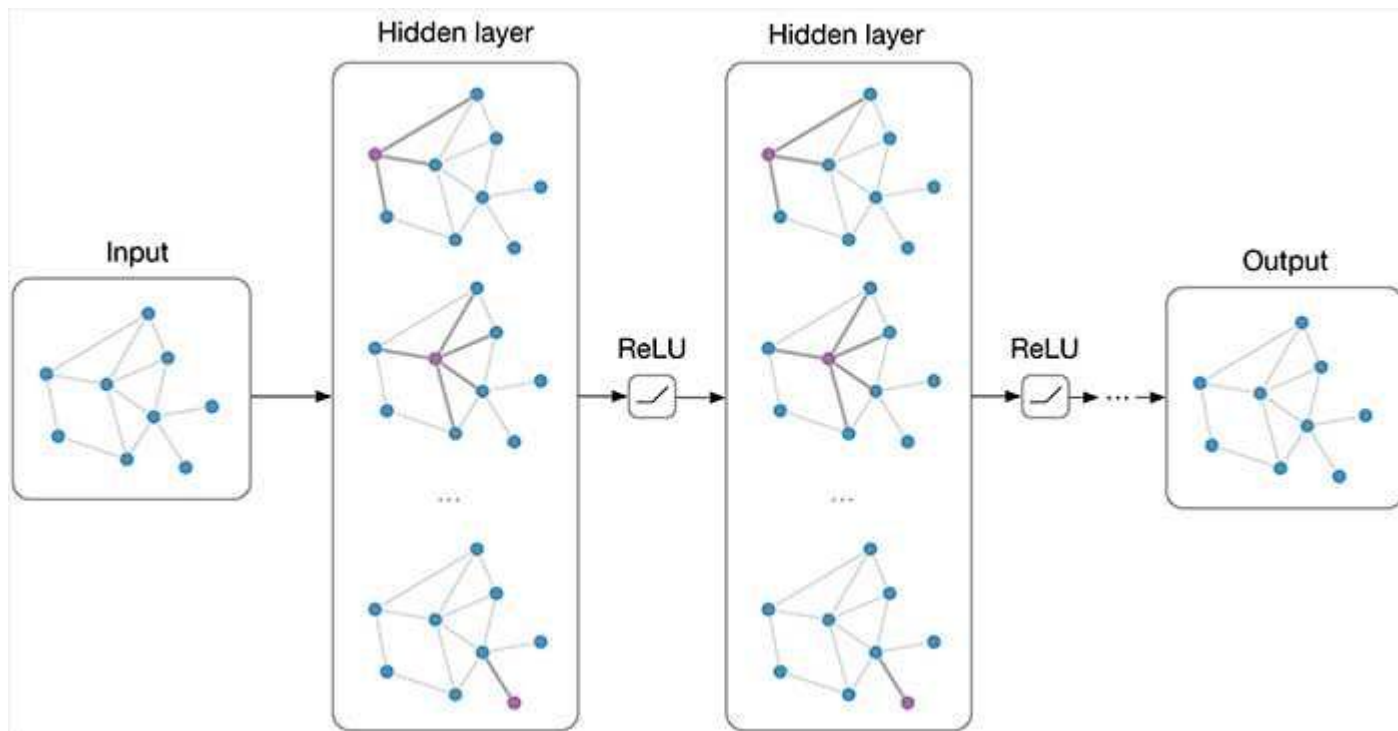
Recall: Transformer, Attention



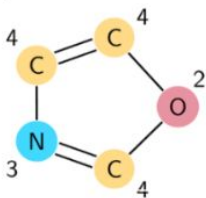
Attention is a complete graph between words



Graph Neural Networks (GNNs)



Merging Graphs and Transformers



Chemistry [1]

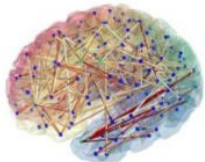
- Learn on molecules and predict chemical properties
- Use in drug repurposing



Simple Particles

Physics [2]

- Learn from interactions of particles in systems
- Accelerate physics research



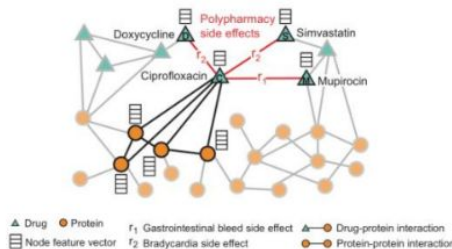
Neuroscience [5]

- Learn functions of brain regions through connectivity
- Accelerate brain-understanding and neuro-disease research



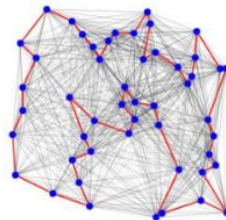
Social networks [3]

- Learn from multi-faceted interactions among users
- Use for commercial and social applications



Medicine [4]

- Learn the effects of multiple drugs on body proteins
- Use for efficient multi-drug medical therapies

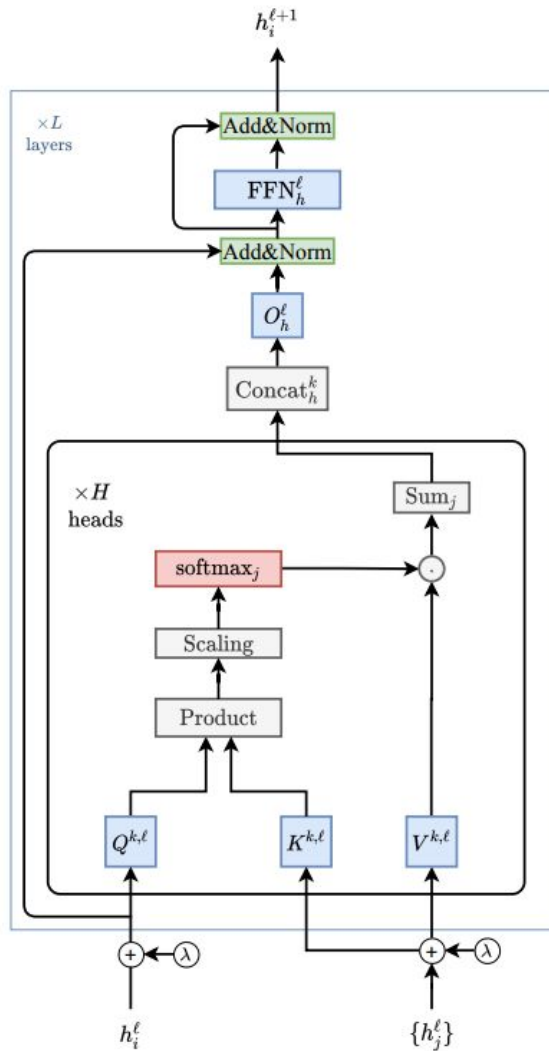


Combinatorial Optimization [6]

- Exploit the fact that most CO problems are rep. as graphs
- Develop better approximated solutions for NP-hard problems

Numerous such examples of graph data.

Proposed architecture

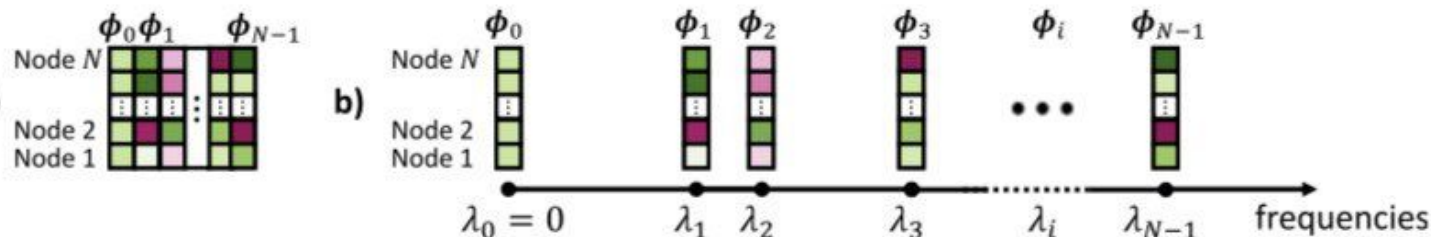


Laplacian Positional Encoding

Eigenvectors are defined via the factorization of the graph Laplacian matrix;

$$\Delta = \mathbf{I} - D^{-1/2} A D^{-1/2} = U^T \Lambda U, \quad (1)$$

where A is the $n \times n$ adjacency matrix, D is the degree matrix, and Λ , U correspond to the eigenvalues and eigenvectors respectively. We use the k smallest non-trivial eigenvectors of a node as its positional encoding and denote by λ_i for node i .



Laplacian Positional Encoding

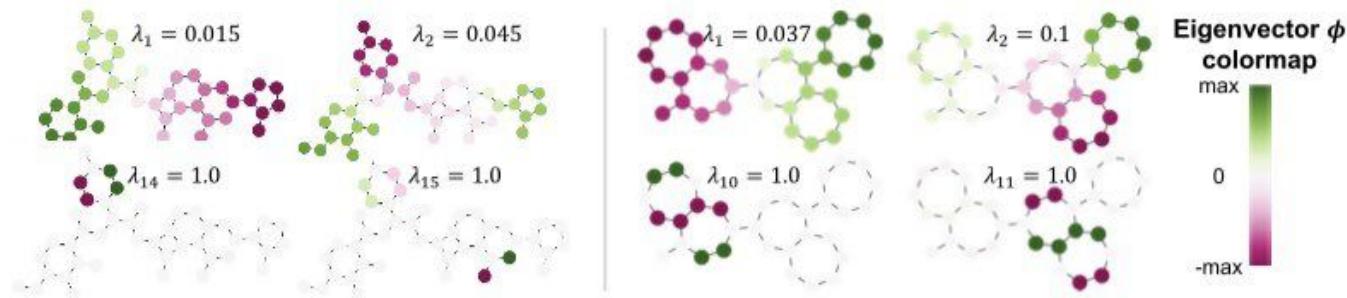
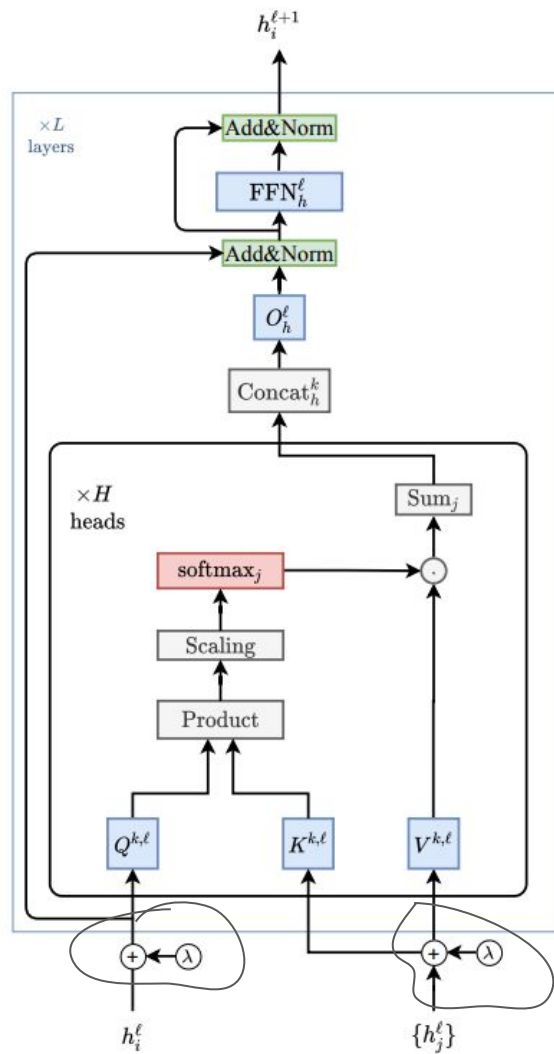


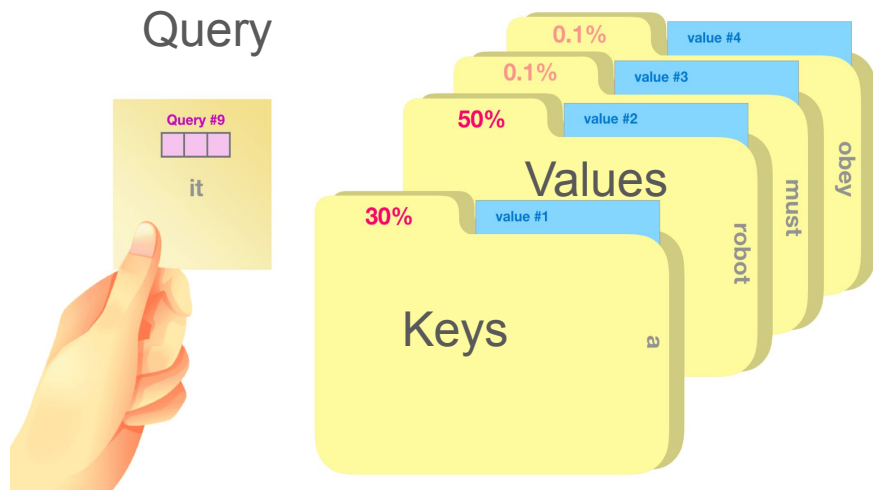
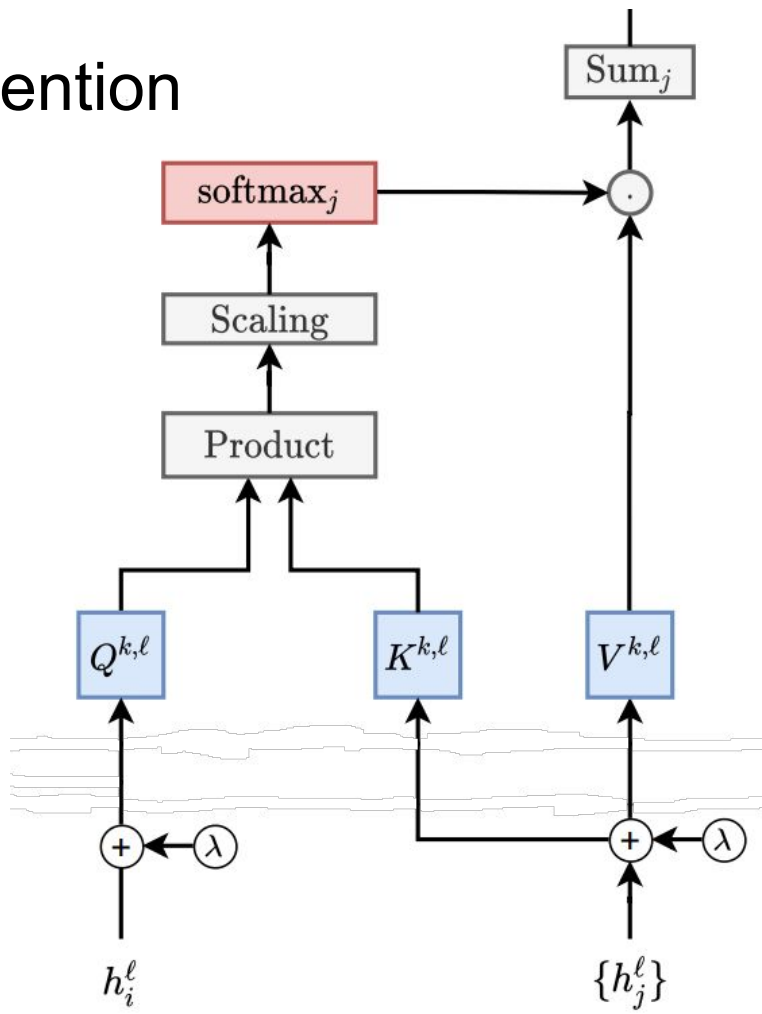
Figure 3: Examples of eigenvalues λ_i and eigenvectors ϕ_i for molecular graphs. The low-frequency eigenvectors ϕ_1, ϕ_2 are spread across the graph, while higher frequencies, such as ϕ_{14}, ϕ_{15} for the left molecule or ϕ_{10}, ϕ_{11} for the right molecule, often resonate in local structures.

Just as the Fourier transform captures the frequency content of a signal, Laplacian eigenvectors capture the structural content of a graph. They help encode distance-aware information, which means that nearby nodes have similar positional features, and farther nodes have dissimilar positional features. Essentially, Laplacian eigenvectors help in understanding the geometrical structure of the graph.

Laplacian Positional Encoding



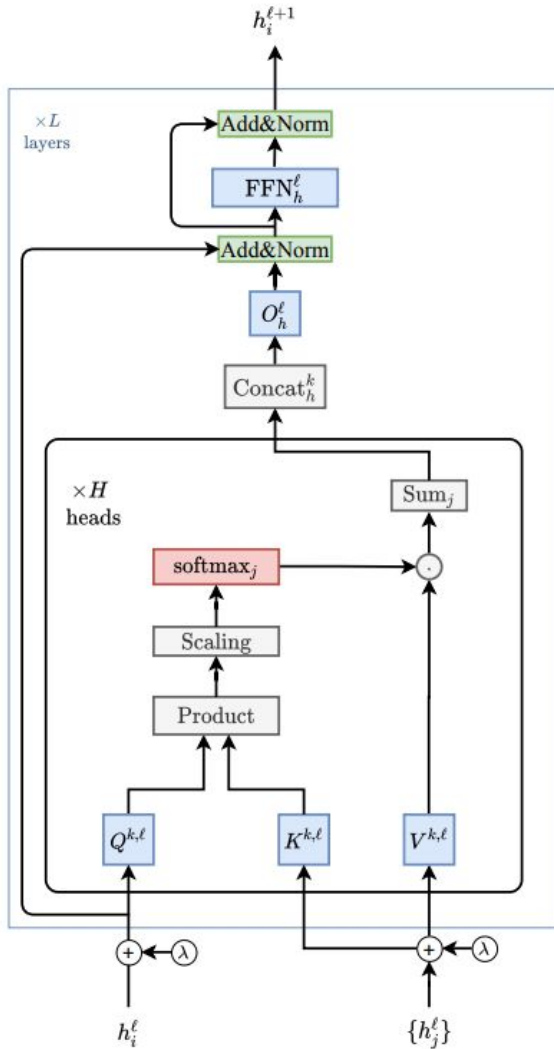
Attention



Attention Vector =
$$\sum_{j \in \mathcal{N}_i} w_{ij}^{k,\ell} V^{k,\ell} h_j^\ell$$

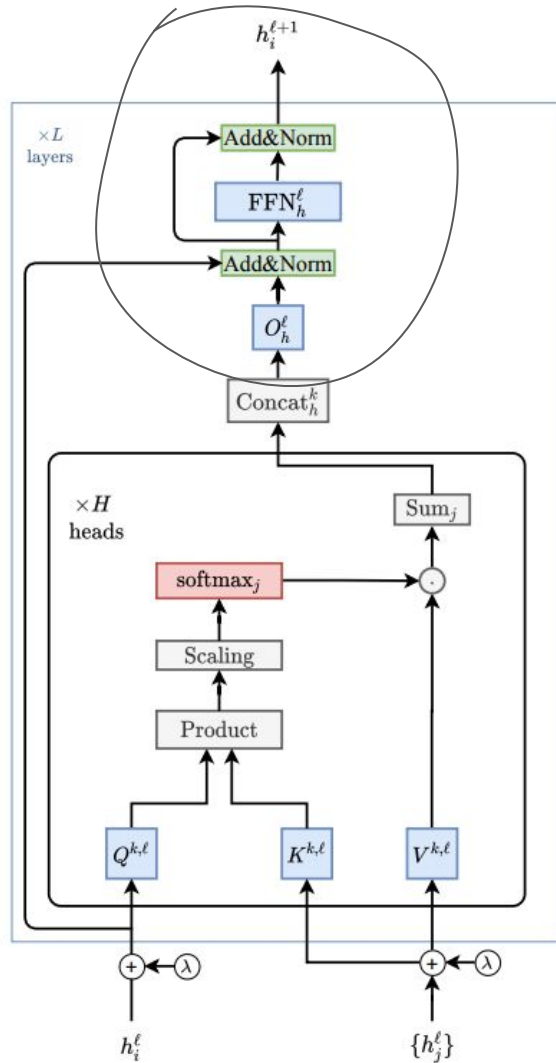
$$w_{ij}^{k,\ell} = \text{softmax}_j \left(\frac{Q^{k,\ell} h_i^\ell \cdot K^{k,\ell} h_j^\ell}{\sqrt{d_k}} \right)$$

Multi-head attention



Feed Forward Network

$$\begin{aligned}\hat{\hat{h}}_i^{\ell+1} &= \text{Norm}\left(h_i^\ell + \hat{h}_i^{\ell+1}\right), \\ \hat{\hat{h}}_i^{\ell+1} &= W_2^\ell \text{ReLU}(W_1^\ell \hat{\hat{h}}_i^{\ell+1}), \\ h_i^{\ell+1} &= \text{Norm}\left(\hat{\hat{h}}_i^{\ell+1} + \hat{\hat{h}}_i^{\ell+1}\right)\end{aligned}$$



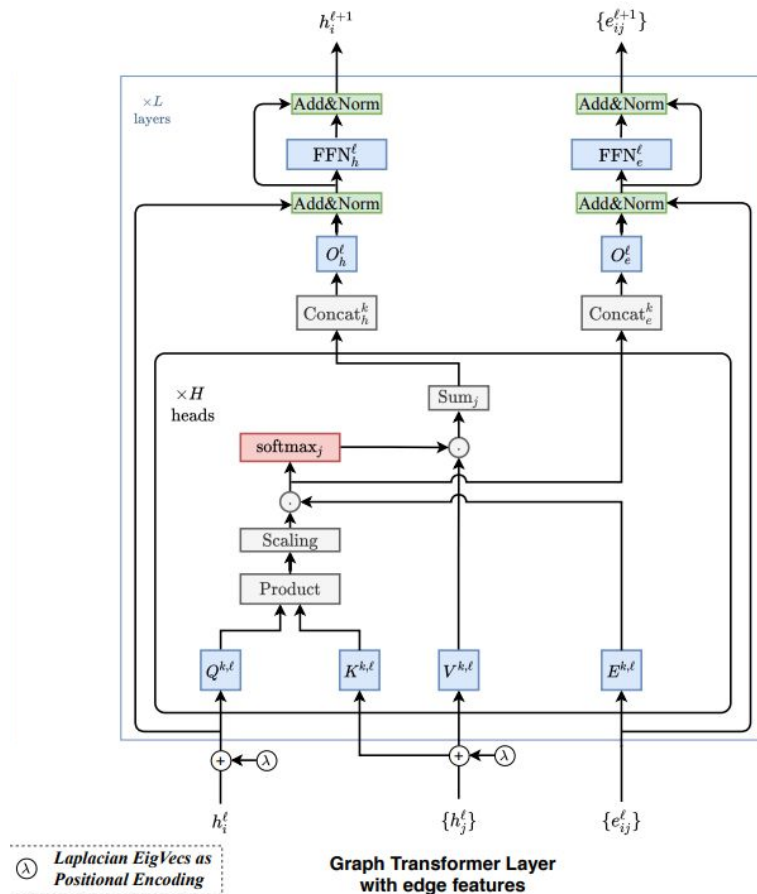
BatchNorm

$$\mu_B = \frac{1}{m} \sum_{i=1}^m x_i \text{ and } \sigma_B^2 = \frac{1}{m} \sum_{i=1}^m (x_i - \mu_B)^2.$$

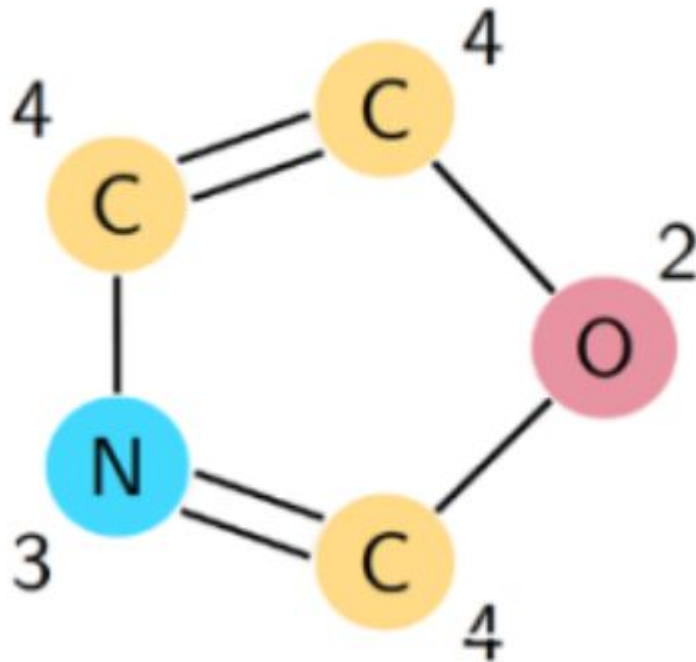
$$\hat{x}_i^{(k)} = \frac{x_i^{(k)} - \mu_B^{(k)}}{\sqrt{(\sigma_B^{(k)})^2 + \epsilon}}$$

What if we have edge features?

- Just multiply the weights in the attention on the edge features! And then softmax
- Update the edge features with the new values



What if we have edge features? Example: molecules



Graph benchmark datasets: ZINC

B

1 2 5 3 4

1 47319 substances subsets endogenous Search within these

ZINC36
Mandelic Acid

ZINC166
Clobutinol

ZINC706
Sanguinarium

ZINC733
Oxedrine

ZINC882
Adenine

ZINC920
Aminobenzoic Acid

ZINC1016
Benzoyl Peroxide, ...

ZINC1070
Bufotenine

ZINC1080
Butylated Hydroxy...

ZINC1082
Trigonelline Hydro...

ZINC1086
Camphoric Ac

ZINC1239
nazine

ZINC1311
Dioscorine

ZINC1360
(+)-Enterolactone

ZINC1392
Ethylparaben

ZINC1507
Gentisic Acid Etha...

ZINC1571
Iotyrosine

ZINC1592
e Hydroc...

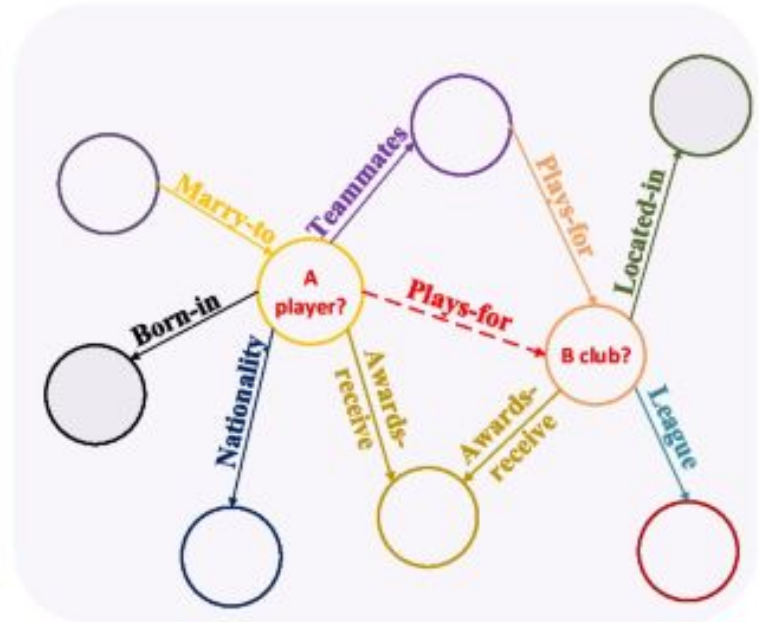
Download All As
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JSON
TXT
MOL2
DB
SDF
SMI
SOLV
DB2

7

Other example of edge features: link prediction

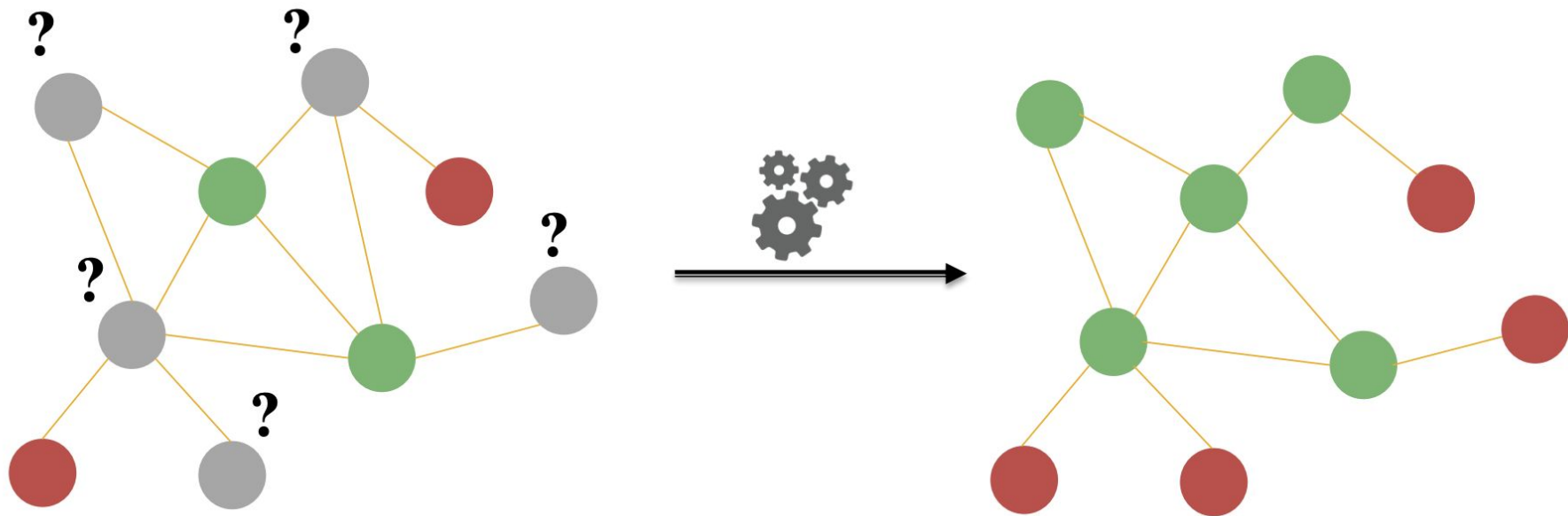


(a) training graph and transductive link prediction



(b) inductive link prediction

Graph benchmark datasets: PATTERN and CLUSTER



Comparison to previous models

Model	ZINC	CLUSTER	PATTERN
GNN BASELINE SCORES from (Dwivedi et al. 2020)			
GCN	0.367 ± 0.011	68.498 ± 0.976	71.892 ± 0.334
GAT	0.384 ± 0.007	70.587 ± 0.447	78.271 ± 0.186
GatedGCN	0.214 ± 0.013	76.082 ± 0.196	86.508 ± 0.085
OUR RESULTS			
GT (Ours)	0.226 ± 0.014	73.169 ± 0.622	84.808 ± 0.068

Comparison to other PEs

Dataset	PE	#Param	Test Perf.±s.d.	Sparse Graph		Epoch/Total
				Train Perf.±s.d.	#Epoch	
Batch Norm: True; Layer Norm: False; $L = 10$						
ZINC	x	588353	0.264±0.008	0.048±0.006	321.50	28.01s/2.52hr
	L	588929	0.226±0.014	0.059±0.011	287.50	27.78s/2.25hr
	W	590721	0.267±0.012	0.059±0.010	263.25	27.04s/2.00hr
CLUSTER	x	523146	72.139±0.405	85.857±0.555	121.75	200.85s/6.88hr
	L	524026	73.169±0.622	86.585±0.905	126.50	201.06s/7.20hr
	W	531146	70.790±0.537	86.829±0.745	119.00	196.41s/6.69hr
PATTERN	x	522742	83.949±0.303	83.864±0.489	236.50	299.54s/19.71hr
	L	522982	84.808±0.068	86.559±0.116	145.25	309.95s/12.67hr
	W	530742	75.489±0.216	97.028±0.104	109.25	310.11s/9.73hr

Analysis of GraphTransformer (GT) using different PE schemes. Notations x: No PE; L: LapPE (ours); W: WLPE (Zhang et al. 2020). Bold: the best performing model for each dataset.

Thank you for your *attention*

