

Graph Neural Networks

Al-guided Protein Science





Limitations of Shallow Embeddings

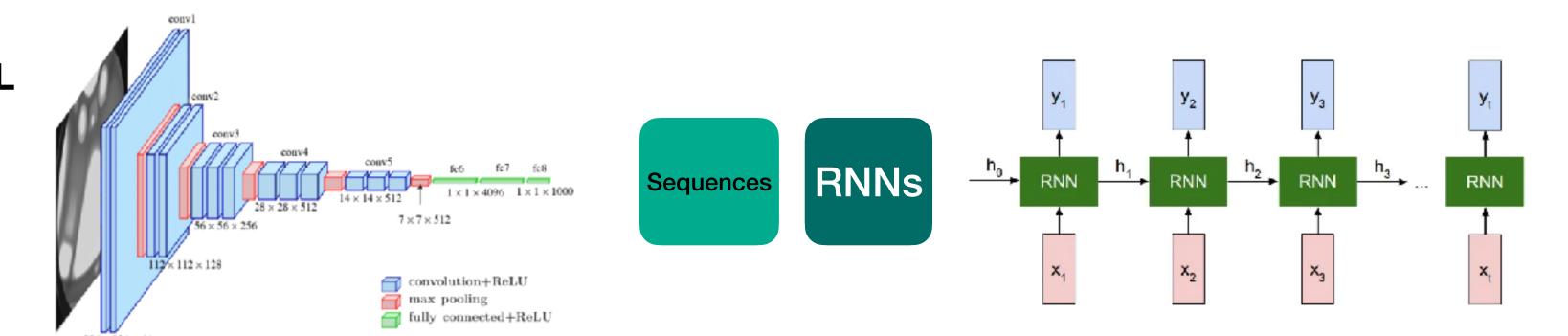
- Complexity grows with the number of nodes (O(|V|)) problem with large graphs
- They do not use node features
- They are **transductive** can only generate embeddings for nodes present during the training phase (no generalizable to unseen nodes)

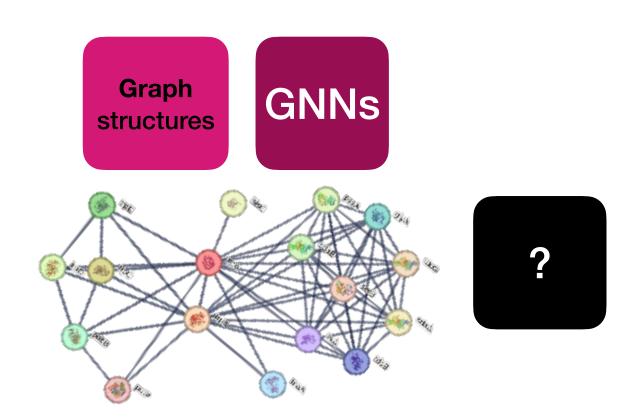
More Complex Encoder Models

- Graph Neural Networks a deep learning framework for representing graph data
 - Incorporate both local and global context
 - Allows node features (e.g., expression)
 - Inductive
- Different to traditional DL



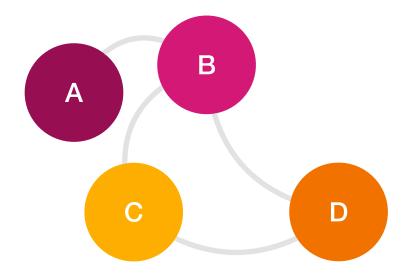






Using the Adjacency Matrix

- Why not using the Adjacency Matrix A?
 - Assumptions for DL
 - Permutation invariance: same output regardless of the order of its input.
 - Permutation equivariance: the output reflects the permutation applied to the input.
 - A has an arbitrary order of nodes



	Α	В	С	D
Α	0	1	0	0
В	1	0	1	1
O	0	1	0	1
D	0	1	1	0

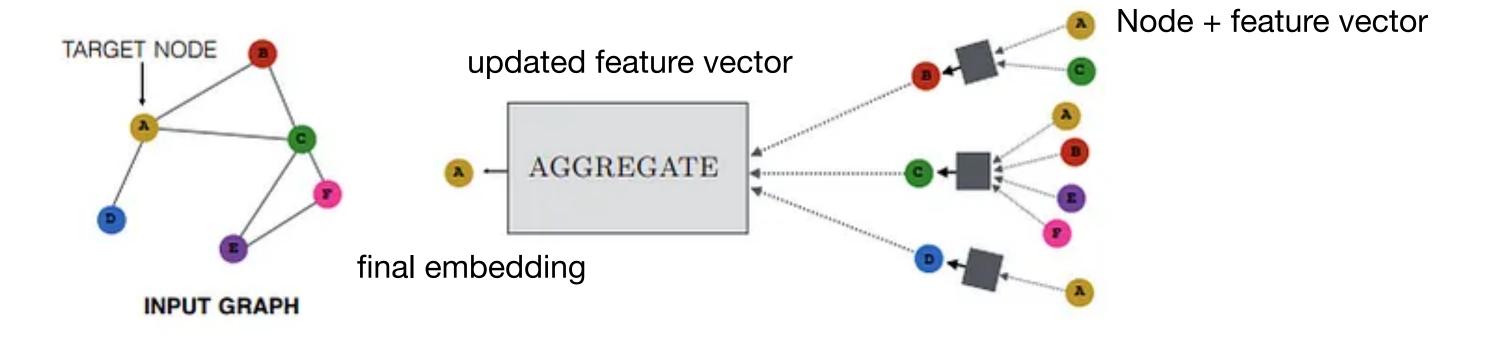
	В	Α	D	С
Α	1	0	0	0
В	0	1	1	1
С	1	0	1	0
D	1	0	0	1

Neural Message Passing

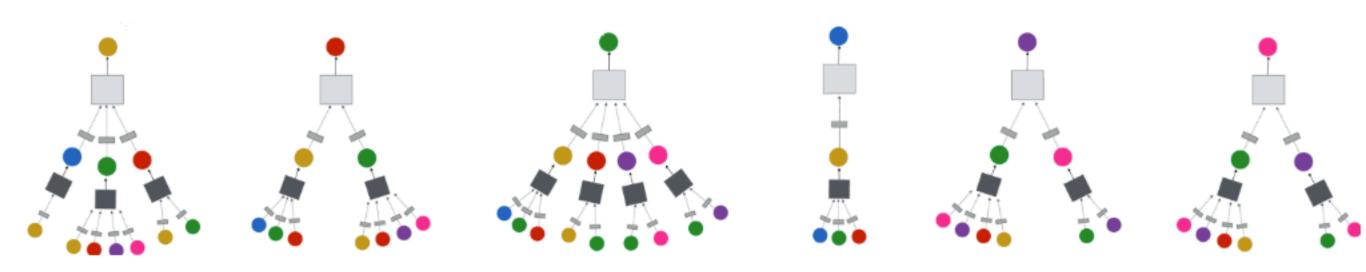
- Enables information exchange and aggregation among nodes capturing the dependencies and interactions among them
- The information collected from neighbours is used to update the representation of the node

At each iteration (layer) of the message passing in a GNN, a hidden embedding $h_u^{(k)}$ is updated for each node u based on information gathered from its graph neighbourhood N(u).

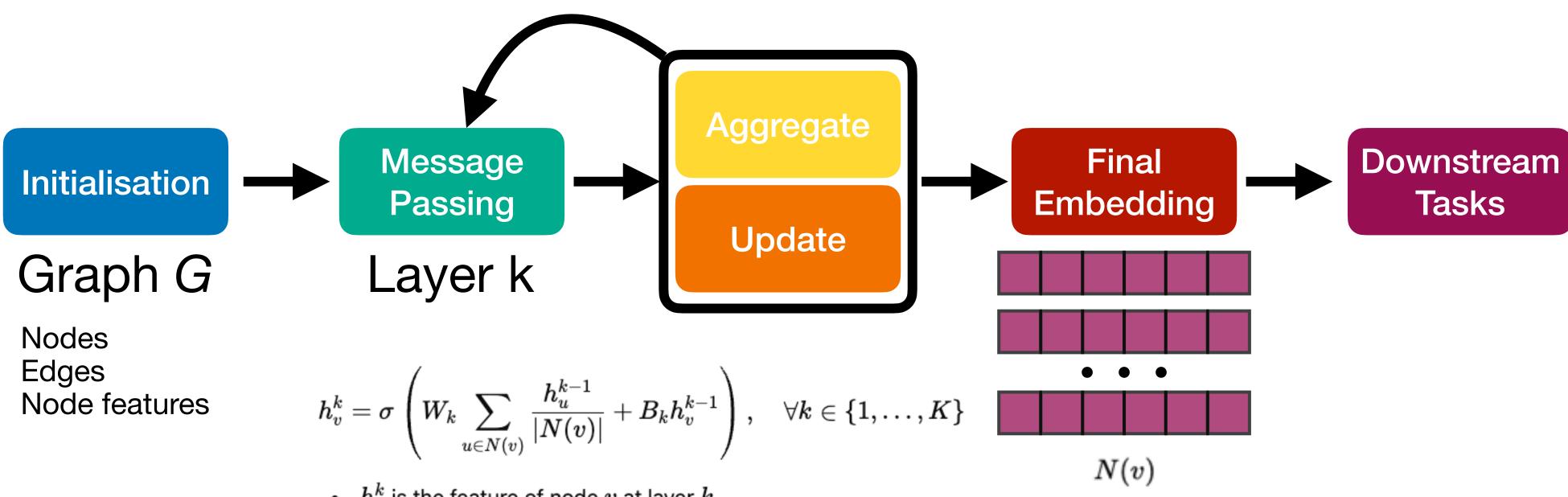
As these iterations progress, each node embedding contains more and more global information



Each node defines a computational graph based on its neighbours



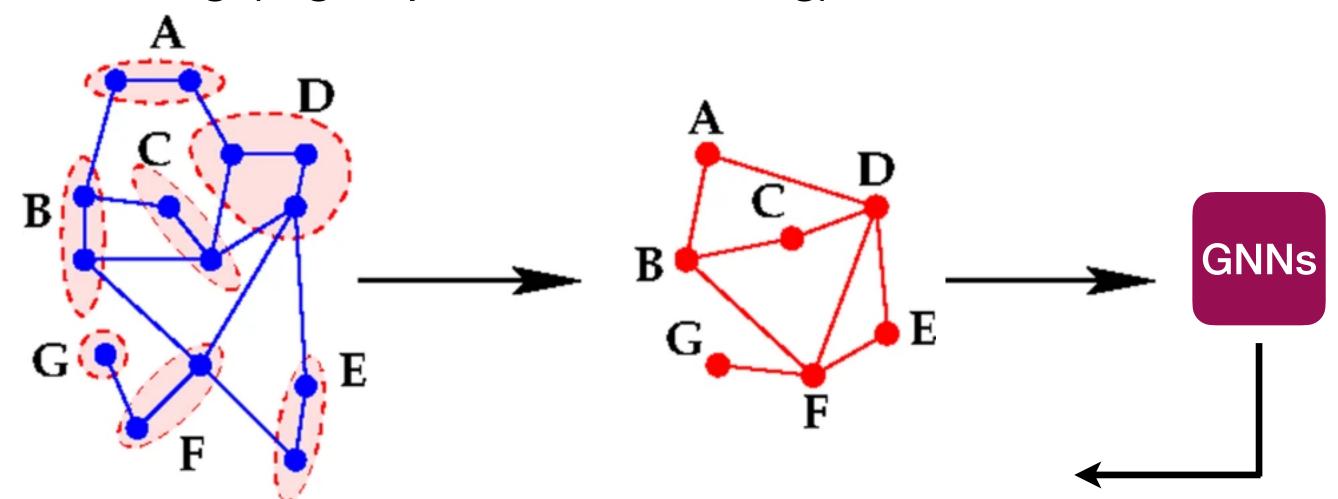
GNN Workflow



- h_v^k is the feature of node v at layer k,
- N(v) is the set of neighbors of node v,
- h_u^{k-1} is the feature of node u at layer k-1,
- |N(v)| is the number of neighbors of node v,
- W_k and B_k are learnable weight matrices and bias at layer k,
- σ is a non-linear activation function (e.g., ReLU or Sigmoid).

Graph Pooling

- Graph-level tasks Define a pooling function that maps a set of node embeddings to an
 embedding representing the full graph.
- Strategies:
 - Take the sum or mean of the node embeddings
 - Combination of LSTMs and attention to pool the node embeddings
 - Clustering or coarsening (e.g., spectral clustering)



Variations of GNNs

- Graph Convolutional Networks (GCN): A popular form of GNN where the aggregation step is similar to a convolution operation on graphs.
- Graph Attention Networks (GAT): Incorporates attention mechanisms, allowing each node to weight the importance of its neighbours differently.
- **GraphSAGE**: A GNN variant that samples a fixed-size neighbourhood for each node to improve scalability for large graphs.

Model	Best for	
GNN	General-purpose graph tasks, node/graph classification, link prediction	
GCN	Semi-supervised node classification, graph classification	
GAT	Tasks where neighbour importance varies (e.g., node classification with varying relevance of neighbours)	
GraphSAGE	Scalable learning on large graphs, node classification, link prediction	

Generative Models

- Variational Graph Autoencoders (VGAE) an extension of Variational Autoencoders (VAEs) for graphs. It uses a GNN encoder and a decoder that reconstructs the graph structure (adjacency matrix or edge probabilities) using the latent variables.
- Graph Convolutional GAN (GraphGAN) a generative adversarial network (GAN) designed to generate graph-structured data. It uses a generator to create graph structures and a discriminator to differentiate between real and fake graphs.
- Graph Recurrent Neural Networks (GRNN) GRNNs apply RNNs to graphs, generating node sequences or edge sequences iteratively. They can be used for generating dynamic graphs, where nodes and edges evolve over time.

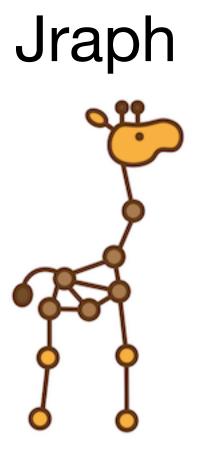
GNN Python Libraries





https://www.dgl.ai/





https://github.com/google-deepmind/jraph

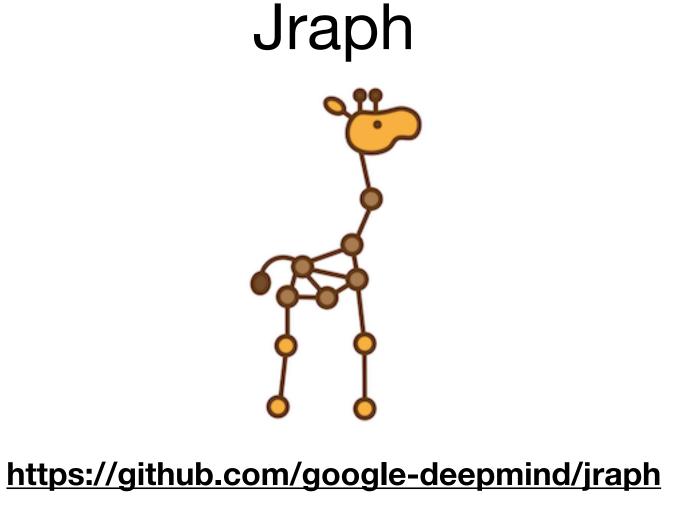
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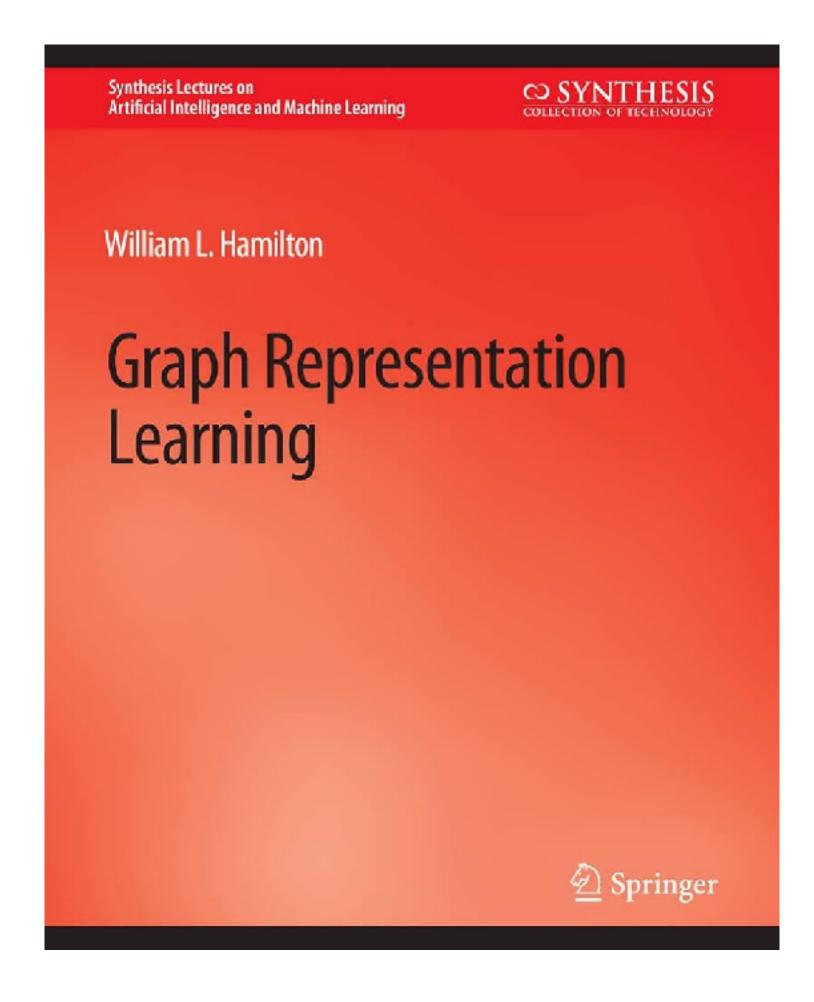


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Acknowledgements

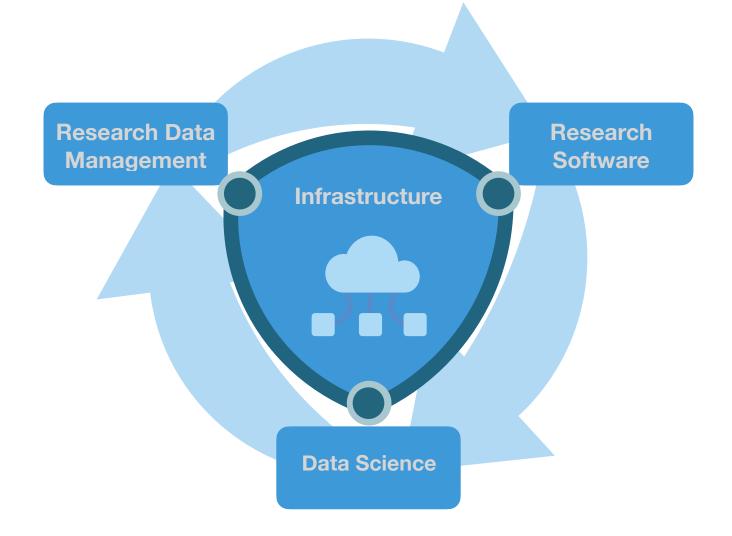


Thank you

Multi-omics Network Analytics Research Group



Informatics Platform









albsad@dtu.dk



@albsantosdel



https://github.com/Multiomics-Analytics-Group



https://multiomics-analytics-group.github.io/

