Lecture 13 Graph Neural Networks

cscis-89 Deep Learning, Fall 2024 Zoran B. Djordjević & Rahul Joglekar

References

- "Graph Representation Learning" by William L. Hamilton, Morgan & Claypool 2020, book available at: https://www.cs.mcgill.ca/~wlh/grl_book/files/GRL_Book.pdf
- "Geometric Deep Learning: Grids, Groups, Graphs, Geodesics, and Gauges" by Michael M. Bronstein, Joan Bruna, Taco Cohen, Petar Veličković, https://arxiv.org/abs/2104.13478
- Bronstein, Michael M. et al. "Geometric deep learning: going beyond Euclidean data." IEEE Signal Processing Magazine 34.4 (2017): 18-42."
 https://arxiv.org/abs/1611.08097
- Liu, Zhiyuan; Zhou, Jie, Introduction to Graph Neural Networks, Morgan & Claypool, 2020.
- The best resource for learning GNNs is the course:
 CS224W: Machine Learning with Graphs, Jure Leskovec, Stanford University
 http://cs224w.stanford.edu
 - These notes borrow heavily from cs224W materials.
- For Python network aware software look for Networkx package https://networkx.org/documentation/stable/tutorial.html
- An excellent book on networks: "Network Science" by Alberto-Laszlo Barabas.
 Available in online version: http://networksciencebook.com/

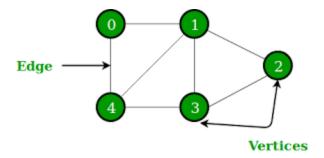
Networks and/or Graphs

Graphs

- Graphs are data structures and a universal language for describing complex systems. In the most general terms, a graph is a collection of objects (i.e., nodes), along with a set of interactions (i.e., edges) between pairs of those objects.
- Graphs are a general language for describing and analyzing entities with relations/interactions.
- We can find graphs (network structures) in nature (molecules), society (social networks), technology (the internet), and everyday settings (roadmaps).
- For example, to encode a social network as a graph we use nodes to represent individuals and use edges to represent the friendship or some other relationship between the individuals.
- In the biological domain, we could use the nodes to represent proteins, and use the edges to represent various biological interactions between proteins.
- To perform machine learning on graphs, we need a specialized form of the neural networks, the graph neural network or GNN.

Graphs vs Grid-like Structures

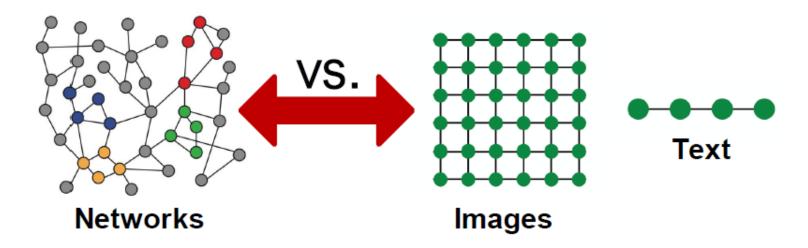
- Graphs or networks are data structures whose key elements are relationships. Graphs
 are often represented as images or diagrams, with edges representing relationships
 and nodes, or vertices representing entities. The diagram below is one representation
 of a graph. On subsequent slides we will see that graphs could be represented by
 matrices and tensors.
- In graphs or networks, the relationships are the first-class citizens.
- In more traditional learning and analytics approaches, we use grid-like structure such as the relational database tables, Pandas DataFrames, or Excel spreadsheets.
- In non-network approach, relationships play much less pronounced role than in the networks or graphs



• In above diagram, circles or nodes (vertices) represent entities. Lines or edges represent relationships. The numbers could be used to indicate node IDs. Nodes could be named and could have many attributes. Edges could be similarly labeled by edge IDs or names. Edges (relationships) could have different attributes as well.

What is different in Graphs

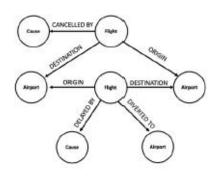
- Networks and graphs could be complex.
- Graphs could have arbitrary size and topological structure without locale spatial uniformity like grids.



- Network or graphs have no fixed node ordering or reference point
- Graphs are often dynamic and have multimodal features
- One can safely argue that images and text are also graphs, just with some additional structural rules or symmetries. Typically, when dealing with such "simplified or ordered" graphs, we insist on relaying on those extra rules and symmetries.

Application of Graphs

Applications of Graphs



Event Graphs

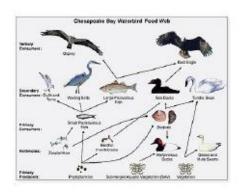


Image credit: Wikipedia

Food Webs



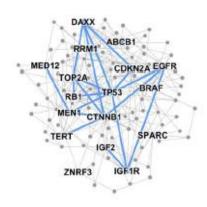
Image credit: SalientNetworks

Computer Networks



Image credit: Pinterest

Particle Networks



Disease Pathways



Image credit: visitlondon.com

Underground Networks

Applications of Graphs





Image credit: Medium

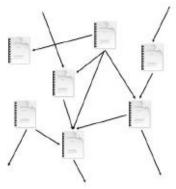
Image credit: Science

Image credit: Lumen Learning

Social Networks



Economic Networks Communication Networks



Citation Networks



Image credit: Missoula Current News

Internet

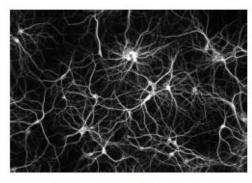
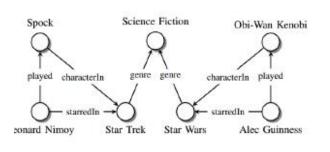
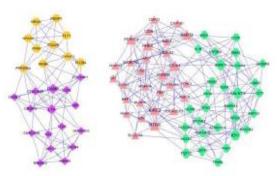


Image credit: The Conversation

Networks of Neurons

Applications of Graphs





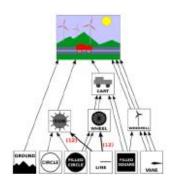


Image credit: Maximilian Nickel et al

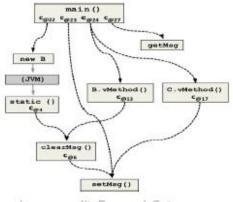
Image credit: ese wustl.edu

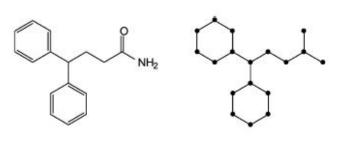
Image credit: math.hws.edu

Knowledge Graphs

Regulatory Networks

Scene Graphs





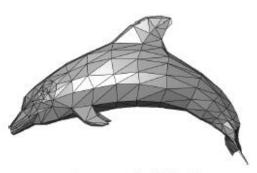


Image credit: ResearchGate

Image credit: MDPI

Image credit: Wikipedia

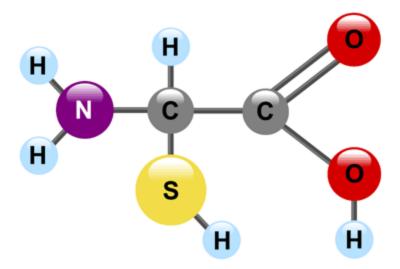
Code Graphs

Molecules

3D Shapes

Molecular Fingerprints and Interfaces

In chemistry and molecular sciences, a prominent problem is how to represent
molecules in a general application-agnostic way, and inferring possible interfaces
between organic molecules, such as proteins. Some molecule representation bear
resemblance to a graph structure, consisting of nodes (atoms) and edges (atomic
bonds).



• Applying Graph Convolutional Networks to structure of molecules is a nascent field that promises to outperform traditional 'fingerprint' methods, which involve the creation of features by domain experts to capture molecular properties.

What could we do with graph data & GNNs

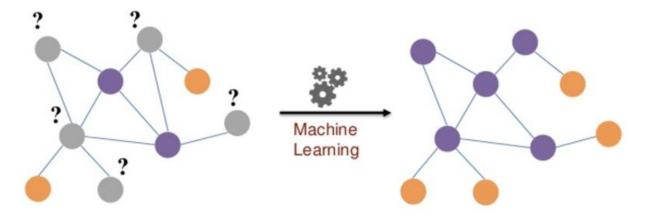
- Assuming one had access to the graph data and it is of sufficient quantity and quality, what could a GNN extract from such data?
- Historical records are not always complete. We could use GNNs to fill in the missing information.
- GNNs could perform node classification to predict node attributes. For example,
 on the list of Titanic passengers, we could predict missing passenger information.
 For example, if the citizenship information was missing from some of the
 passenger data, we could use node classification to uncover the citizenship labels.
- GNNs could perform edge prediction (or link prediction) to uncover hidden or missing links between nodes. In the Titanic case, we could use it to find nonobvious relationships between the passengers, like extended family relationships (cousins on the ship), and business relationships.
- **Graph-level tasks like classification and regression** involve predicting the label or a quantity associated with the entire graph.
- There are also unsupervised tasks that involve GNNs. These involve Graph Auto Encoders that embed graphs and do the opposite process of generating a graph from an embedding. There is also a class of models that use adversarial methods to generate graphs.

Prediction Tasks

- A slew of graph algorithms and analytical methods has existed for decades, for a variety of use cases. Many of these methods are quite powerful, are used at scale, and are very successful.
- PageRank, the first algorithm used by Google to order web search results, is
 probably the most famous example of a graph algorithm used to great success by
 an enterprise.
- Applications where graph neural networks shine are problems that require
 predictive models. We either use the GNN to train such a model, or we use it to
 produce a representation of the graph data that can be used in a downstream
 model.

Node-Level vs Edge-Level tasks

 Nodes in graphs often have attributes and labels. Node classification is the nodelevel task analog to the traditional machine learning classification: given a graph with several node classes, predict the class of an unlabeled node.



- For example, we want to find out what are the true colors of gray nodes?
- Edge-level tasks are very similar to node-level tasks. Link classification involves
 predicting a link's label in a supervised or semi-supervised way. Edge Prediction is
 inferring a link between nodes, where one may not be provided in the graph under
 study.

Types of Graphs and Terminology

Key Terms

- **Graphs** are formally defined by a set of vertices and a set of edges between these vertices: G = G (V, E). Fundamentally, graphs are just a way to encode data. Every property of a graph represents some real element, or concept. Graphs can be used to represent complex concepts.
- **Vertices** represent items, entities, or objects, which have quantifiable attributes and relationships to other items, entities, or objects. We refer to a set $\{vi\}$ of vertices as V. In a graph, not all vertices have to be homogenous with the same set of properties.
- Edges represent and characterize relationships between items, entities, or objects. Formally, a single edge can be defined with respect to two vertices. We label the set $\{e_{ij}\}$ of edges as E. e_{ij} is a single edge between the i-th and j-th vertices. There could be more edges connecting the same pair of vertices.
- Neighborhoods are subgraphs within a graph and represent distinct groups of vertices and edges. Most commonly, the neighborhood N_{v_i} is centered around v_i , and contains v_i itself, its adjoining edges (e_{ij}) , and the vertices that are directly connected to node (vertex) v_i . Neighborhoods can be iteratively grown from a single vertex by considering the vertices attached (via edges) to the current neighborhood. Note that a neighborhood can be defined subject to certain vertex and edge feature criteria (i.e., all vertices within 2 hops of the central vertex, rather than 1 hop).
- **Features** are quantifiable attributes which characterize a phenomenon that is under study. In the graph domain, features characterize vertices and edges. In social networks, we might have features for each person (vertex) specifying the person's age, popularity, and social media usage. Similarly, we might have a feature for each relationship (edge) which quantifies how well two people know each other, or the type of relationship they have (familial, collegial, etc.). In practice, there might be many different features for each vertex or edge, and they are usually represented by numeric feature vectors referred to as F_{v_i} and $F_{e_{ij}}$, respectively.

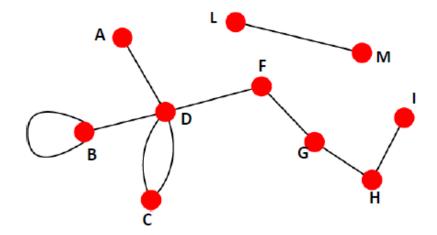
Undirected vs. Directed Graphs

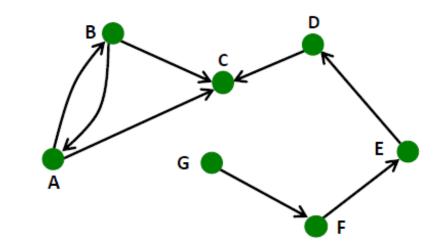
Undirected graphs

- Links: undirected (symmetrical, reciprocal)
- Examples:
 - Collaborations
 - Friendship on Facebook
- Node degree, k_i:the number of edges adjacent to node i. e. g., k_i=4
- Avg. degree: $\bar{k} = \frac{1}{N} \sum_{i=1}^{N} k_i$

Directed graphs:

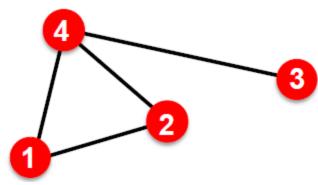
- Links: directed (arcs)
- Examples:
 - Phone calls
 - Followings on Twitter
- In directed networks we define an in-degree and out-degree.
- The (total) degree of a node is the sum of in- and out-degrees.





Representing Graphs: Adjacency Matrix

Undirected Graph



Directed graph

3

- $A_{ij} = 1$ if there is a link from node i to node j
- $A_{ij} = \mathbf{0}$ otherwise

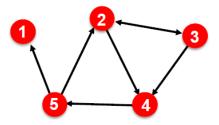
$$\bullet \quad A_{undir} = \begin{pmatrix} 0 & 1 & 0 & 1 \\ 1 & 0 & 0 & 1 \\ 0 & 0 & 0 & 1 \\ 1 & 1 & 1 & 0 \end{pmatrix}$$

$$A_{dir} = \begin{pmatrix} 0 & 0 & 0 & 1 \\ 1 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 \\ 0 & 1 & 1 & 0 \end{pmatrix}$$

- The adjacency matrix for an undirected graph A_{undir} is a symmetric matrix.
- The adjacency matrix for a directed graph A_{dir} is not symmetric.
- Coefficients in the adjacency matrix do not have to be 0-s and 1-s. They could represent weights and be any rational numbers.
- Adjacency matrixes are sparse. Usually, they have very few 1-s and many 0-s.

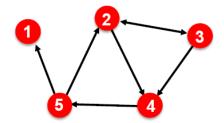
Representing Graphs: Lists of Edges and Adjacency Lists

Represent graph as a list of edges:



- (2, 3)
- (2, 4)
- (3, 2)
- (3, 4)
- (4, 5)
- (5, 2)
- (5, 1)

- Represent graph as an adjacency list:
- Easier to work with if network is Large and Sparse
- Adjacency list allows us to quickly retrieve all neighbors of a given node.



- 1:
- 2: 3, 4
- 3: 2, 4
- 4: 5
- 5: 1, 2

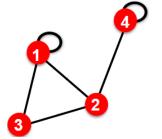
Node and Edge Attributes, Types of Graphs

Possible options for node and edge attributes:

- Weight (e.g., frequency of communication)
- Ranking (best friend, second best friend...)
- Type (friend, relative, co worker)
- Sign: Friend vs. Foe, Trust vs. Distrust
- Properties depending on the structure of the rest of the graph: Number of common friends.

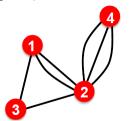
Graphs could be **connected**: Any two vertices can be joined by a path A **disconnected graph** is made up by two or more connected components

Self-edges (with self-loops, undirected)



$$A = \begin{pmatrix} 1 & 1 & 1 & 0 \\ 1 & 0 & 1 & 1 \\ 1 & 1 & 0 & 0 \\ 0 & 1 & 0 & 1 \end{pmatrix}, A_{ii} \neq 0, A_{ij} = A_{ji}$$

Multigraph (undirected)

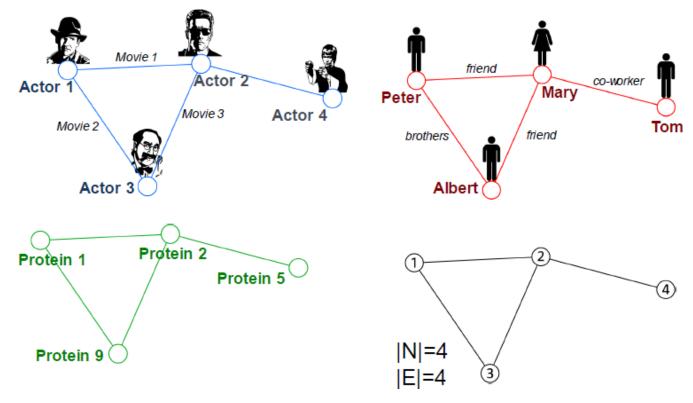


$$A = \begin{pmatrix} 1 & 1 & 1 & 0 \\ 1 & 0 & 1 & 1 \\ 1 & 1 & 0 & 0 \\ 0 & 1 & 0 & 1 \end{pmatrix}, A_{ii} \neq 0, A_{ij} = A_{ji}$$

$$A = \begin{pmatrix} 0 & 2 & 1 & 0 \\ 2 & 0 & 1 & 3 \\ 1 & 1 & 0 & 0 \\ 0 & 3 & 0 & 0 \end{pmatrix}, A_{ii} = 0, A_{ij} = A_{ji}$$

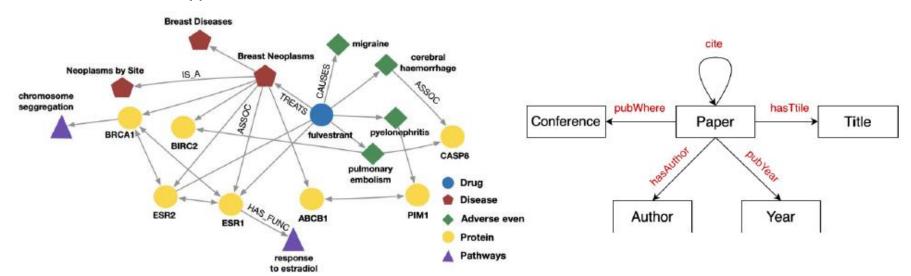
Component of a Graph, Graphs are Abstractions

- Objects: nodes, vertices N or V
- Interactions: links, edges E
- System: network, graph G(N,E) or G(V,E)
- Graphs are abstraction. The same graph could be used for a variety of applications
- Graphs are a common language for many domains. All problems depicted below can be represented by the same graph with 4 vertices and 4 edges.



Many Graphs are Heterogeneous Graphs

- A heterogeneous graph is defined as: G = (V, E, R, T)
 - Vertices (nodes) with node types $v_i \in V$
 - Edges with relation types (v_i, r, v_j)
 - Node type $T(v_i)$
 - Relation type $r \in R$



Biomedical Knowledge Graphs

Example node: Migraine

Example edge: (fulvestrant, Treats, Breast Neoplasms)

Example node type: Protein

Example edge type (relation): Causes

Academic publications graph

Example node: ICML

Example edge: GraphCN, NeurIPS

Example node: Author

Example edge type (relation) pubYear

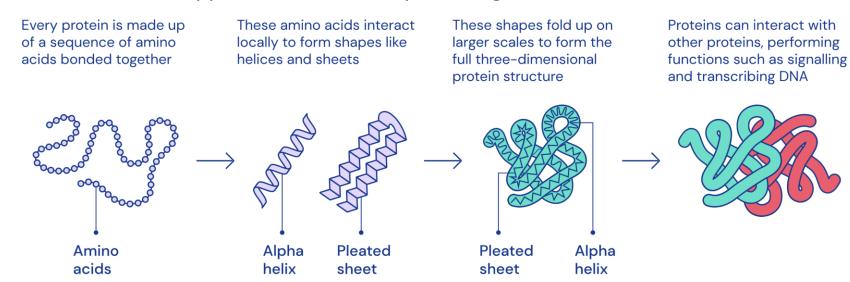
Examples of Uses of GNNs

Graph Neural Networks

- A graph neural network (GNN) is an algorithm that allows us to learn from graphs, including predicting or classifying properties of nodes, edges and graph features.
- Graph neural networks are similar to conventional neural networks, except the learning takes place on graph data.
- Specialized neural network architecture for graphs was first introduced in the late 1990s, gained traction in the early 2000s, and greatly accelerated after 2015.
- Today, there are many examples of enterprise systems (Google, Alibaba, others) that employ GNNs at scale.
- GNNs are profitably used in speeding up drug discovery.
- GNNs is still a rapidly developing field, with many remaining challenges

Example: Protein Folding

- A protein chain in solution folds into a 3D structure. Knowledge of that structure is
 essential for understanding how a protein interacts with other proteins.
- One successful application of GNNs is predicting such 3D structures.



AlphaFold: Using AI for scientific discovery | DeepMind

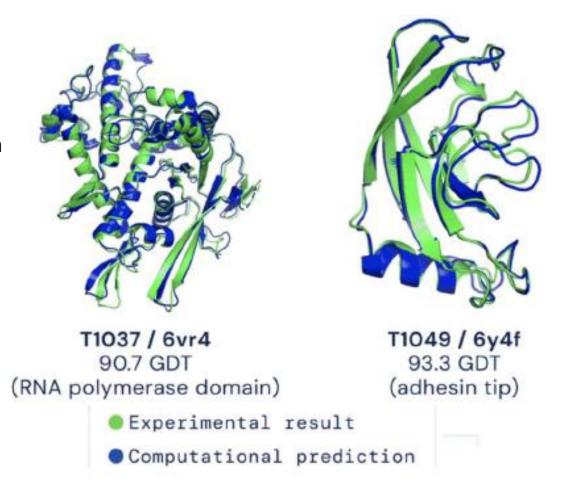
AlphaFold

 AlphaFold is a project of DeepMind a Google company. DeepMind developed GNNs which predict the structure (folding) of proteins. https://www.nature.com/articles/s41586-019-1923-7.epdf

Sequence 0.4 0.3 0.2 Deep neural Distance and torsion Gradient descent on and MSA protein-specific potential network distribution predictions features Noisy restarts 64 bins deep b 2D covariation featur 0.8 0.7 0.6 0.5 0.4 0.3 0.2 Ried L × f 1D sequence and profile features TM score TM score - r.m.s.d. 800 1,000 1,200 400 600 120 100 80 60 40 220 residual convolution blocks Prediction x-1 Gradient descent steps

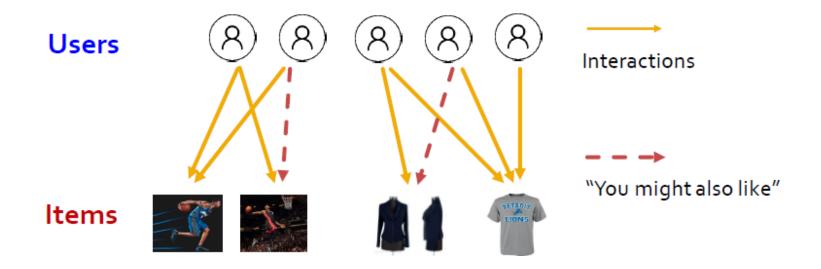
Protein Folding

- GNNs could computationally predict proteins' 3D structure based solely on protein's amino acid sequence.
- Key idea: "Spatial graph"
- Nodes: Amino acids in a protein sequence
- Edges: Proximity between amino acids (residues)



Recommender Systems

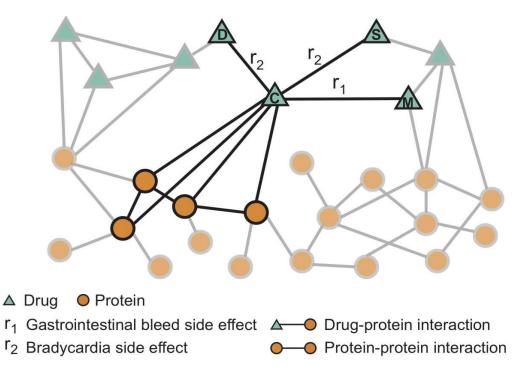
- We can build recommender systems using GNNs.
- Users interacts with items
 - Watch movies, buy merchandise, listen to music
 - Nodes: Users and items
 - Edges: User item interactions
- Goal: Recommend items users might like



Drug Side Effects

- Many patients take multiple drugs to treat complex or co-existing diseases:
 - 46% of people ages 70-79 take more than 5 drugs
 - Many patients take more than 20 drugs to treat heart disease, depression, insomnia, etc.
- Given a pair of drugs, GNNs can predict adverse side effects
- Modeling Polypharmacy Side Effects with Graph Convolutional Networks (arxiv.org)
- Nodes: Drugs & Proteins
- **Edges**: Interactions

Query: How likely will Simvastatin and Ciprofloxacin, when taken together, break down muscle tissue?



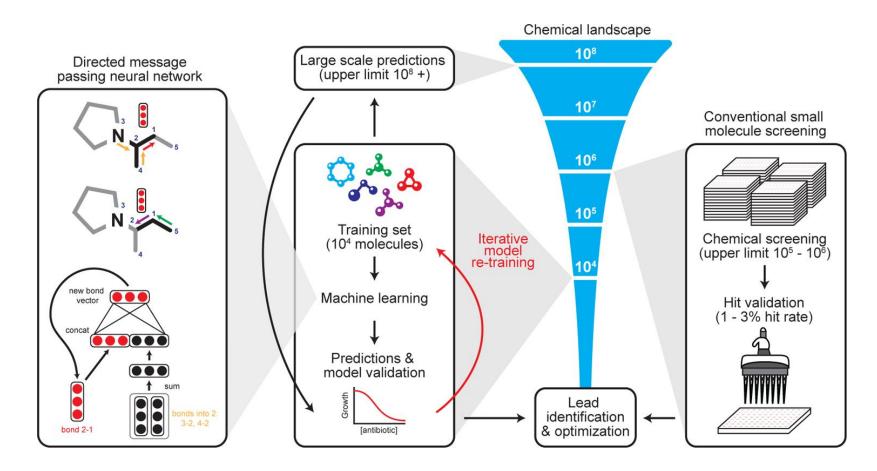
Drug Discovery

Antibiotics are small molecular graphs

Nodes: Atoms

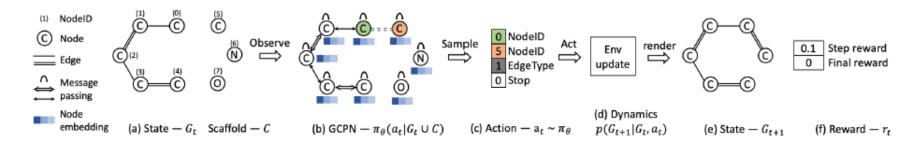
Edges: Chemical bonds

 A classification model based on Graph Neural Network can predict promising molecules from a large pool of candidates (https://pubmed.ncbi.nlm.nih.gov/32084340/)

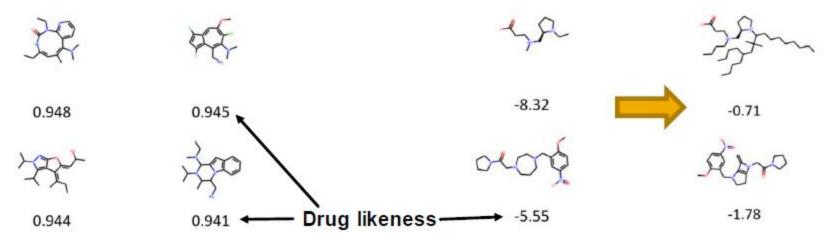


Molecule Generation - Optimization

GNNs can generate novel molecules

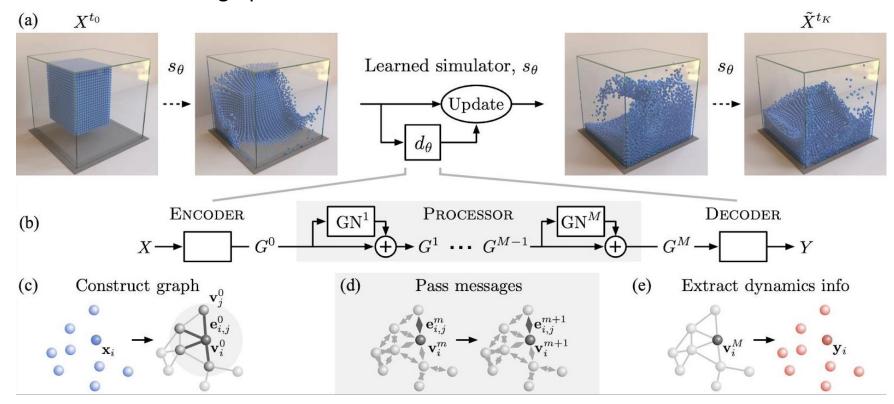


- Use case 1: Generate novel molecules with high drug likeness value
- Use case 2: Optimize existing molecules to have desirable properties



Physical Simulations

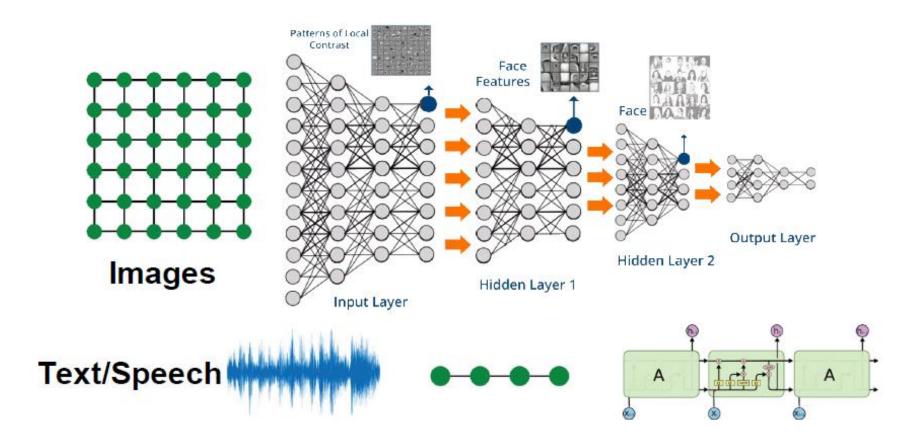
- We could treat a physical structure as a dynamic graph
 - Nodes: particles
 - Edges: interactions between particles
 - Predict: how graph evolves over time



Graph Analysis & Embeddings

Legacy Neural Networks

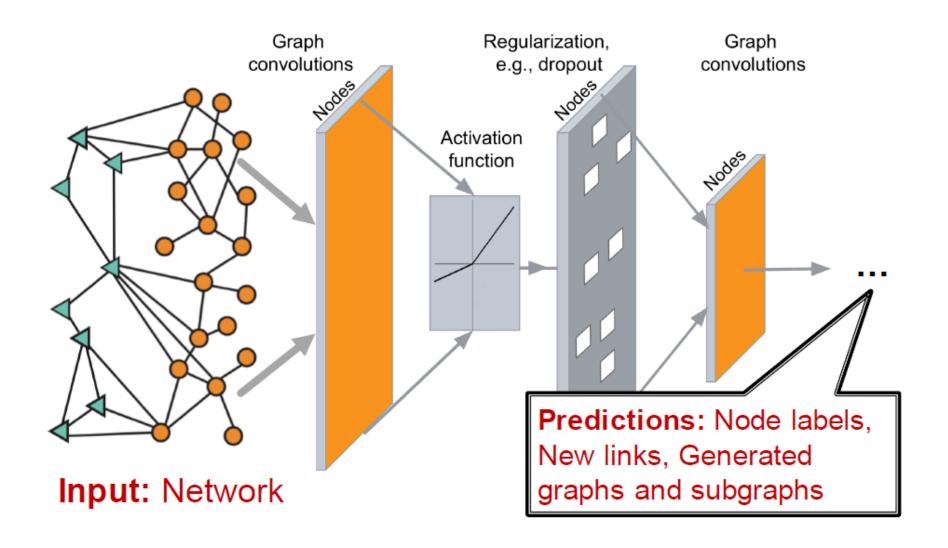
"Classical" deep learning toolbox is designed for simple sequences & grids



Graph Analysis

- A Graph Analysis (e.g., Graph Neural Network's forward pass) can be thought of as two processes:
 - a) converting input graphs into useful embeddings, and/or
 - b) performing some downstream task (e.g., classification) on the embeddings, which converts the embeddings into some useful output.
- Typically, in Graph Analysis we produce three output types:
 - (1) **Vertex-level outputs,** when we require a prediction (e.g., a distinct class or predicted property value) for each vertex in a graph.
 - (2) **Edge-level outputs,** when we make a prediction for one or many properties of edges in a graph.
 - (3) **Graph-level outputs,** when we require a prediction for a property or feature of an entire graph. For example, solubility in a certain liquid is a property of the entire molecule (graph).

Forward Transformation of Graphs through GNNs

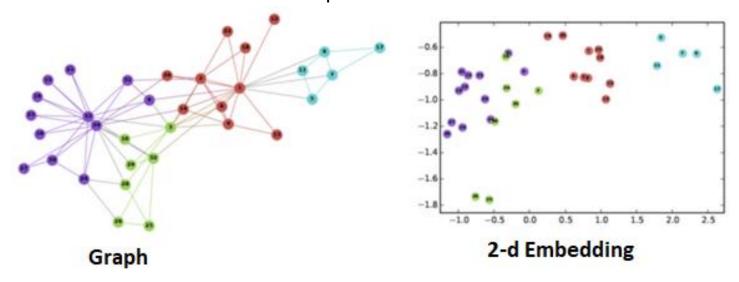


Embeddings

- **Embeddings** are compressed feature representations. By reducing large feature vectors associated with vertices and edges into low dimensional embeddings, we make it possible to classify vertices and edges with lower-order models.
- A key measure of an embedding's quality is whether the (cosine or some other) similarity between nodes or edges in the original space (graph) is retained in the embedding space.
- Embeddings can be created (or learned) for vertices, edges, neighborhoods, or the entire graphs.
- Embeddings are also referred to as *representations*, encodings, *latent vectors*, or high-level *feature vectors* depending on the context.

Graph Representation

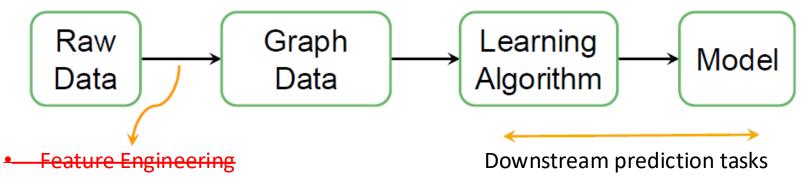
- The challenge of dealing with non-Euclidian graphs is to represent them accurately in a way that can be fed as an input into a "conventional" neural network layers.
- In GNNs, the starting layers produce embeddings which transform a graph structure into a vector or matrix representation.



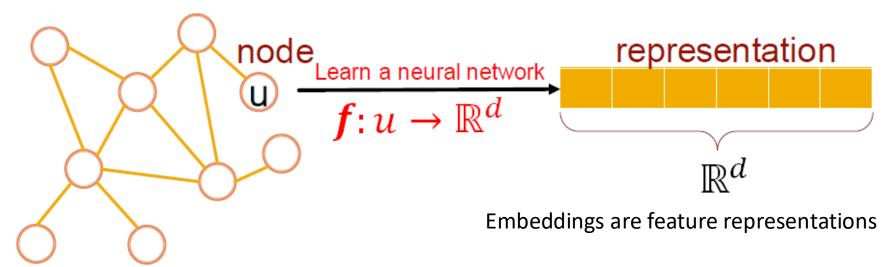
• Above is an illustration of graph embeddings. We start (left) with a non-Euclidean graph structure and end up with embedding vectors (right) that can be projected into a 2D Euclidean space.

Machine Learning Lifecycle: Representation Learning

Representation Learning – Automatically learn the features (embeddings)

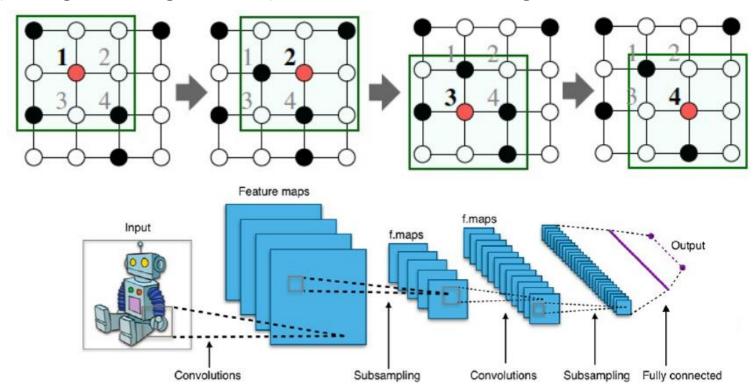


 Initial layers of GNNs map nodes to d-dimensional vectors (embeddings) such that similar nodes in the network are embedded close together.



Implementing a NN on a Graph

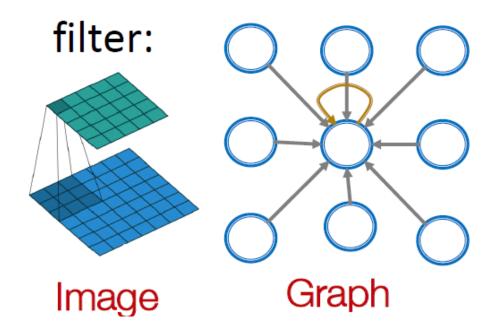
- We understand CNNs well. So, our goal is to generalize convolutions beyond simple lattices.
- We want to use, leverage node features/attributes (e.g., text, images).
- CNNs on images relied on the input organized as a regular grid and filters (flashlights, sliding windows) we move over those images.



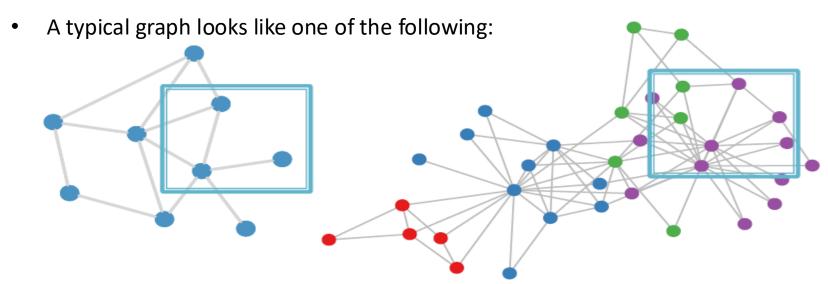
The question is, could we do something similar on a graph?

CNN vs. GNN

- CNN can be seen as a special GNN with fixed neighbor size and ordering:
 - In CNNs, the size of the filter is pre-defined.
 - GNN process arbitrary graphs with different neighborhoods for each node.



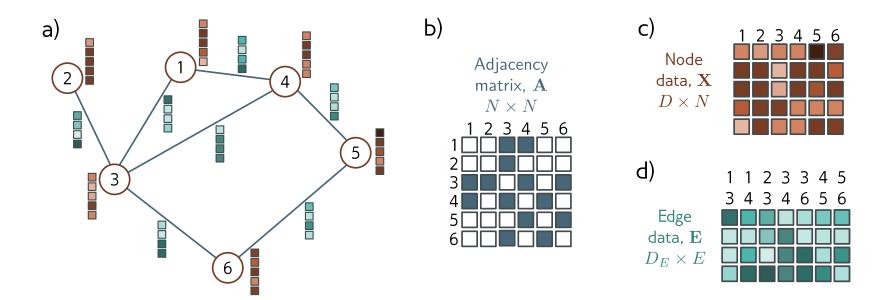
A typical graph is Permutation Invariant



- How do you systematically move a "window" over the graph.
- There is no fixed notion of locality or simple way to define a sliding window on the graph
- Graph is permutation invariant. Relabeling the nodes should not change anything. We want any schema we define to keep on working under arbitrary permutation of node labels.
- Permutation invariance means that a graph does not have a canonical order of the nodes!
- We can have many different order plans. Any labeling of nodes should be equivalent to any other. We cannot take the adjacency matrix too seriously. Relabeling will result in very different adjacency matrix. Graph and node representation should be the same for any node labeling.

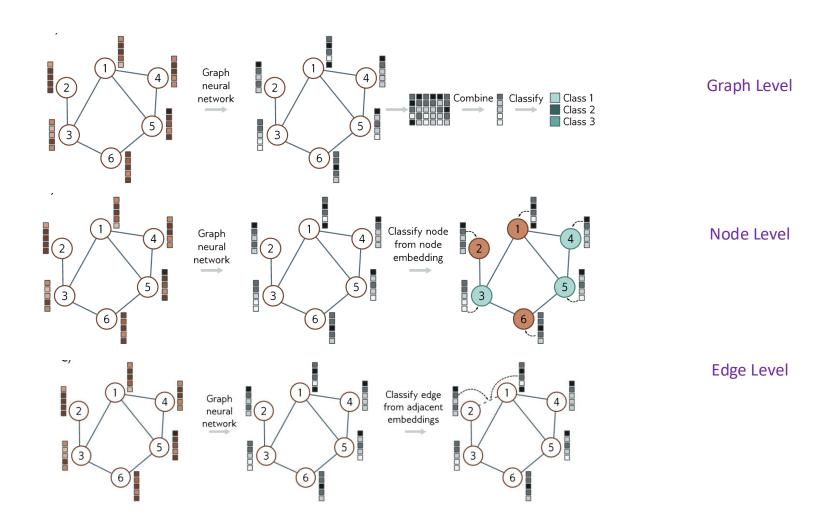
Implementing GNNs

Thinking in Graphs



- a) A Graph with 6 Nodes and respective Node and Edge features
- b) Adjacency Matrix
- c) Node Data Matrix
- d) Edge Data Matrix

Thinking in Graphs

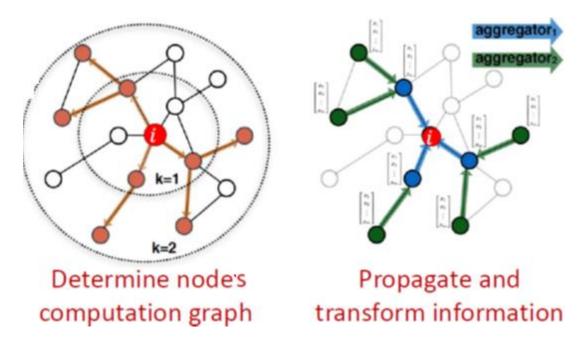


Central Idea of GNN

The central idea of GNN's is the neighborhood aggregation scheme which generates node embeddings by combining information from neighborhoods!

Permutation Invariance, Graph Convolutional Networks

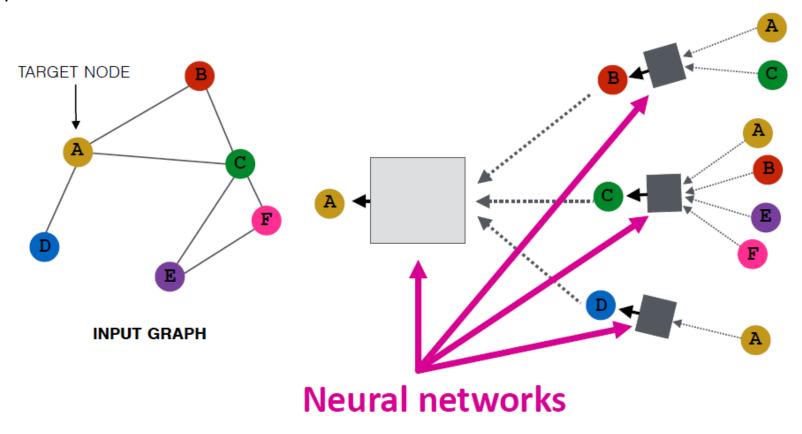
- We design graph neural networks that are permutation invariant by passing and aggregating information from every node's neighbors!
- Key idea: Node's neighborhood defines a local computation graph.



- We propagate information across the graph and compute node features.
- We generate node representations (embeddings) based on local network neighborhoods.
- Calculations are iterative and are repeated until the features are stabilized.

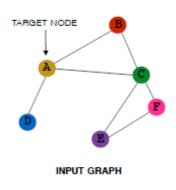
Aggregate Neighbor's Features

- Key idea: Generate node embeddings based on local network neighborhoods.
- Intuition: Nodes aggregate information from their neighbors using an iterative process we will call the neural network.



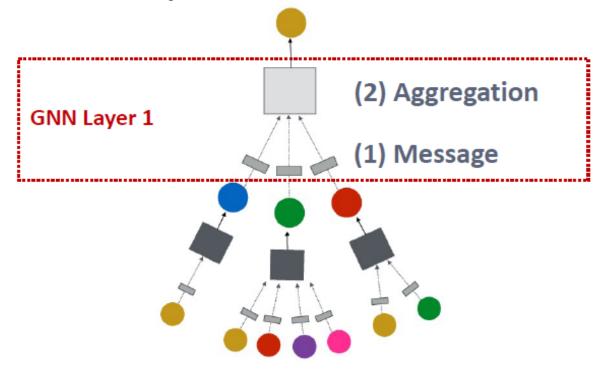
GNN Layer = Message + Aggregation

- There are two key notions in this process: messages and aggregations
- Different GNN architectures define messages and aggregations in different ways



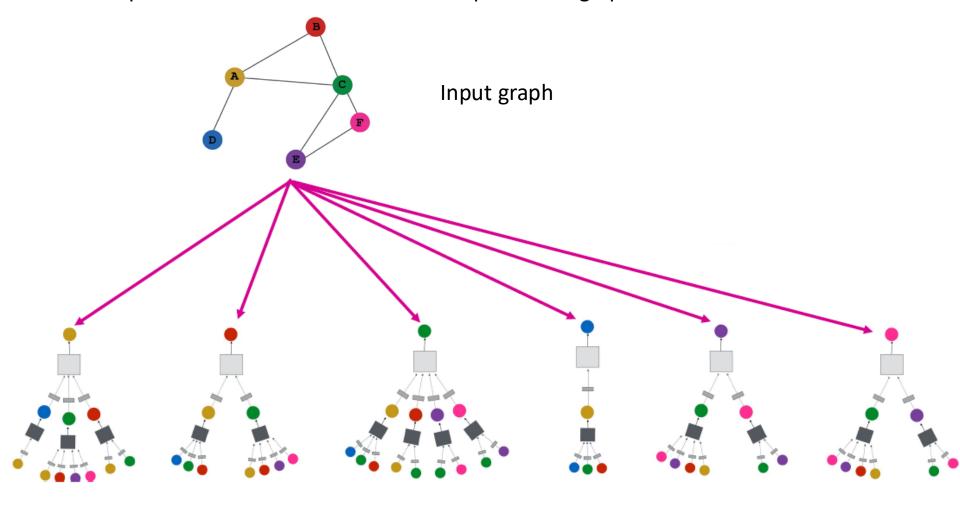
GNN Layer = Message + Aggregation

- Different instantiations under this perspective
- GCN, GraphSAGE, GAT, ...



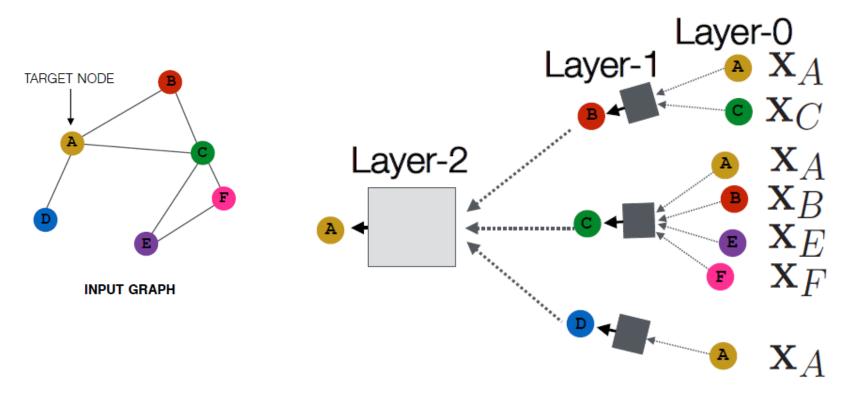
Aggregating Neighbors

- **Intuition:** Network neighborhood defines a computation graph.
- Every node defines a (local) computational graph based on its neighborhood!
- Graph with 6 nodes will result in 6 computational graphs.



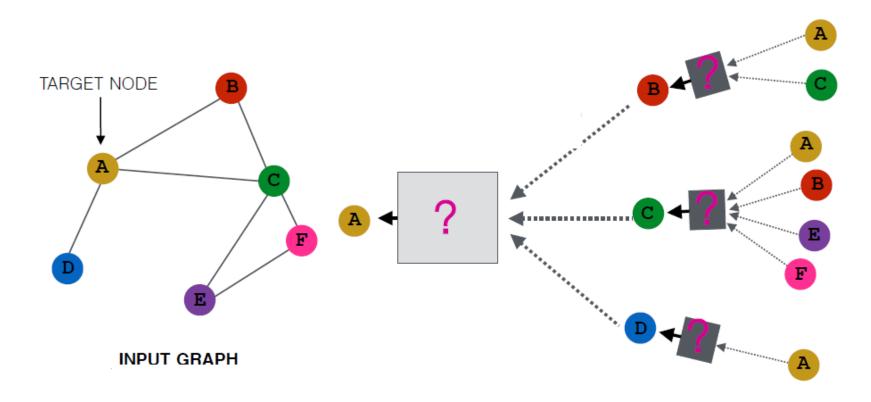
Depth of the Model: How many layers

- What we proposed on the previous slide implies recursion. All nodes source in inputs from all their neighbors and so on.
- Model can be of arbitrary depth:
 - Nodes have embeddings at each layer
 - Layer-0 embedding of node v is its input feature x_v
 - Layer-k embedding gets information from nodes that are k hops away

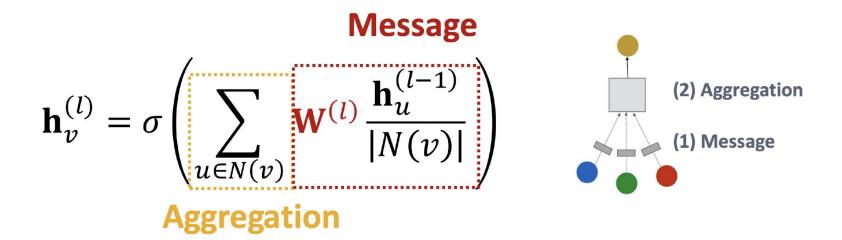


Neighborhood Aggregation

- How we perform aggregation of information (generate embeddings) from the neighborhood is somewhat arbitrary.
- Different types of GNNs perform aggregation in different way.
- Different GNN types are define by the "content" of the box?

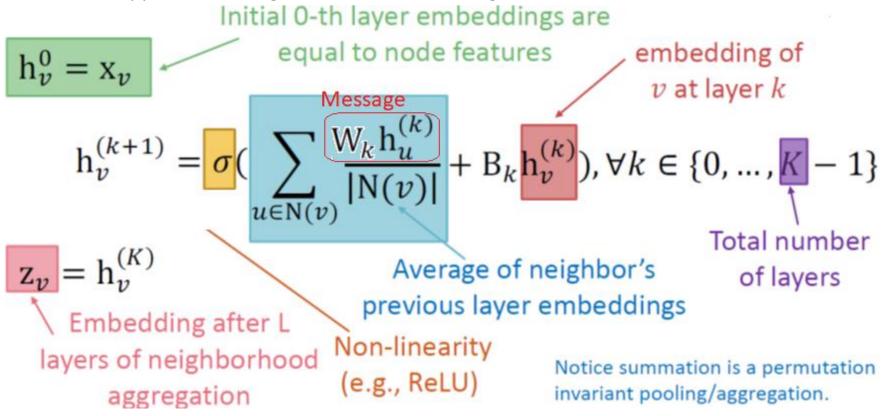


Basic Approach : Message Passing and Aggregation (Simplified)



Basic approach

Basic approach: Average information from neighbors



- Message passing and neighbor aggregation in graph convolution networks is permutation equivariant. Every target node v will have the same computational graph irrespective of different labels (permutations) attached to nodes.
- W_k is trainable weight matrix for neighborhood aggregation. B_k weight matrix for transforming hidden vector of self. h_v^k the hidden representation of node v at layer k.

Training the model, Matrix Formulation

- How do we train the GCN to generate embeddings?
- Need to define a loss function on the embeddings.
- We can feed embeddings h_v^k into any loss function and run SGD to train the weight parameters.
- Many aggregations can be performed efficiently by (sparse) matrix operations.
- We define the matrix of hidden embeddings as

$$H^{(k)} = \left[\begin{array}{ccc} h_1^{(k)} & \dots & h_V^{(k)} \end{array} \right] \ ^T$$

• Let D is the "degree" matrix. It is a diagonal matrix listing the number of neighbors $N\left(v\right)$ of any node v

$$D_{v,v} = \text{Deg}(v) = |N(v)|$$

• The inverse of $D: D^{-1}$ is also diagonal:

$$D_{v,v}^{-1} = 1/|N(v)|$$

• $\sum_{u \in N(v)} \frac{h_u^{(k-1)}}{|N(v)|}$ can be written as $H^{(k+1)} = D^{-1}AH^{(k)}$

Matrix formulation

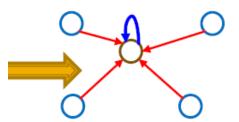
One can show that the update function could be written in the matrix form as:

$$H^{(k+1)} = \sigma(D^{-1}AH^{(k)}W_k^T + H^{(k)}B_k^T)$$

- Red: neighborhood aggregation
- Blue: self transformation

$$H^{(k)} = \left[\begin{array}{ccc} h_1^{(k)} & \dots & h_V^{(k)} \end{array} \right] \quad ^T$$

is hidden state (embeddings) of all nodes in layer k



- In practice, this implies that efficient sparse matrix multiplication can be used (matrix $D^{-1}A$ is sparse)
- Note: not all GNNs can be expressed in matrix form. Sometimes, aggregation function is too complex to be written as a matrix expression.
- Number of layers (k) in GNNs is not arbitrarily large. Typically, we go through a small number recursions. Practice has shown that too many recursions would result in smoothing out of node and edge features and all states of all nodes will start approaching the same value.

Node Similarity

- When calculating Cross Entropy for the Loss Function we will use node similarity. The simplest **node similarity** can be defined as "nodes u, v are similar if they are connected by an edge."
- This means: $z_v^T z_u = A_{u,v}$ which is the (u,v) entry of the graph adjacency matrix A
- Therefore, $Z^TZ = A$
- Node could be similar even if they are not adjacent.
- There are several stochastic techniques for assessing that similarity using techniques like the deep (random) walk over node vicinity.
- Most importantly, two nodes are similar if they have high value of the cosine similarity between their embedded (latent) vectors.

Loss Function

• Similar nodes have similar embeddings. The loss function is defined as sum over all node pairs:

$$L = \sum_{(z_u, z_v)} CE(y_{u,v}, (z_u, z_v))$$

- Where $y_{u,v}$ is node similarity and can be 1 when node u and v are **similar** or could be less than 1
- CE stands for the cross entropy.
- Node similarity can be calculated in different ways, using
 - Random walks (node2vec, DeepWalk, struc2vec)
 - Matrix factorization
 - Node proximity in the graph

Loss Function in Supervised Training

- We could directly train the model for a supervised task (e.g., node classification) such as discovering whether a drug is safe or toxic.
- Safety of a drug is in this case a binary feature of the drug.
- Drug-drug and drug-protein interactions are edges of the graph.
- In these problems, we usually use simpler Loss Function:

$$L = \sum_{v \in V} c_v \log(\sigma(z_v^T \theta)) + (1 - c_v) \log(1 - \sigma(z_v^T \theta))$$

• Here z_v is node embedding, heta is classifcation weights and c_v is node class label.

Inductive Capability

- Inductive node embedding -> generalize to entirely unseen graphs
 - We could train on protein interaction graph from model organism A and generate embeddings on newly collected data about organism B.
- Many application settings constantly encounter previously unseen nodes:
 - We need to generate new embeddings "on the fly"

Graph Attention Networks

• In GANs, recursive equation for embedding vector of node v in layer k is defined as:

$$h_v^{(k)} = \sigma(\sum_{u \in N(v)} W^{(k-1)} \alpha_{vu} h_u^{(k-1)})$$

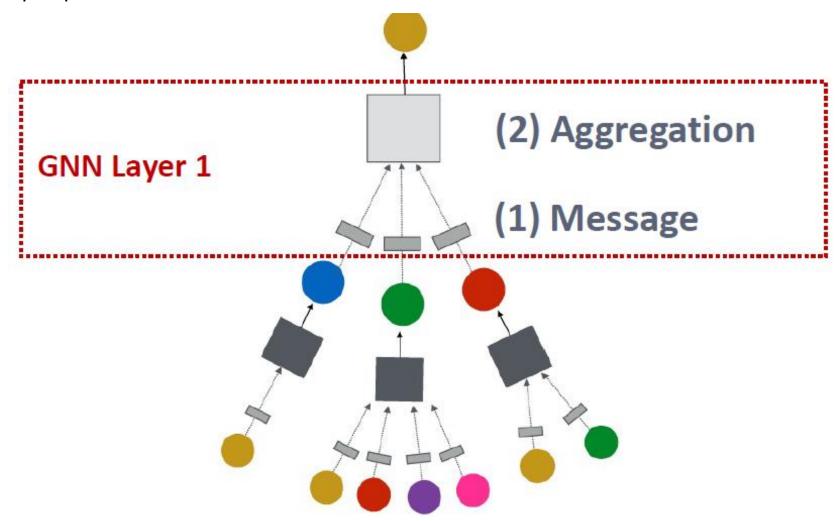
- In Graph Convolutional Networks(GCNs) $\alpha_{vu}={}^1\!/_{N(v)}$ is the weighting factor (importance) of the message coming from note u to node v
- α_{vu} is defined **explicitly** based on the structural properties of the graph (node degree) and the assumption that all neighbors $u \in N(v)$ are equally important to node v.
- Attention model starts with assumption that not all node's neighbors are equally important
- Attention is inspired by cognitive attention and the attention mechanism we will study
 in NLP
- The **attention** α_{vu} focuses on the important parts of the input data and fades out the rest.
- Idea: the NN should devote more computing power on that small but important part of the data.
- Through training, depending on the context, network will learn attention coefficients and find which neighbors are more important.

Message Passing GNNs

- The basic intuition behind the GNN message-passing framework is straightforward: at each iteration, every node aggregates information from its local neighborhood, and as these iterations progress each node embedding contains more and more information from further reaches of the graph. To be precise: after the first iteration (k=1), every node embedding contains information from its 1-hop neighborhood, i.e., every node embedding contains information about the features of its immediate graph neighbors, which can be reached by a path of length 1 in the graph; after the second iteration (k=2) every node embedding contains information from its 2-hop neighborhood; and in general, after k iterations every node embedding contains information about its k-hop neighborhood.
- But what kind of "information" do these node embeddings actually encode? Generally, this information comes in two forms. On the one hand there is structural information about the graph. For example, after k iterations of GNN message passing, the embedding $h_u^{(k)}$ of node u might encode information about the degrees of all the nodes in u's k-hop neighborhood. This structural information can be useful for many tasks. For instance, when analyzing molecular graphs, we can use degree information to infer atom types and different structural motifs, such as benzene rings.
- In addition to structural information, the other key kind of information captured by GNN node embedding is feature-based. After k iterations of GNN message passing, the embeddings for each node also encode information about all the features in their k-hop neighborhood. This local feature-aggregation behavior of GNNs is analogous to the behavior of the convolutional kernels in convolutional neural networks (CNNs). However, whereas CNNs aggregate feature information from spatially-defined patches in an image, GNNs aggregate information based on local graph neighborhoods

GNN Layer = Message + Aggregation

 We can generalize operations in all GNN as different instantiations under this perspective:

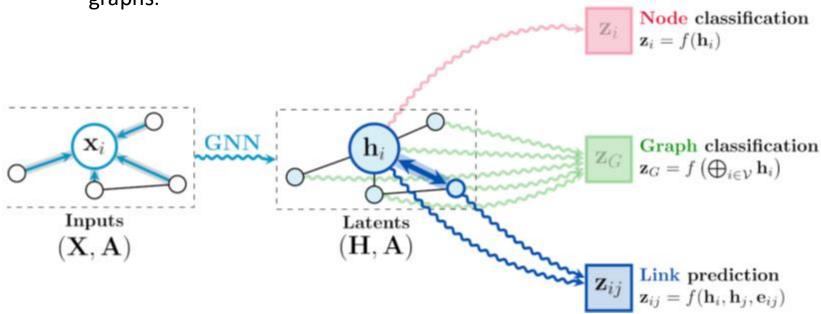


Types of GNNs

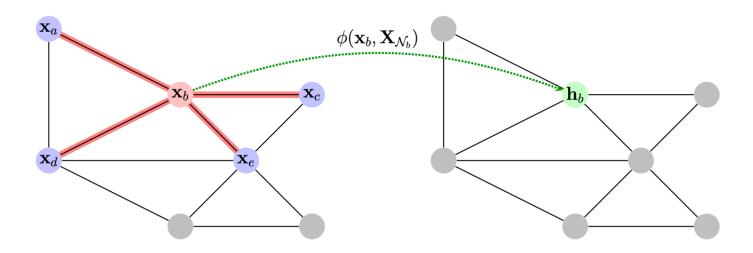
General blueprint for learning on graphs

- Given some functions X on the domain made of nodes and/or edges of a graph with topology described by adjacency matrix A, we calculate embeddings or latent values H.
- Starting with latent values *H* and adjacency matrix *A*, we could make:
 - predictions on properties of nodes or perform node classifications,
 - predictions on the properties of edges or classification of edges, and make

 predictions on the properties of the entire graphs or classification of graphs.

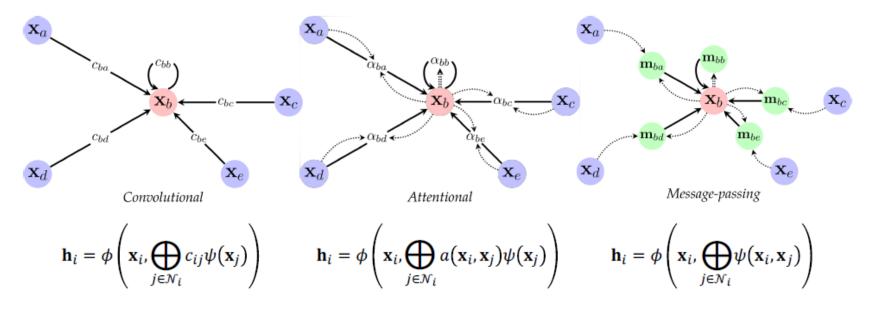


Recipe for graph neural networks, visualized



- Results of any calculations described on the previous slide in any layer of any GNN are vector $\pmb{embeddings}$ or the $\pmb{latent\ vectors\ }h_b$
- We build permutation equivariant functions $\mathbf{F}(\mathbf{X}, \mathbf{A})$ on graphs by shared application of a local permutation-invariant function $\phi(x_i, X_{N_i})$
- Common terminology used in GNNs is:
 - F is a "GNN layer"
 - ϕ is "diffusion", "propagation" or "message passing"
- How do we implement ϕ is a very intense area of research!
- Fortunately, almost all of layers can be classified across three "flavors"

The three "flavors" of GNN layers



- Most of GNNs can be classified as Convolutional, Attention or Message-passing.
- Symbol \oplus stands for some generalized aggregation. It could be summation, multiplication, concatenation, whatever.
- **Convolutional GNNs** are most frequently encountered in social networks, where edges encode similarity between nodes. Highly scalable. Most industry apps are Conv GNNs
- Attention GNNs aggregate features of neighbors with implicit weights (attention) $\alpha_{ij} = a(x_i, x_j)$. Useful as "middle ground" w.r.t. capacity, scale, interpretability. Edges need not encode similarity between nodes. These networks are still computing only a **scalar** per edge.
- Message-passing GNNs send arbitrary vectors (messages) across edges. Messages computed as $m_{ij} = \psi(x_i, x_j)$. Most generic GNN layer. Edges give "recipe" for passing data. Have scalability or learnability issues. Ideal for computational chemistry, reasoning and simulation tasks

Node Classification with GNNs

Node Classification with Graph Neural Networks

- This example demonstrate an implementation of a Graph Neural Network (GNN) model.
- The model is used for a node prediction task on the Cora dataset of scientific papers. The model will predict the subject of a paper given its words and citations network.
- We implement a Graph Convolution Layer from scratch to provide better understanding of how they work. However, there are specialized TensorFlow-based libraries that provide rich GNN APIs, such as Spectral, StellarGraph, and GraphNets.

Setup: We need networkx, a Python library for network manipulation and analysis !pip install network

```
import os
import pandas as pd
import numpy as np
import networkx as nx
import matplotlib.pyplot as plt
import tensorflow as tf
from tensorflow import keras
from tensorflow.keras import layers
```

Fetch the Dataset

- The Cora dataset consists of 2,708 scientific papers classified into one of seven classes. The citation network consists of 5,429 links. Each paper has a binary word vector of size 1,433, indicating the presence of a word with a given index.
- The dataset has two tab-separated files: cora.cites and cora.content.
- The cora.cites includes the citation records with two columns: cited_paper_id (target) and citing_paper_id (source).
- The cora.content includes the paper content records with 1,435 columns: paper id, subject, and 1,433 binary features.

Download the dataset

```
zip_file = keras.utils.get_file(
    fname="cora.tgz",
    origin="https://linqs-data.soe.ucsc.edu/public/lbc/cora.tgz",
    extract=True,
)
data_dir = os.path.join(os.path.dirname(zip_file), "cora")
```

• The dataset ends up in you home directory, subdirectory .keras/datasets/cora. On my Windws machine that is: C:\Users\Zoran\.keras\datasets\cora

Process and visualize the dataset, citations

Load the citations data, file cora.cites into a Pandas DataFrame.

```
citations = pd.read_csv(
    os.path.join(data_dir, "cora.cites"),
    sep="\t",
    header=None,
    names=["target", "source"],
)
print("Citations shape:", citations.shape)
Citations shape: (5429, 2)
```

• Next we display a sample of the citations DataFrame. The target column includes the paper ids cited by the paper ids in the source column.

```
citations.sample(frac=1).head()
```

	target	source	
65	35	1956	
5037	578646	1128291	
1195	6184	1120731	
3927	100197	447250	
4686	250566	1113926	

Visualize papers

We will load the papers data into a Pandas DataFrame.

```
column_names = ["paper_id"] + [f"term_{idx}" for idx in range(1433)] + ["subject"]
papers = pd.read_csv(
    os.path.join(data_dir, "cora.content"), sep="\t", header=None,
names=column_names,
)
print("Papers shape:", papers.shape)
Papers shape: (2708, 1435)
```

• The DataFrame papers includes the paper_id and the subject columns, as well as 1,433 binary column representing whether a term exists in the paper or not. A sample of papers reads:

<pre>print (papers.sample(5).T)</pre>						
	615	1863	2594	2079	1005	
paper_id	101263	107252	1105603	101660	416964	
term_0	0	0	0	0	0	
term_1	0	0	0	0	0	
term_2	0	0	0	0	0	
term_3	0	0	0	0	0	
• • •	• • •	• • •	• • •	• • •	• • •	
term_1429	0	0	0	0	0	
term_1430	0	0	0	0	0	
term_1431	0	0	0	0	0	
term_1432	0	0	0	0	0	
subject	Theory	Theory	Neural_Networks	Rule_Learning	Neural_Networks	

[1435 rows x 5 columns]

• Note that we are presenting a transposed matrix sample (5). Every paper is a column.

The count of the papers in each subject

```
print(papers.subject.value_counts())
Neural_Networks 818
Probabilistic_Methods 426
Genetic_Algorithms 418
Theory 351
Case_Based 298
Reinforcement_Learning 217
Rule_Learning 180
Name: subject, dtype: int64
```

Next, we convert the paper ids and the subjects into zero-based indices.

```
class_values = sorted(papers["subject"].unique())
class_idx = {name: id for id, name in enumerate(class_values)}
paper_idx = {name: idx for idx, name in
enumerate(sorted(papers["paper_id"].unique()))}

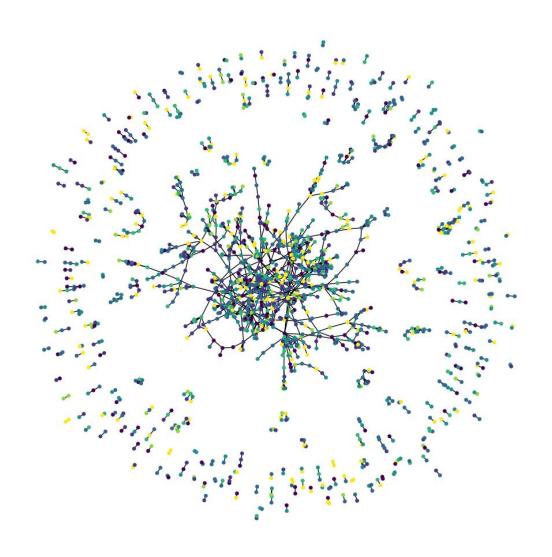
papers["paper_id"] = papers["paper_id"].apply(lambda name: paper_idx[name])
citations["source"] = citations["source"].apply(lambda name: paper_idx[name])
citations["target"] = citations["target"].apply(lambda name: paper_idx[name])
papers["subject"] = papers["subject"].apply(lambda value: class idx[value])
```

The citation graph

 Now let's visualize the citation graph. Each node in the graph represents a paper, and the color of the node corresponds to its subject. We only show a sample of the papers in the dataset.

```
plt.figure(figsize=(10, 10))
colors = papers["subject"].tolist()
cora_graph = nx.from_pandas_edgelist(citations.sample(n=1500))
subjects =
list(papers[papers["paper_id"].isin(list(cora_graph.nodes))]["subject"])
nx.draw_spring(cora_graph, node_size=15, node_color=subjects)
```

The citation graph



Split the dataset into stratified train and test sets

```
train_data, test_data = [], []

for _, group_data in papers.groupby("subject"):
    # Select around 50% of the dataset for training.
    random_selection = np.random.rand(len(group_data.index)) <= 0.5
    train_data.append(group_data[random_selection])
    test_data.append(group_data[~random_selection])

train_data = pd.concat(train_data).sample(frac=1)

test_data = pd.concat(test_data).sample(frac=1)

print("Train data shape:", train_data.shape)

print("Test data shape:", test_data.shape)</pre>
```

Train data shape: (1325, 1435) Test data shape: (1383, 1435)

Train and Evaluate Experiment

```
hidden_units = [32, 32]
learning_rate = 0.01
dropout_rate = 0.5
num_epochs = 300
batch_size = 256
```

This function compiles and trains an input model using the given training data.

```
def run experiment(model, x train, y train):
    # Compile the model.
    model.compile(
        optimizer=keras.optimizers.Adam(learning rate),
        loss=keras.losses.SparseCategoricalCrossentropy(from logits=True),
        metrics=[keras.metrics.SparseCategoricalAccuracy(name="acc")],
    # Create an early stopping callback.
    early stopping = keras.callbacks.EarlyStopping(
        monitor="val acc", patience=50, restore best weights=True
    # Fit the model.
    history = model.fit(
        x=x train,
       y=y train,
        epochs=num epochs,
        batch size=batch size,
        validation split=0.15,
        callbacks=[early stopping],
    return history
```

The loss and accuracy curves

This function displays the loss and accuracy curves of the model during training.

```
def display learning curves (history):
    fig, (ax1, ax2) = plt.subplots(1, 2, figsize=(15, 5))
    ax1.plot(history.history["loss"])
    ax1.plot(history.history["val loss"])
    ax1.legend(["train", "test"], loc="upper right")
    ax1.set xlabel("Epochs")
    ax1.set ylabel("Loss")
    ax2.plot(history.history["acc"])
    ax2.plot(history.history["val acc"])
    ax2.legend(["train", "test"], loc="upper right")
    ax2.set xlabel("Epochs")
    ax2.set ylabel("Accuracy")
   plt.show()
```

Feedforward Network (FFN) Model

We will use this module in the baseline and the GNN models.

```
def create_ffn(hidden_units, dropout_rate, name=None):
    fnn_layers = []

for units in hidden_units:
        fnn_layers.append(layers.BatchNormalization())
        fnn_layers.append(layers.Dropout(dropout_rate))
        fnn_layers.append(layers.Dense(units, activation=tf.nn.gelu))

return keras.Sequential(fnn_layers, name=name)
```

Prepare the data for the baseline model

```
feature_names = set(papers.columns) - {"paper_id", "subject"}
num_features = len(feature_names)
num_classes = len(class_idx)

# Create train and test features as a numpy array.
x_train = train_data[list(feature_names)].to_numpy()
x_test = test_data[list(feature_names)].to_numpy()
# Create train and test targets as a numpy array.
y_train = train_data["subject"]
y_test = test_data["subject"]
```

Implement a baseline classifier

 We add five FFN blocks with skip connections, so that we generate a baseline model with roughly the same number of parameters as the GNN models to be built later.

```
def create_baseline_model(hidden_units, num_classes, dropout_rate=0.2):
    inputs = layers.Input(shape=(num features,), name="input features")
    x = create ffn(hidden units, dropout rate, name=f"ffn block1")(inputs)
    for block idx in range(4):
        # Create an FFN block.
        x1 = create ffn(hidden units, dropout rate,
name=f"ffn block{block idx + 2}")(x)
        # Add skip connection.
        x = layers.Add(name=f"skip connection{block idx + 2}")([x, x1])
    # Compute logits.
    logits = layers.Dense(num classes, name="logits")(x)
    # Create the model.
    return keras.Model(inputs=inputs, outputs=logits, name="baseline")
baseline model = create baseline model (hidden units, num classes,
dropout rate)
baseline model.summary()
```

baseline model.summary()

Model: "baseline"

Layer (type)	Output Shape	Param #	Connected to
input_features (InputLayer)	[(None, 1433)]	0	[]
ffn_block1 (Sequential)	(None, 32)	52804	['input_features[0][0]']
ffn_block2 (Sequential)	(None, 32)	2368	['ffn_block1[0][0]']
skip_connection2 (Add)	(None, 32)	0	['ffn_block1[0][0]', 'ffn_block2[0][0]']
ffn_block3 (Sequential)	(None, 32)	2368	['skip_connection2[0][0]']
skip_connection3 (Add)	(None, 32)	0	<pre>['skip_connection2[0][0]', 'ffn_block3[0][0]']</pre>
ffn_block4 (Sequential)	(None, 32)	2368	['skip_connection3[0][0]']
skip_connection4 (Add)	(None, 32)	0	<pre>['skip_connection3[0][0]', 'ffn_block4[0][0]']</pre>
ffn_block5 (Sequential)	(None, 32)	2368	['skip_connection4[0][0]']
skip_connection5 (Add)	(None, 32)	0	<pre>['skip_connection4[0][0]', 'ffn_block5[0][0]']</pre>
logits (Dense)	(None, 7)	231	['skip_connection5[0][0]']

Total params: 62,507 Trainable params: 59,065 Non-trainable params: 3,442

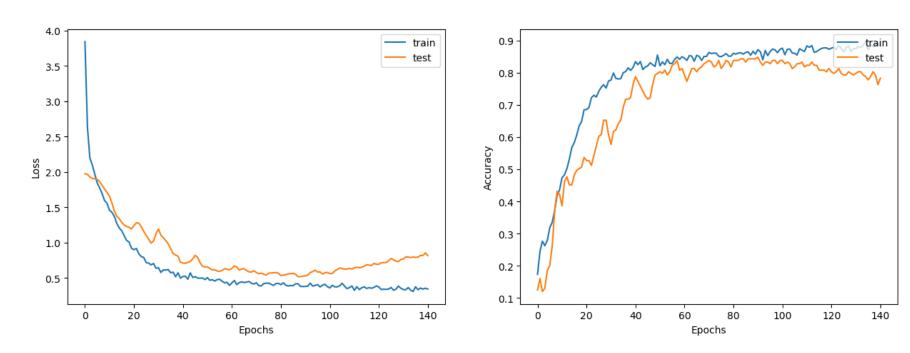
Train the baseline classifier

history = run experiment(baseline model, x train, y train) Epoch 1/300 0.1741 - val loss: 1.9753 - val acc: 0.1256 Epoch 2/300 0.2451 - val loss: 1.9667 - val acc: 0.1608 Epoch 3/300 Epoch 140/300 0.8899 - val loss: 0.8551 - val acc: 0.7638 Epoch 141/300 0.8837 - val loss: 0.8169 - val acc: 0.7839

The learning curves

Let's plot the learning curves.

display learning curves (history)



Evaluate the baseline model on the test data split.

```
_, test_accuracy = baseline_model.evaluate(x=x_test, y=y_test, verbose=0) print(f"Test accuracy: {round(test accuracy * 100, 2)}%")
```

Examine the baseline model predictions

• We will create new data instances by randomly generating binary word vectors with respect to the word presence probabilities.

```
def generate random instances(num instances):
    token probability = x train.mean(axis=0)
    instances = []
    for in range (num instances):
        probabilities = np.random.uniform(size=len(token probability))
        instance = (probabilities <= token probability).astype(int)</pre>
        instances.append(instance)
    return np.array(instances)
def display class probabilities (probabilities):
    for instance idx, probs in enumerate (probabilities):
        print(f"Instance {instance idx + 1}:")
        for class idx, prob in enumerate(probs):
            print(f"- {class values[class idx]}: {round(prob * 100, 2)}%")
```

```
new instances = generate random instances(num classes)
logits = baseline model.predict(new instances)
probabilities =
keras.activations.softmax(tf.convert to tensor(logits)).numpy()
display class probabilities (probabilities)
1/1 [=======] - 2s 2s/step
Instance 1:
- Case Based: 5.96%
- Genetic Algorithms: 1.55%
- Neural Networks: 13.87%
- Probabilistic Methods: 1.23%
- Reinforcement Learning: 2.96%
- Rule Learning: 28.57%
- Theory: 45.86%
Instance 2:
- Case Based: 14.55%
- Genetic Algorithms: 11.0%
- Neural Networks: 34.77%
- Probabilistic Methods: 3.09%
- Reinforcement Learning: 0.92%
- Rule Learning: 0.85%
- Theory: 34.83%
```

The baseline model predictions

Instance 3:

- Case_Based: 0.47%
- Genetic_Algorithms: 97.14%
- Neural_Networks: 1.07%
- Probabilistic Methods: 0.22%
- Reinforcement_Learning: 0.6%
- Rule Learning: 0.09%
- Theory: 0.4%

Instance 4:

- Case Based: 1.16%
- Genetic_Algorithms: 0.71%
- Neural Networks: 91.4%
- Probabilistic Methods: 0.19%
- Reinforcement Learning: 1.05%
- Rule Learning: 0.49%
- Theory: 4.99%

Instance 5:

- Case_Based: 0.36%
- Genetic_Algorithms: 81.57%
- Neural_Networks: 14.92%
- Probabilistic_Methods: 2.47%
- Reinforcement_Learning: 0.52%
- Rule_Learning: 0.05%
- Theory: 0.1%

Instance 6:

- Case Based: 0.17%
- Genetic_Algorithms: 0.94%
- Neural Networks: 18.5%
- Probabilistic_Methods: 0.3%
- Reinforcement Learning: 79.23%
- Rule Learning: 0.35%
- Theory: 0.51%

Instance 7:

- Case_Based: 2.87%
- Genetic_Algorithms: 4.82%
- Neural Networks: 32.97%
- Probabilistic Methods: 5.3%
- Reinforcement Learning: 4.51%
- Rule Learning: 9.11%
- Theory: 40.43%

Graph Neural Network Model

- Preparing and loading the graphs data into the model for training is the most challenging part in GNN models, which is addressed in different ways by the specialized libraries. In this example, we show a simple approach for preparing and using graph data that is suitable if your dataset consists of a single graph that fits entirely in memory.
- The graph data is represented by the graph_info tuple, which consists of the following three elements:
 - 1. node_features: This is a [num_nodes, num_features] NumPy array that includes the node features. In this dataset, the nodes are the papers, and the node_features are the word-presence binary vectors of each paper.
 - 2. edges: This is [num_edges, num_edges] NumPy array representing a sparse adjacency matrix of the links between the nodes. In this example, the links are the citations between the papers.
 - 3. edge_weights (optional): This is a [num_edges] NumPy array that includes the edge weights, which quantify the relationships between nodes in the graph. In this example, there are no weights for the paper citations.

graph_info tuple

```
# Create an edges array (sparse adjacency matrix) of shape [2, num edges].
edges = citations[["source", "target"]].to numpy().T
# Create an edge weights array of ones.
edge weights = tf.ones(shape=edges.shape[1])
# Create a node features array of shape [num nodes, num features].
node features = tf.cast(
    papers.sort values("paper id")[list(feature names)].to numpy(),
dtype=tf.dtypes.float32
# Create graph info tuple with node features, edges, and edge weights.
graph info = (node features, edges, edge weights)
print("Edges shape:", edges.shape)
print("Nodes shape:", node features.shape)
```

Edges shape: (2, 5429) Nodes shape: (2708, 1433)

Graph convolution layer

- We implement a graph convolution module as a Keras Layer. Our GraphConvLayer performs the following steps:
- **1. Prepare**: The input node representations are processed using a FFN to produce a *message*. You can simplify the processing by only applying linear transformation to the representations.
- 2. Aggregate: The messages of the neighbors of each node are aggregated with respect to the <code>edge_weights</code> using a permutation invariant pooling operation, such as sum, mean, and max, to prepare a single aggregated message for each node. See, for example, <code>tf.math.unsorted_segment_sum</code> APIs used to aggregate neighbor messages.
- 3. Update: The node_repesentations and aggregated_messages—both of shape [num_nodes, representation_dim]— are combined and processed to produce the new state of the node representations (node embeddings). If combination_type is gru, the node_repesentations and aggregated_messages are stacked to create a sequence, then processed by a GRU layer. Otherwise, the node_repesentations and aggregated_messages are added or concatenated, then processed using a FFN.
- The technique implemented use ideas from Graph Convolutional Networks, GraphSage, Graph Isomorphism Network, Simple Graph Networks, and Gated Graph Sequence Neural Networks. Two other key techniques that are not covered are Graph Attention Networks and Message Passing Neural Networks.

```
class GraphConvLayer(layers.Layer):
   def init (
        self,
        hidden units,
        dropout rate=0.2,
        aggregation type="mean",
        combination type="concat",
        normalize=False,
        *args,
        **kwargs,):
        super(GraphConvLayer, self). init (*args, **kwargs)
        self.aggregation type = aggregation type
        self.combination type = combination type
        self.normalize = normalize
        self.ffn prepare = create ffn(hidden units, dropout rate)
        if self.combination type == "gated":
            self.update fn = layers.GRU(
                units=hidden units,
                activation="tanh",
                recurrent activation="sigmoid",
                dropout=dropout rate,
                return state=True,
                recurrent dropout=dropout rate,
        else:
            self.update fn = create ffn(hidden units, dropout rate)
```

```
def prepare (self, node repesentations, weights=None):
        # node repesentations shape is [num edges, embedding dim].
       messages = self.ffn prepare(node repesentations)
        if weights is not None:
           messages = messages * tf.expand dims(weights, -1)
        return messages
def aggregate(self, node indices, neighbour messages, node repesentations):
        # node indices shape is [num edges].
        # neighbour messages shape: [num edges, representation dim].
        # node repesentations shape is [num nodes, representation dim].
        num nodes = node repesentations.shape[0]
        if self.aggregation type == "sum":
            aggregated message = tf.math.unsorted segment sum(
                neighbour messages, node indices, num segments=num nodes
        elif self.aggregation type == "mean":
            aggregated message = tf.math.unsorted segment mean(
                neighbour messages, node indices, num segments=num nodes
        elif self.aggregation type == "max":
            aggregated message = tf.math.unsorted segment max(
                neighbour messages, node indices, num segments=num nodes
        else:
            raise ValueError(f"Invalid aggregation type: {self.aggregation type}.")
        return aggregated message
```

```
def update(self, node repesentations, aggregated messages):
       # node repesentations shape is [num nodes, representation dim].
       # aggregated messages shape is [num nodes, representation dim].
       if self.combination type == "gru":
           # Create a sequence of two elements for the GRU layer.
           h = tf.stack([node repesentations, aggregated messages], axis=1)
       elif self.combination type == "concat":
           # Concatenate the node repesentations and aggregated messages.
           h = tf.concat([node_repesentations, aggregated_messages], axis=1)
       elif self.combination type == "add":
           # Add node repesentations and aggregated messages.
           h = node repesentations + aggregated messages
       else:
           raise ValueError(f"Invalid combination type: {self.combination type}.")
       # Apply the processing function.
       node embeddings = self.update fn(h)
       if self.combination type == "gru":
           node embeddings = tf.unstack(node embeddings, axis=1)[-1]
       if self.normalize:
           node embeddings = tf.nn.l2 normalize(node embeddings, axis=-1)
       return node embeddings
```

```
def call(self, inputs):
       """Process the inputs to produce the node embeddings.
       inputs: a tuple of three elements: node repesentations, edges, edge weights.
       Returns: node embeddings of shape [num nodes, representation dim].
       node repesentations, edges, edge weights = inputs
       # Get node indices (source) and neighbour indices (target) from edges.
       node indices, neighbour indices = edges[0], edges[1]
       # neighbour repesentations shape is [num edges, representation dim].
       neighbour repesentations = tf.gather(node repesentations, neighbour indices)
       # Prepare the messages of the neighbours.
       neighbour messages = self.prepare(neighbour repesentations, edge weights)
       # Aggregate the neighbour messages.
       aggregated messages = self.aggregate(
           node indices, neighbour messages, node repesentations
       # Update the node embedding with the neighbour messages.
       return self.update(node repesentations, aggregated messages)
```

Graph neural network node classifier

- The GNN classification model proceeds, as follows:
 - 1. Apply preprocessing using FFN to the node features to generate initial node representations.
 - 2. Apply one or more graph convolutional layer, with skip connections, to the node representation to produce node embeddings.
 - 3. Apply post-processing using FFN to the node embeddings to generat the final node embeddings.
 - 4. Feed the node embeddings in a Softmax layer to predict the node class.
- Each graph convolutional layer added captures information from a further level of neighbours. However, adding many graph convolutional layer can cause oversmoothing, where the model produces similar embeddings for all the nodes.
- Note that the graph_info passed to the constructor of the Keras model, and used as a property of the Keras model object, rather than input data for training or prediction. The model will accept a batch of node_indices, which are used to lookup the node features and neighbours from the graph_info.

Class GNNNodeClassifier

```
class GNNNodeClassifier(tf.keras.Model):
   def init (
        self,
       graph info,
       num classes,
       hidden units,
        aggregation type="sum",
        combination type="concat",
        dropout rate=0.2,
        normalize=True,
        *args,
        **kwarqs,
    ):
        super(GNNNodeClassifier, self). init (*args, **kwargs)
        # Unpack graph info to three elements: node features, edges, and edge weight.
        node features, edges, edge weights = graph info
        self.node features = node features
        self.edges = edges
        self.edge weights = edge weights
        # Set edge weights to ones if not provided.
        if self.edge weights is None:
            self.edge weights = tf.ones(shape=edges.shape[1])
        # Scale edge weights to sum to 1.
        self.edge weights = self.edge weights / tf.math.reduce sum(self.edge weights)
```

Class GNNNodeClassifier

```
# Create a process layer.
       self.preprocess = create ffn(hidden units, dropout rate, name="preprocess")
       # Create the first GraphConv layer.
       self.conv1 = GraphConvLayer(
           hidden units,
           dropout rate,
           aggregation type,
           combination type,
           normalize,
           name="graph conv1",
       # Create the second GraphConv layer.
       self.conv2 = GraphConvLayer(
           hidden units,
           dropout rate,
           aggregation type,
           combination type,
           normalize,
           name="graph conv2",
       # Create a postprocess layer.
       self.postprocess = create ffn(hidden units, dropout rate, name="postprocess")
       # Create a compute logits layer.
       self.compute logits = layers.Dense(units=num classes, name="logits")
```

Class GNNNodeClassifier

```
def call(self, input node indices):
       # Preprocess the node features to produce node representations.
       x = self.preprocess(self.node features)
       # Apply the first graph conv layer.
       x1 = self.conv1((x, self.edges, self.edge weights))
       # Skip connection.
       x = x1 + x
       # Apply the second graph conv layer.
       x2 = self.conv2((x, self.edges, self.edge weights))
       # Skip connection.
       x = x2 + x
       # Postprocess node embedding.
       x = self.postprocess(x)
       # Fetch node embeddings for the input node indices.
       node embeddings = tf.gather(x, input node indices)
       # Compute logits
       return self.compute logits(node embeddings)
```

Instantiate GNNNodeClassifier

• Let's test instantiating and calling the GNN model. Notice that if you provide N node indices, the output will be a tensor of shape [N, num_classes], regardless of the size of the graph.

```
gnn_model = GNNNodeClassifier(
    graph_info=graph_info,
    num_classes=num_classes,
    hidden_units=hidden_units,
    dropout_rate=dropout_rate,
    name="gnn_model",
)
print("GNN output shape:", gnn_model([1, 10, 100]))
gnn_model.summary()
```

gnn model.summary()

```
GNN output shape: tf.Tensor(
\lceil \lceil -0.08258238 -0.23063104 \ 0.05941907 \ 0.15656857 -0.07656706 -0.15783894 
  -0.107356191
[-0.1807271 \quad -0.26390904 \quad 0.00740898 \quad 0.15194565 \quad -0.11337246 \quad -0.15723836
 -0.165755121
[-0.05183263 \quad 0.10546789 \quad -0.09391725 \quad -0.15574552 \quad -0.01967955 \quad 0.06629369
  -0.14769788]], shape=(3, 7), dtype=float32)
Model: "qnn model"
Layer (type)
                             Output Shape
______
preprocess (Sequential) (2708, 32)
                                                        52804
 graph conv1 (GraphConvLayer multiple
                                                        5888
 graph conv2 (GraphConvLayer multiple
                                                        5888
postprocess (Sequential) (2708, 32)
                                                        2368
 logits (Dense)
                             multiple
                                                        231
Total params: 67,179
Trainable params: 63,481
Non-trainable params: 3,698
```

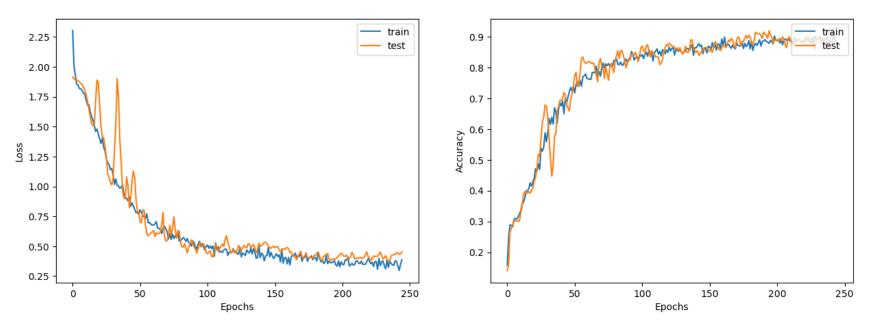
Train the GNN model

Note that we use the standard supervised cross-entropy loss to train the model.
However, we can add another self-supervised loss term for the generated node
embeddings that makes sure that neighbouring nodes in graph have similar
representations, while faraway nodes have dissimilar representations.

```
x train = train data.paper id.to numpy()
history = run experiment(gnn model, x train, y train)
Epoch 1/300
0.1581 - val loss: 1.9164 - val acc: 0.1407
Epoch 2/300
0.2469 - val loss: 1.9004 - val acc: 0.1608
Epoch 3/300
0.2886 - val loss: 1.8916 - val acc: 0.2663
. . . .
Epoch 244/300
0.8845 - val loss: 0.4381 - val acc: 0.8894
Epoch 245/300
0.8845 - val loss: 0.4511 - val acc: 0.8794
```

The learning curves

display learning curves (history)



 Now we evaluate the GNN model on the test data split. The results may vary depending on the training sample, however the GNN model always outperforms the baseline model in terms of the test accuracy.

```
x_test = test_data.paper_id.to_numpy()
_, test_accuracy = gnn_model.evaluate(x=x_test, y=y_test, verbose=0)
print(f"Test accuracy: {round(test accuracy * 100, 2)}%")
```

Test accuracy: 76.36%

GNN model predictions

 Let's add the new instances as nodes to the node_features, and generate links (citations) to existing nodes.

```
# First we add the N new instances as nodes to the graph
# by appending the new instance to node features.
num nodes = node features.shape[0]
new node features = np.concatenate([node features, new instances])
# Second we add the M edges (citations) from each new node to a set
# of existing nodes in a particular subject
new node indices = [i + num nodes for i in range(num classes)]
new citations = []
for subject idx, group in papers.groupby("subject"):
    subject papers = list(group.paper id)
    # Select random x papers specific subject.
    selected paper indices1 = np.random.choice(subject papers, 5)
    \# Select random y papers from any subject (where y < x).
    selected paper indices2 = np.random.choice(list(papers.paper id), 2)
    # Merge the selected paper indices.
    selected paper indices = np.concatenate(
        [selected paper indices1, selected paper indices2], axis=0
    # Create edges between a citing paper idx and the selected cited papers.
    citing paper indx = new node indices[subject idx]
    for cited paper idx in selected paper indices:
       new citations.append([citing paper indx, cited paper idx])
new citations = np.array(new citations).T
new edges = np.concatenate([edges, new citations], axis=1)
```

Update the node features and the edges

• Now let's update the node features and the edges in the GNN model.

```
print("Original node features shape:", gnn model.node features.shape)
print("Original edges shape:", gnn model.edges.shape)
gnn model.node features = new node features
gnn model.edges = new edges
gnn model.edge weights = tf.ones(shape=new edges.shape[1])
print("New node features shape:", gnn model.node features.shape)
print("New edges shape:", gnn model.edges.shape)
logits = gnn model.predict(tf.convert to tensor(new node indices))
probabilities = keras.activations.softmax(tf.convert to tensor(logits)).numpy()
display class probabilities (probabilities)
Original node_features shape: (2708, 1433)
Original edges shape: (2, 5429)
New node_features shape: (2715, 1433)
New edges shape: (2, 5478)
1/1 [=======] - 1s 1s/step
Instance 1:
- Case Based: 58.17%
- Genetic_Algorithms: 19.66%
- Neural Networks: 3.94%
- Probabilistic_Methods: 0.85%
- Reinforcement_Learning: 3.0%
- Rule_Learning: 3.86%
- Theory: 10.52%
```

Display class probabilities

Instance 2:

- Case_Based: 0.49%
- Genetic_Algorithms: 97.4%
- Neural_Networks: 0.74%
- Probabilistic_Methods: 0.06%
- Reinforcement_Learning: 0.83%
- Rule_Learning: 0.01%
- Theory: 0.47%

Instance 3:

- Case Based: 0.17%
- Genetic_Algorithms: 1.37%
- Neural_Networks: 51.64%
- Probabilistic_Methods: 45.65%
- Reinforcement_Learning: 0.15%
- Rule_Learning: 0.04%
- Theory: 0.97%

Instance 4:

- Case_Based: 0.04%
- Genetic Algorithms: 0.87%
- Neural Networks: 87.44%
- Probabilistic_Methods: 10.25%
- Reinforcement_Learning: 0.24%
- Rule_Learning: 0.03%
- Theory: 1.13%

Instance 5:

- Case_Based: 0.05%
- Genetic_Algorithms: 99.08%
- Neural_Networks: 0.13%
- Probabilistic_Methods: 0.01%
- Reinforcement_Learning: 0.71%
- Rule_Learning: 0.0%
- Theory: 0.02%

Instance 6:

- Case Based: 1.88%
- Genetic_Algorithms: 2.49%
- Neural Networks: 42.68%
- Probabilistic_Methods: 0.59%
- Reinforcement_Learning: 33.32%
- Rule_Learning: 10.92%
- Theory: 8.12%

Instance 7:

- Case_Based: 0.39%
- Genetic_Algorithms: 54.4%
- Neural Networks: 5.32%
- Probabilistic Methods:
- 17.31%
- Reinforcement_Learning: 0.39%
- Rule_Learning: 0.09%
- Theory: 22.11%

Notice that the probabilities of the expected subjects (to which several citations are added) are higher compared to the baseline model implying that Graph modeling of citation problem exposes more information about the data.