

Machine Learning on Graphs

COMP9312_23T2



UNSW
SYDNEY

Outline

- Machine Learning on Graphs
- Node Feature Engineering

Data Structure & Algorithms

Case studies on Community Detection:

Connected Component, K-Core, K-Truss, Clique, ...

Clustering/partition algorithms, ...

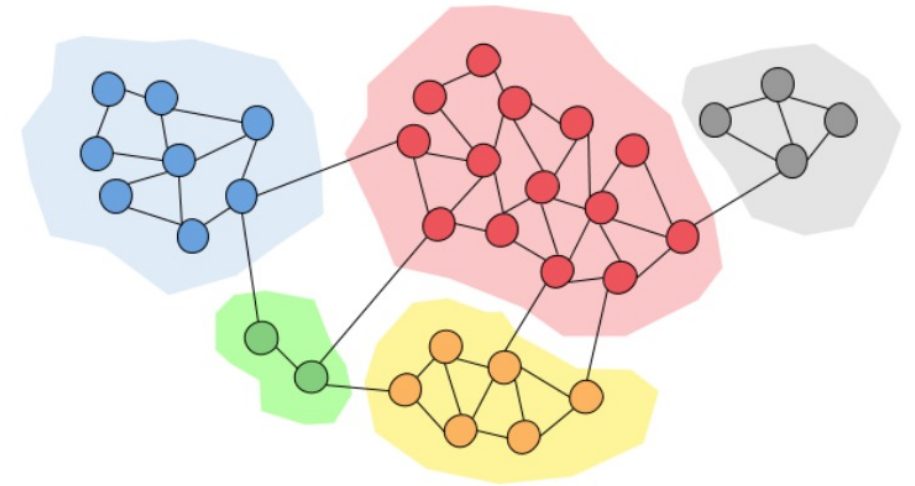
Step 1:
Formulate
problem



Step 2:
Design
Algorithm

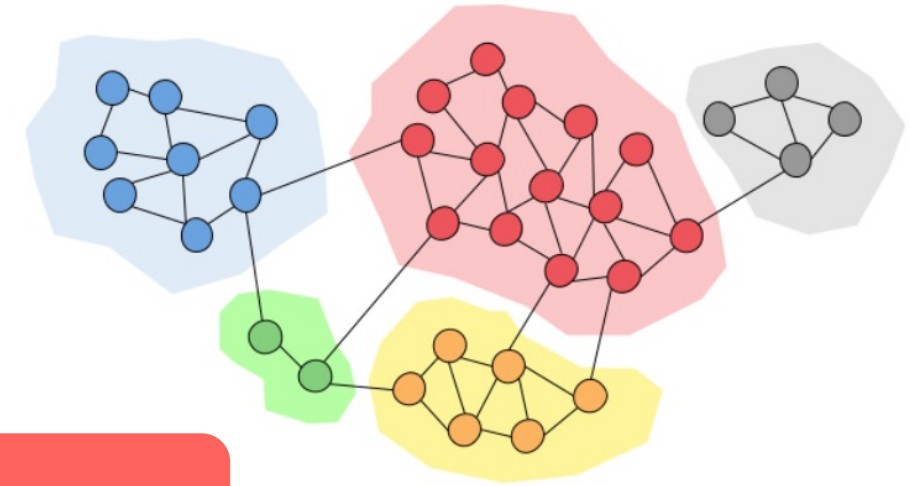


It is hard in many applications.



Learning-based Algorithms

- It is hard to define a good community.
- It is not hard to judge a community.



Step 1:
Decide
parameters



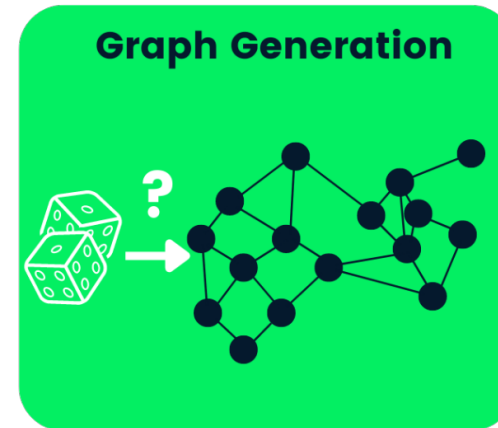
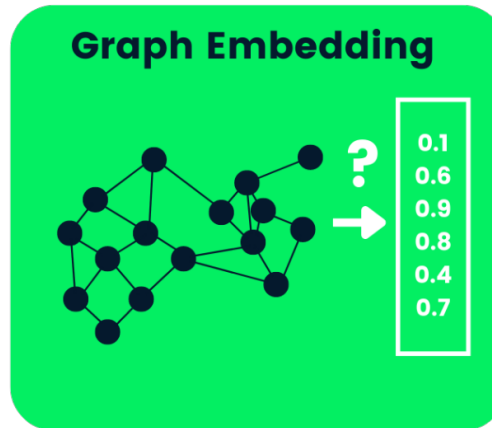
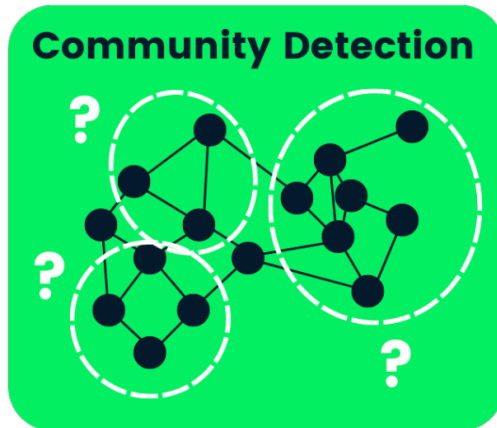
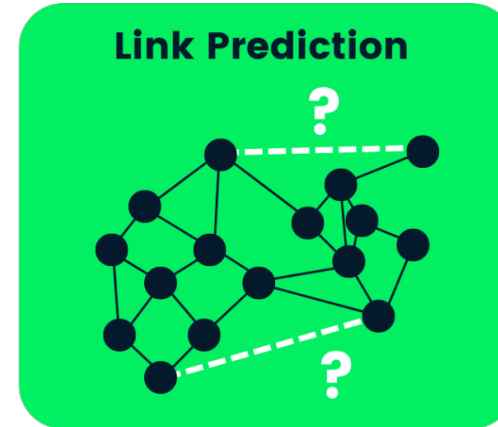
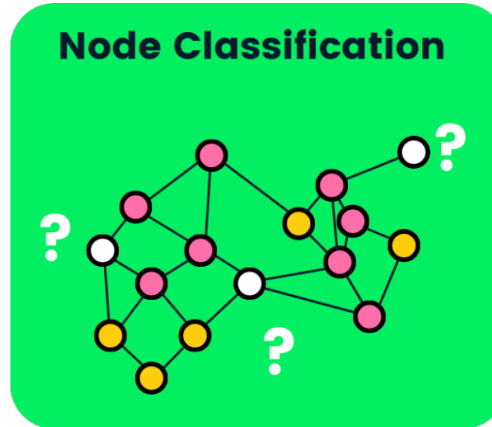
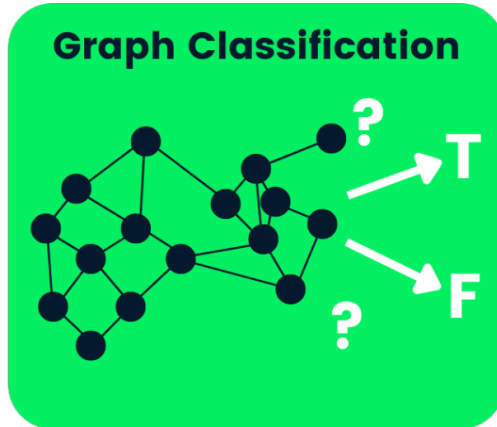
Step 2:
Define loss
function



Step 3:
Training

Efficiency VS Effectiveness

Application



<https://www.datacamp.com/tutorial/comprehensive-introduction-graph-neural-networks-gnns-tutorial>

Application

Node classification: Predict a property of a node

Example: Categorize online users / items

Link prediction: recommendation

Example: Knowledge graph completion

Graph classification: Categorize different graphs

Example: Molecule property prediction

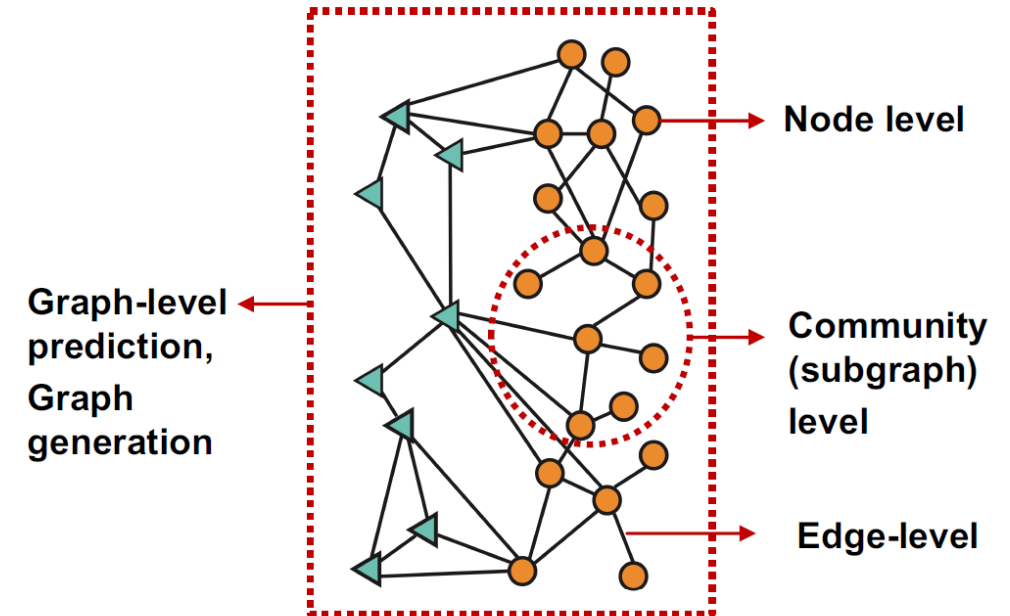
Clustering: Detect if nodes form a community

Example: Social circle detection

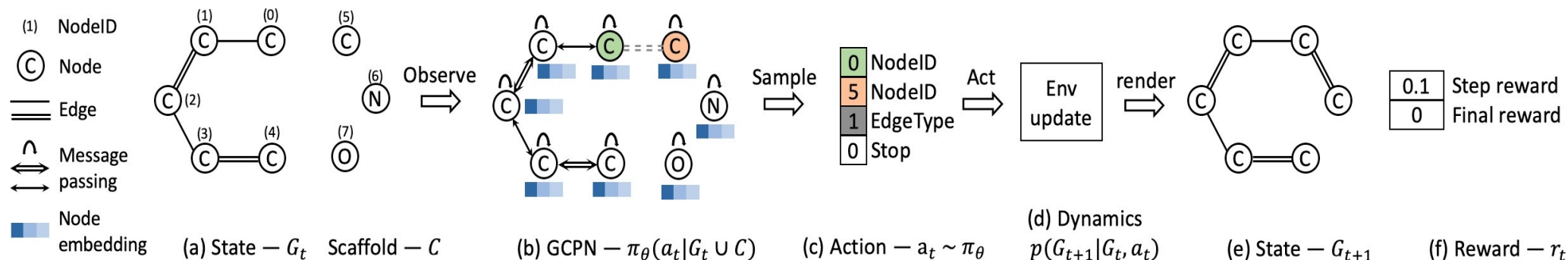
Other tasks:

Graph generation: Drug discovery

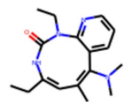
Graph evolution: Physical simulation



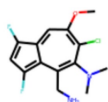
Application: Molecule Generation



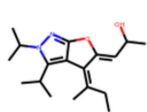
Use case 1: Generate novel molecules with high drug likeness



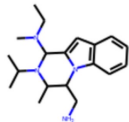
0.948



0.945

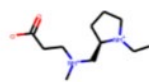


0.944

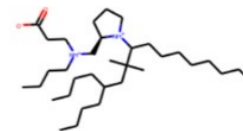


0.941

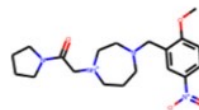
Use case 2: Optimize existing molecules to have desirable properties



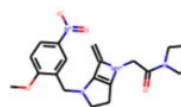
-8.32



-0.71



-5.55

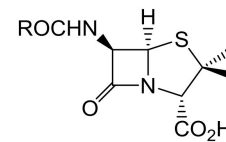


-1.78

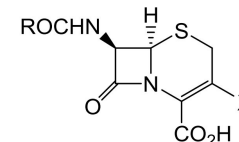
Application: Drug Discovery

Antibiotics are small molecular graphs

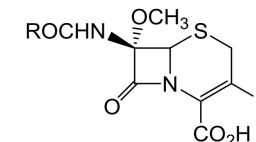
- Nodes: Atoms
- Edges: Chemical bonds



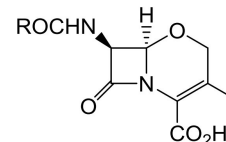
penicillins



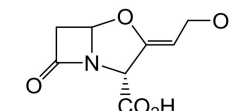
cephalosporins



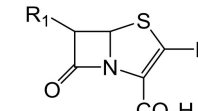
cephamycins



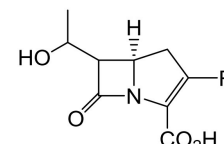
oxacephems



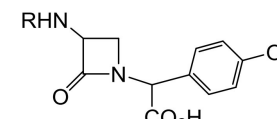
clavulanic acid
(an oxapenem)



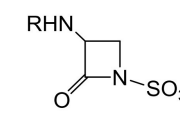
penems



carbapenems



nocardicin



monobactams

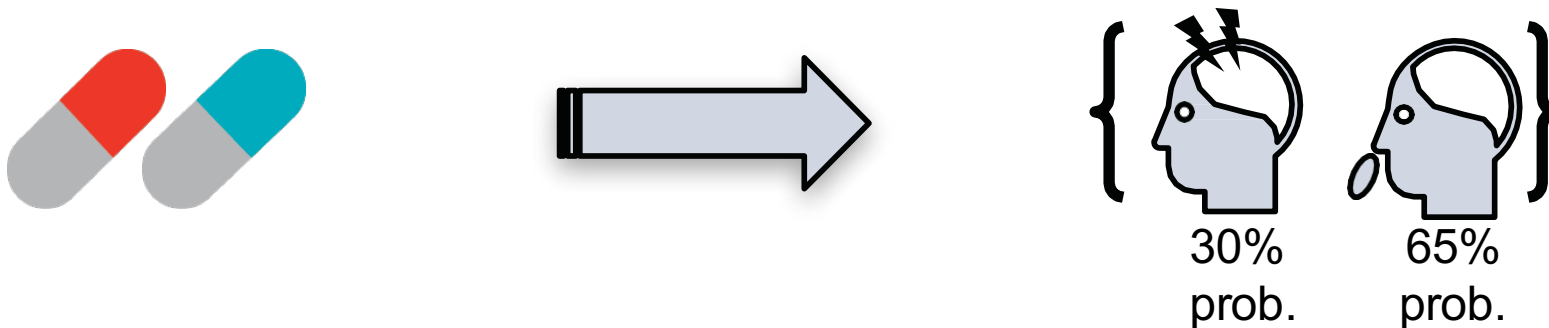
Konaklieva, Monika I. "Molecular targets of β -lactam-based antimicrobials: beyond the usual suspects." *Antibiotics* 3.2 (2014): 128-142.

Application: Drug Side Effects

Many patients take multiple drugs to treat complex or co-existing diseases:

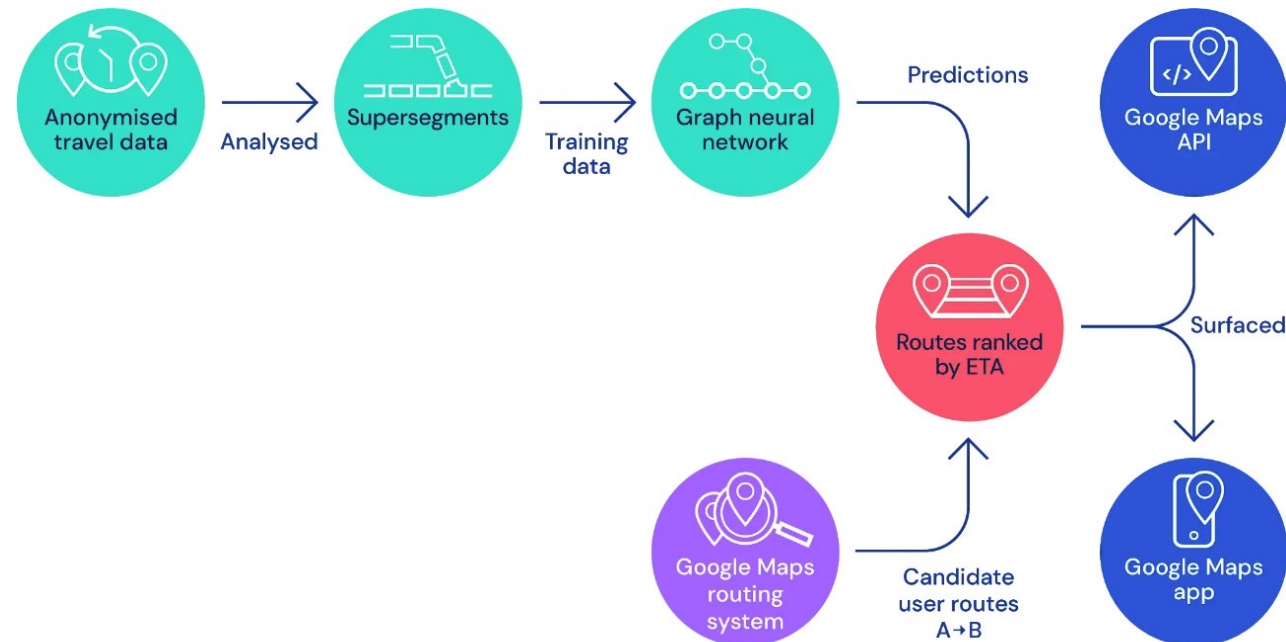
- 46% of people ages 70-79 take more than 5 drugs
- Many patients take more than 20 drugs to treat heart disease, depression, insomnia, etc.

Task: Given a pair of drugs predict adverse side effects



Application: Google Map

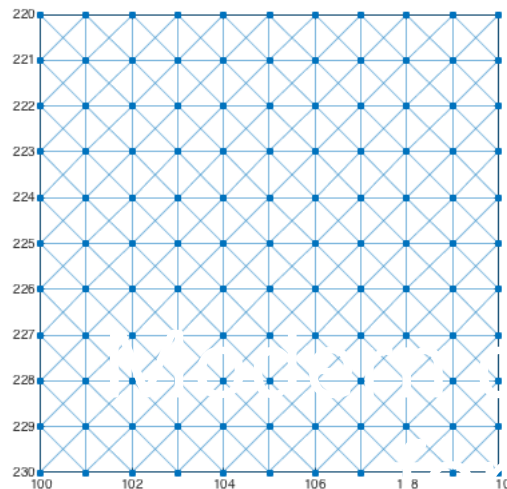
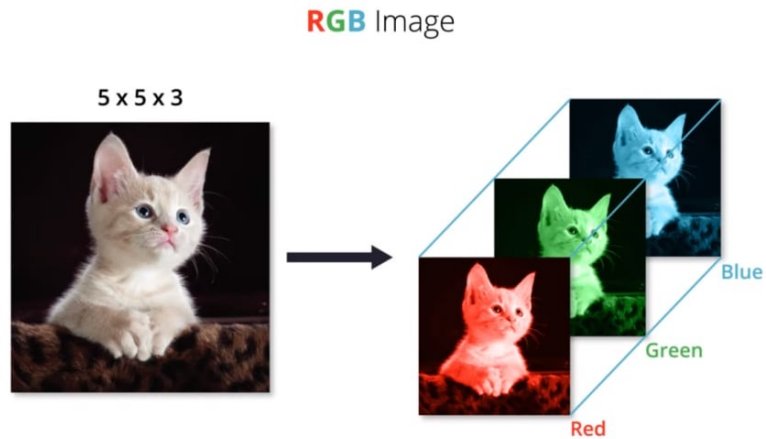
Predict via Graph Neural Networks



THE MODEL ARCHITECTURE FOR DETERMINING OPTIMAL ROUTES AND THEIR TRAVEL TIME.

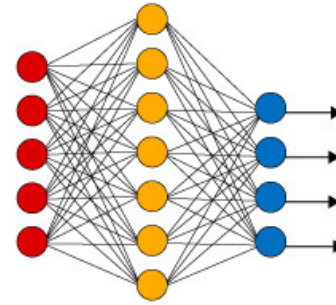
Image credit: [DeepMind](#)

ML/DL on traditional data

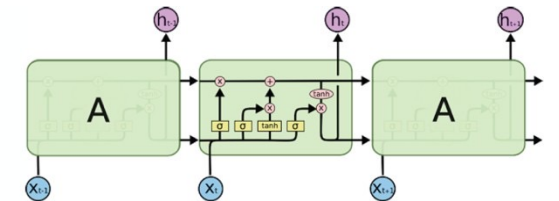
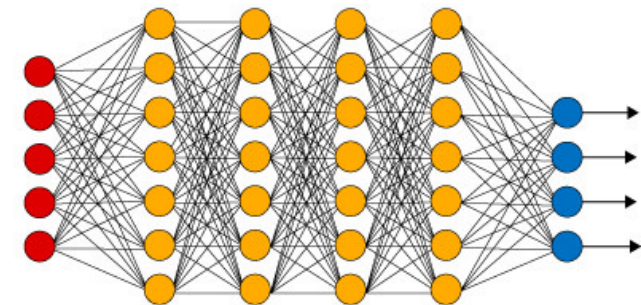


Images

Simple Neural Network



Deep Learning Neural Network

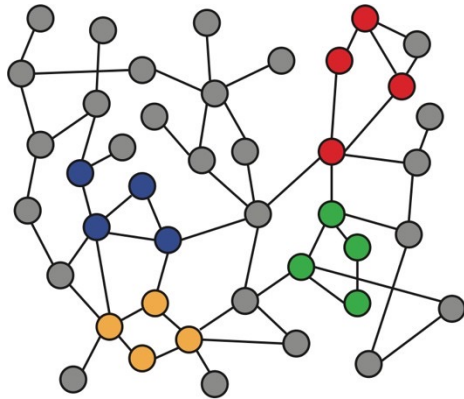


Text/Speech/Audio

Challenges

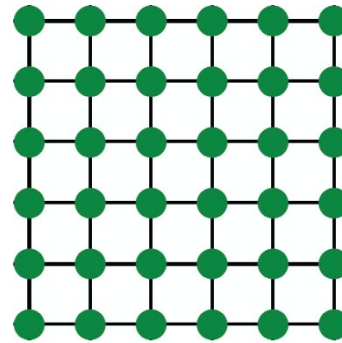
Graphs are complex

- Arbitrary size and complex topological structure (*i.e.*, no spatial locality like grids)
- No fixed node ordering or reference point
- Often dynamic and have multimodal features



Graph

VS

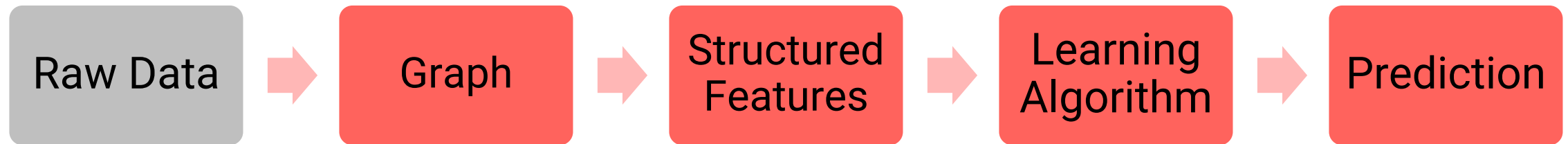


Image



Text

Graph Neural Networks



Adjacency Matrix? Adjacency List? CSR?

Machine learning needs features!

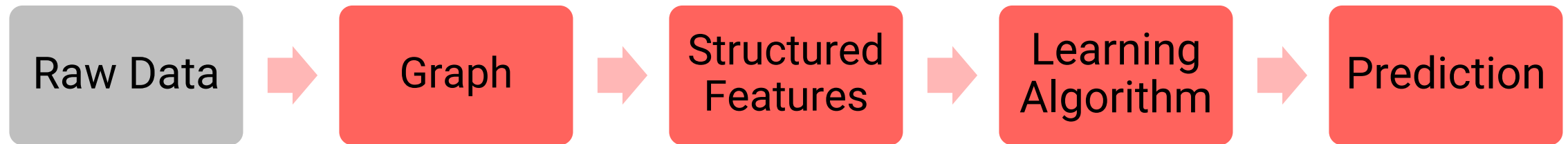
How to get features

1. Feature Engineering

Covered in this topic

2. Graph Representation Learning

Optional topic of node embedding

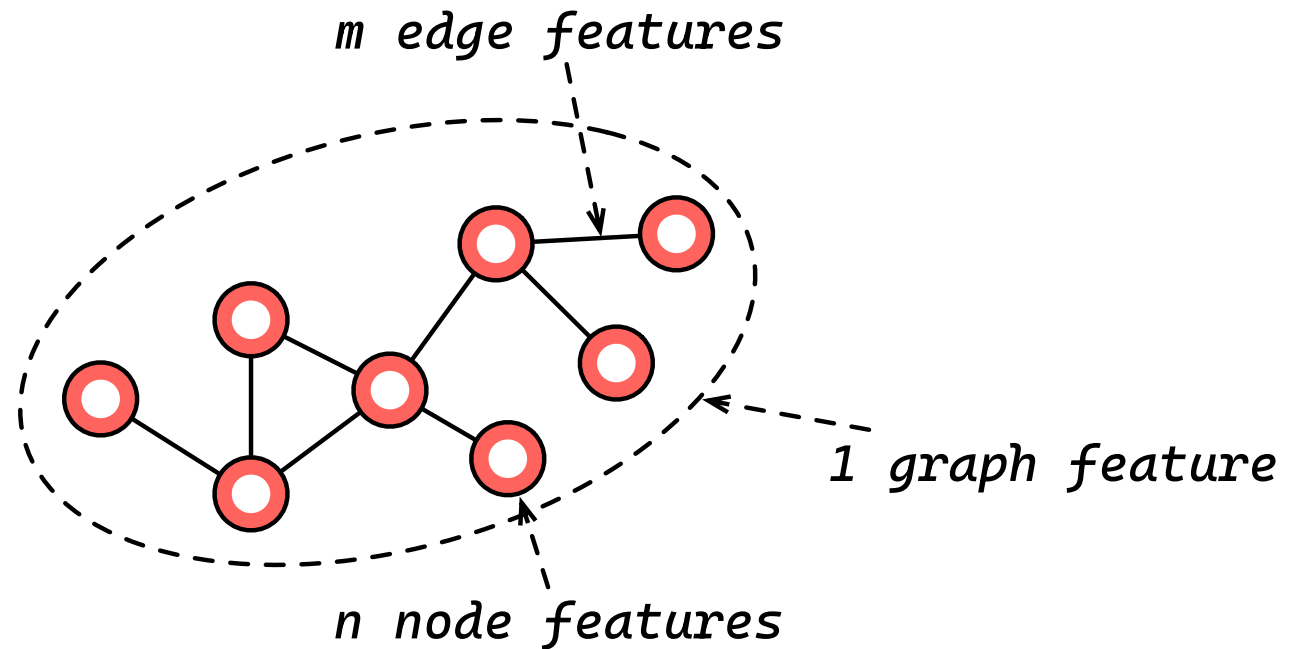


Adjacency Matrix? Adjacency List? CSR?

Machine learning needs features!

Different types of graph features

- Node Level
- Edge Level
- Graph Level



Traditional ML on Graphs

Good features effectively represent the graph structure and achieve good performance.

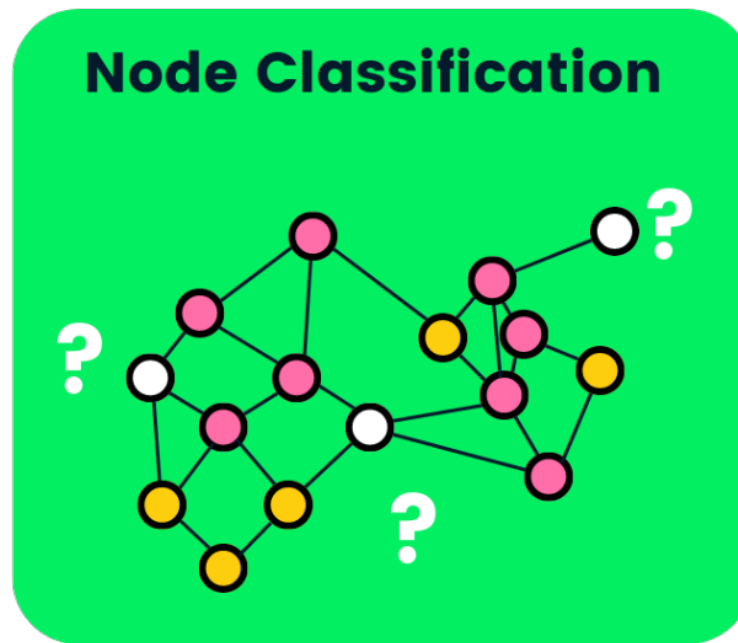
1. Design features for **nodes**/edges/graphs.
2. Get features additional features from training data.
3. Use features to train parameters.

Testing: predict using the feature of query node/link/graph

Node-Level Features

Goal:

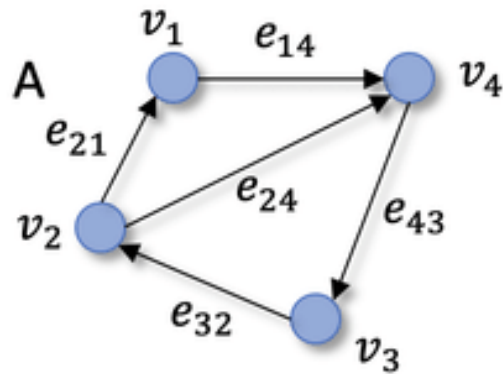
Characterize the structure and position of a node in the network:



A typical application: node classification

Adjacency Matrix?

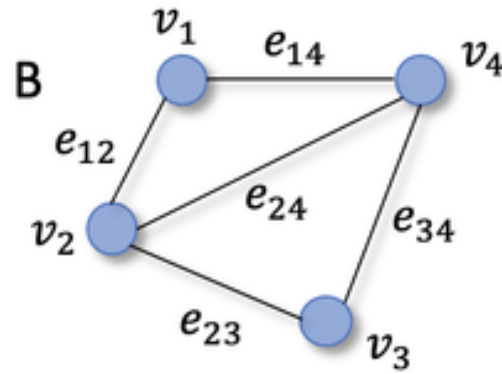
Directed graph $G(V,E)$



E

	v_1	v_2	v_3	v_4
v_1	0	0	0	1
v_2	1	0	0	1
v_3	0	1	0	0
v_4	0	0	1	0

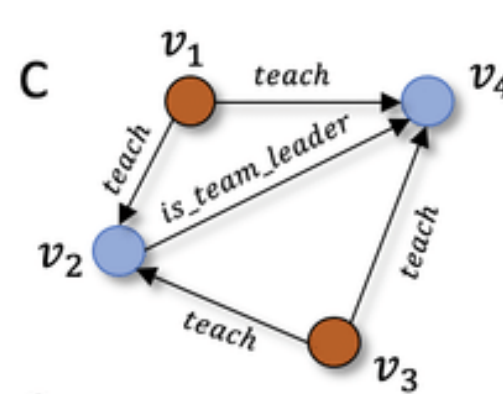
Undirected graph $G(V,E)$



F

	v_1	v_2	v_3	v_4
v_1	0	1	0	1
v_2	1	0	1	1
v_3	0	1	0	1
v_4	1	1	1	0

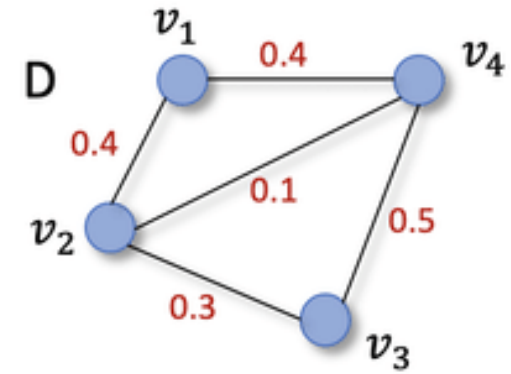
Knowledge graph $G(V,E)$



G

	v_1	v_2	v_3	v_4
v_1	0	1	0	1
v_2	0	0	0	1
v_3	0	1	0	1
v_4	0	0	0	0

Weighted graph $G(V,E)$

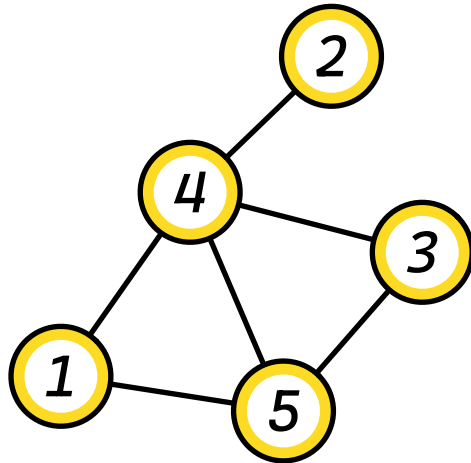


H

	v_1	v_2	v_3	v_4
v_1	0	0.4	0	0.4
v_2	0.4	0	0.3	0.1
v_3	0	0.3	0	0.5
v_4	0.4	0.1	0.5	0

Not working for big graphs!

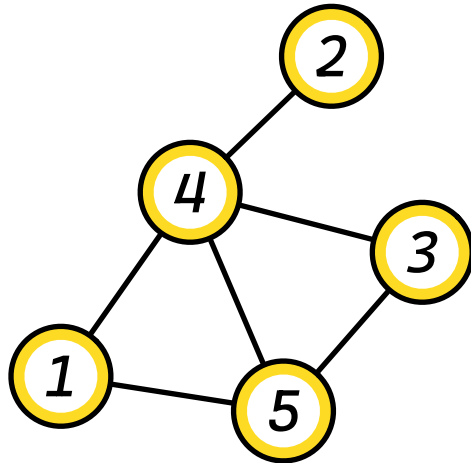
Adjacency List?



<i>v1</i>	4	5		
<i>v2</i>	4			
<i>v3</i>	4	5		
<i>v4</i>	1	2	3	5
<i>v5</i>	1	3	4	

Feature dimension need to be consistent

Adjacency List?



How about this?

<i>v1</i>	<i>4</i>	<i>5</i>	<i>0</i>	<i>0</i>
<i>v2</i>	<i>4</i>	<i>0</i>	<i>0</i>	<i>0</i>
<i>v3</i>	<i>4</i>	<i>5</i>	<i>0</i>	<i>0</i>
<i>v4</i>	<i>1</i>	<i>2</i>	<i>3</i>	<i>5</i>
<i>v5</i>	<i>1</i>	<i>3</i>	<i>4</i>	<i>0</i>

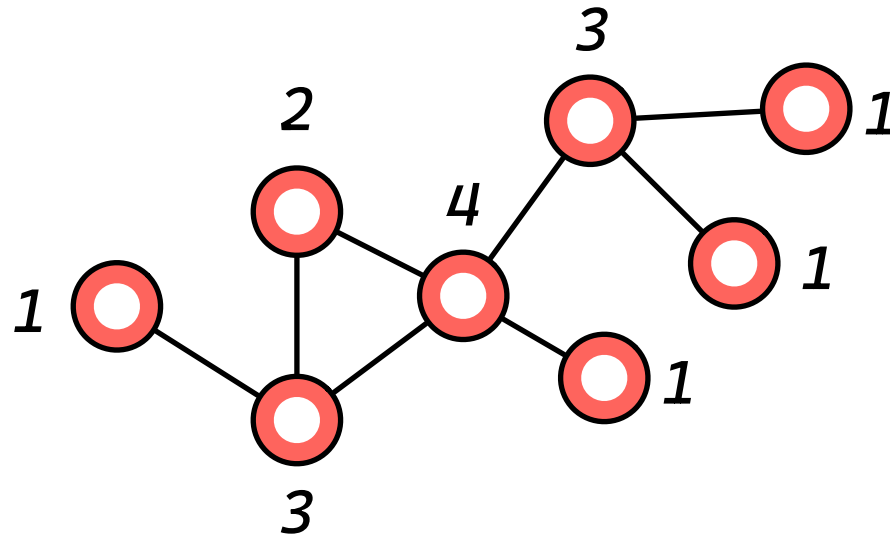
Feature dimension need to be consistent

Node-Level Features: Overview

- Node degree
- Clustering coefficient
- Graphlets
- Node centrality
- ...

Node Degree

Degree of a node: the number of neighbors.
Treat all neighbors equally.

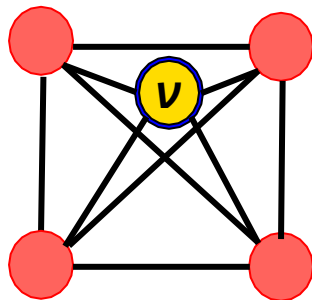


Node Centrality: Clustering Coefficient

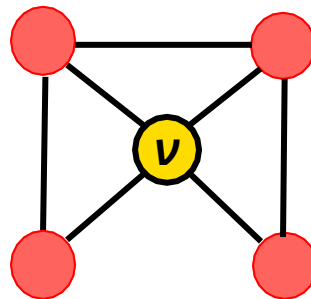
Measures how connected v 's neighboring nodes are:

$$e_v = \frac{\#(\text{edges among neighboring nodes})}{\binom{k_v}{2}} \in [0,1]$$

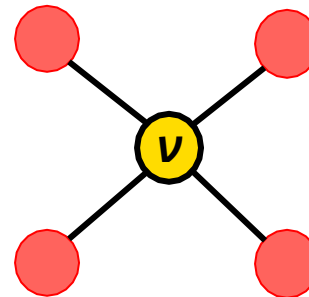
Can be also understand as #triangles/#possible triangles



$$e_v = 1$$



$$e_v = 0.5$$



$$e_v = 0$$

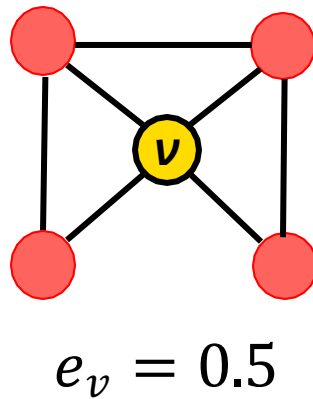
Ego-network:
the induced subgraph
of the node and all its
neighbors

Computing Clustering Coefficient

Can you design an algorithm to compute the clustering coefficient of all nodes in a graph with n nodes and m edges?

Node Features: Graphlets

Observation: Clustering coefficient counts the #(triangles) in the ego-network.



Three triangles in 6 possible triplets

We can generalize the above by counting #(pre-specified subgraphs, i.e., **graphlets**).

Node Features: Graphlets

Graphlets are small subgraphs.

We aim to describe network structure **around** the node based on graphlets.

Analogy: **Degree**

counts **#(edges)** that a node touches.

Clustering coefficient

counts **#(triangles)** that a node is involved.

Graphlet Degree Vector (GDV):

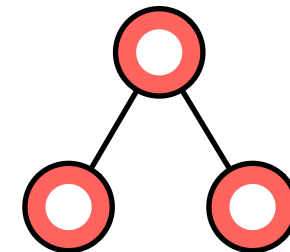
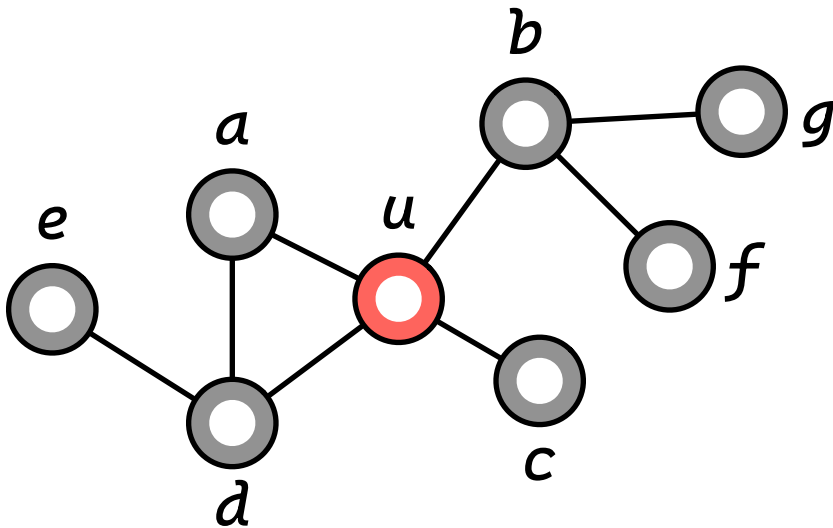
Graphlet-base features for nodes

GDV counts **#(graphlets)** that a node is involved.

Node Features: Graphlets

How to represent a node by graphlets?

Let's start by considering (connected) graphlets with three nodes:

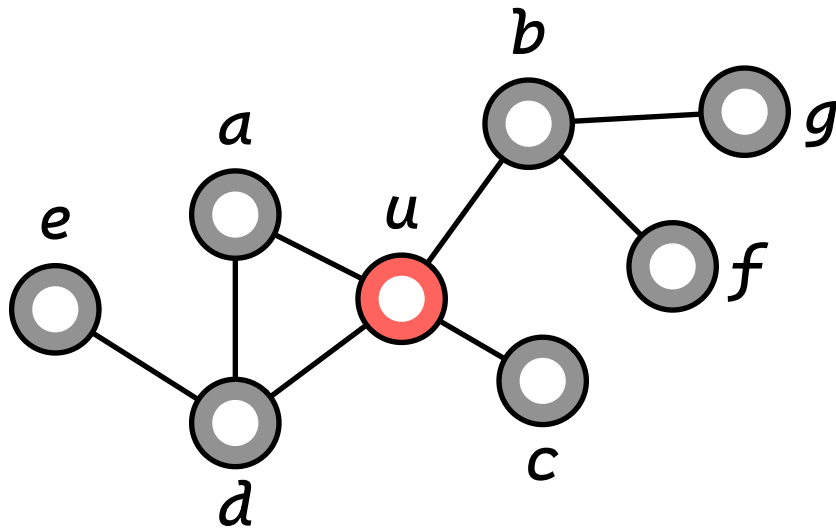


Choose a specific pattern (wedge)

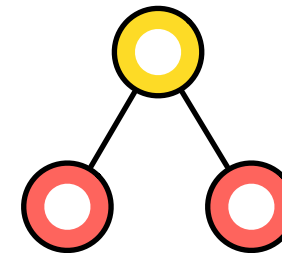
How many subgraphs containing u that are isomorphic to the pattern?

Node Features: Graphlets

How many subgraphs containing u that are isomorphic to the pattern?



11 after removing symmetric cases



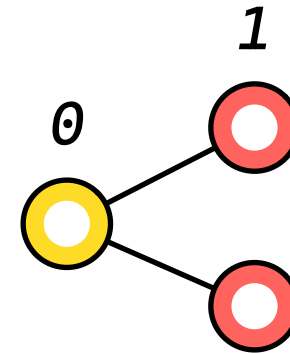
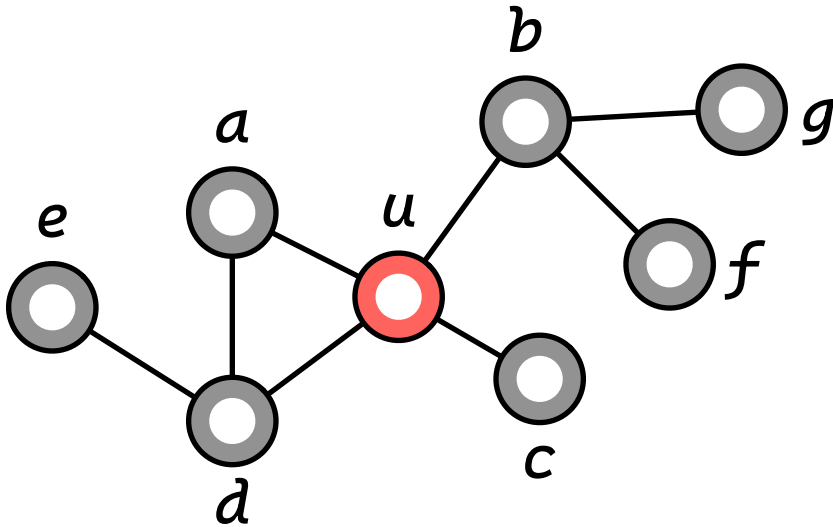
Choose a specific pattern (wedge)

We use **11** as the feature of u

Node Features: Graphlets

Move forward by utilizing different types:

6 for type 0
5 for type 1

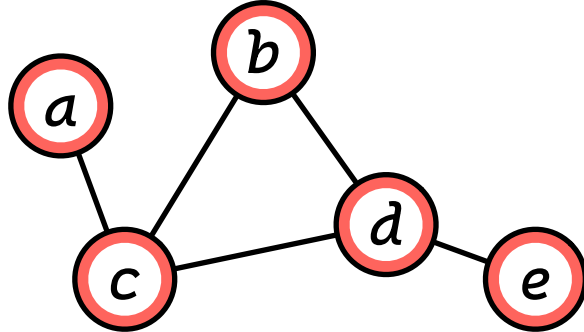


Choose a specific pattern (wedge)

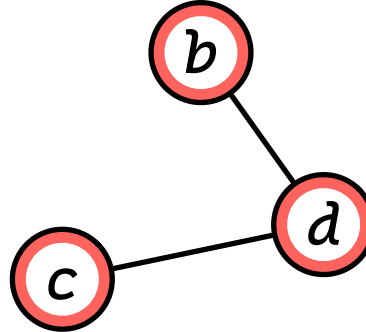
We use $[6, 5]$ as the feature of u

Node Features: Graphlets

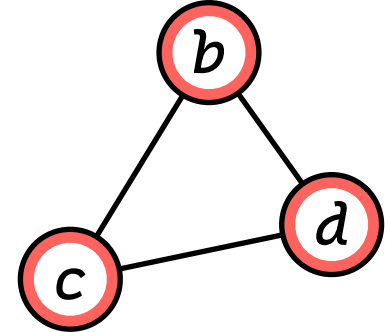
Move forward by only considering induced matching instances:



A graph



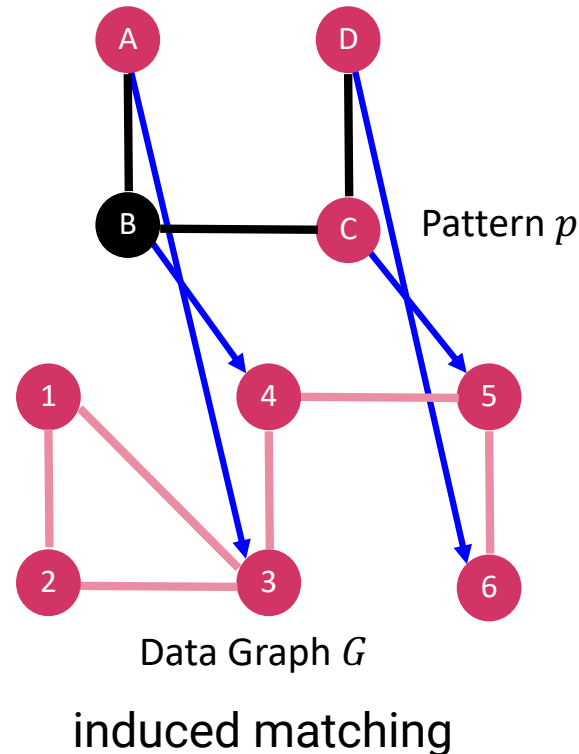
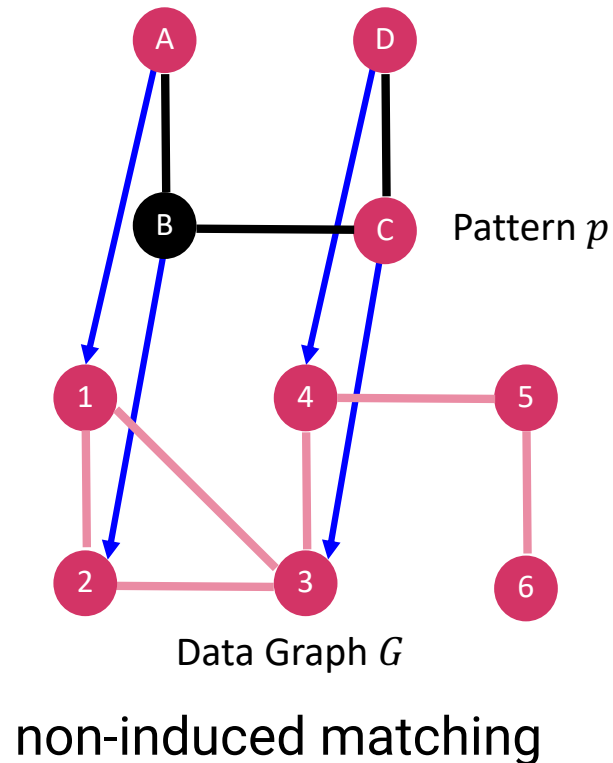
A non-induced subgraph



An induced subgraph

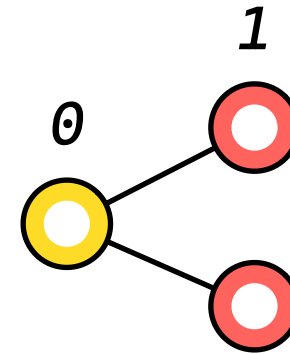
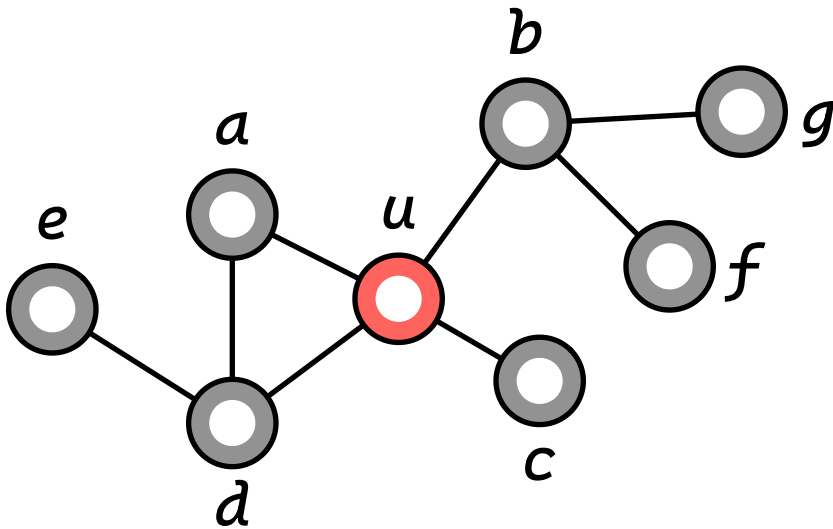
Node Features: Graphlets

Move forward by only considering induced matching instances:



Node Features: Graphlets

Move forward by only considering **induced** matching instances:



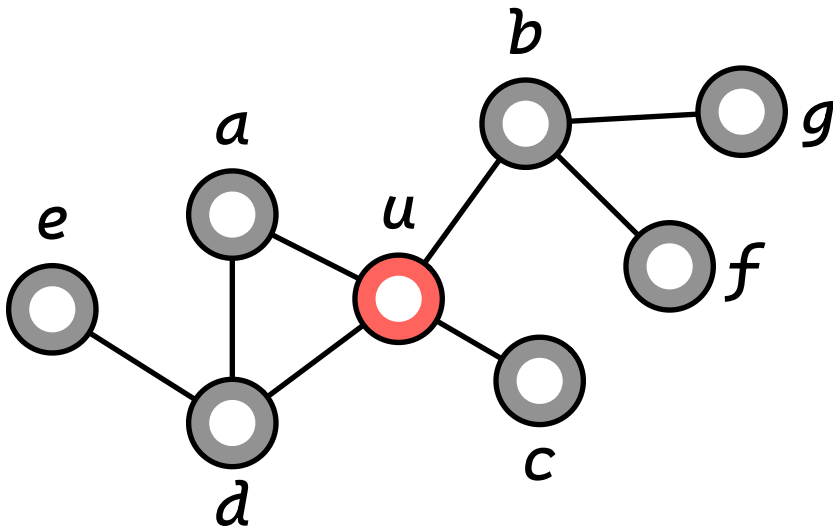
Choose a specific pattern (wedge)

We use **$[5, 3]$** as the feature of u

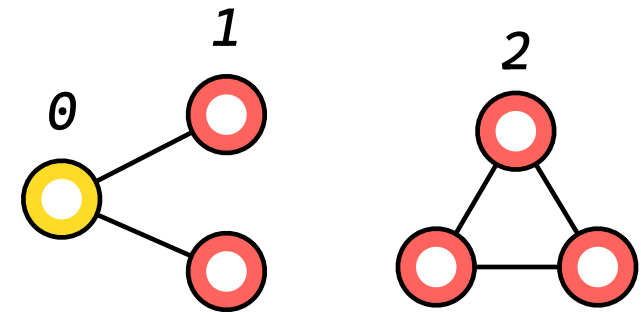
Node Features: Graphlets

Move forward by utilizing all 3-graphlets:

6 for type 0
5 for type 1
1 for type 2



Type 0:



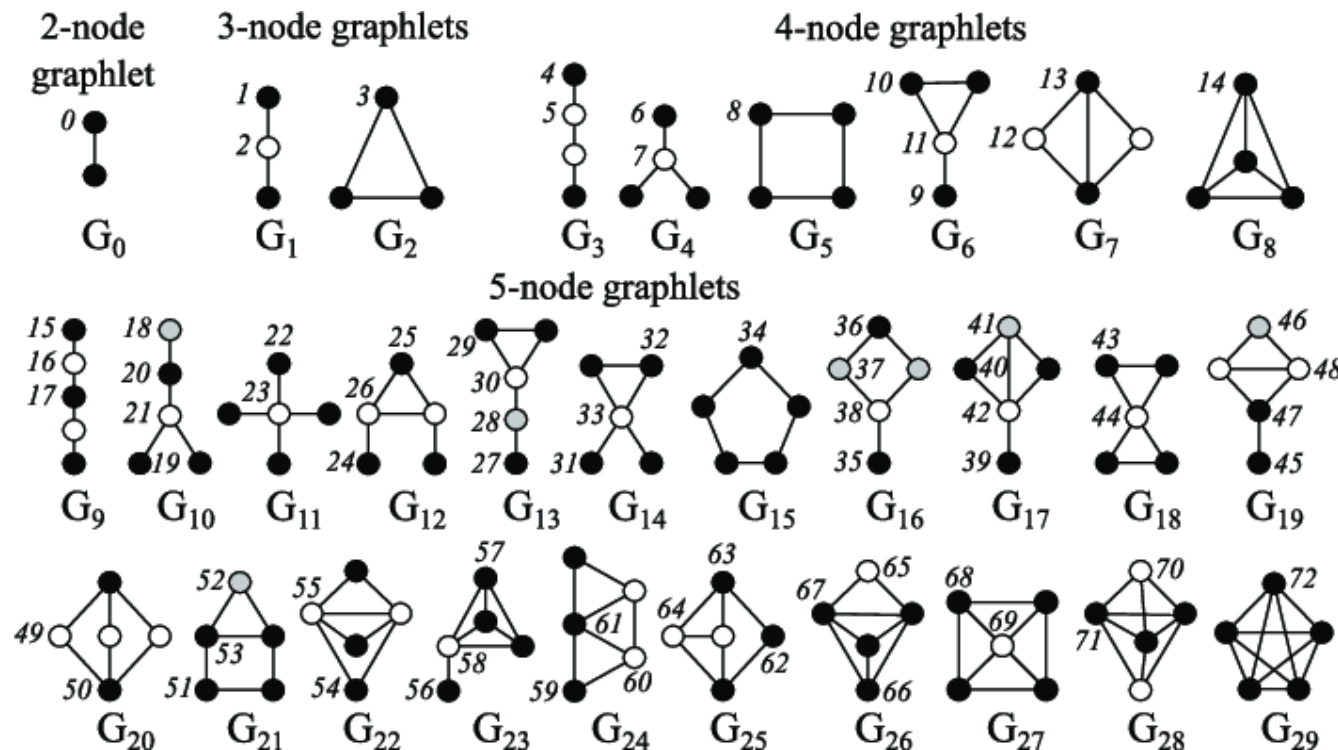
There are three types in all 3-graphlets.

We use $[5, 3, 1]$ as the feature of u

Node Features: Graphlets

Consider all graphlets with ≤ 5 nodes

How many node roles in all connected non-isomorphic subgraphs?



There are **73** different graphlets of up to 5 nodes.

To get the node feature, compute the number of induced matching instances for each role id.

Node Features: Graphlets

Graphlet Degree Vector (GDV): A count vector of graphlets rooted at a given node.

Considering graphlets of size 2-5 nodes we get:

- **Vector of 73 coordinates** is a signature of a node that describes the topology of node's neighborhood

Graphlet degree vector provides a measure of a **node's local network topology**:

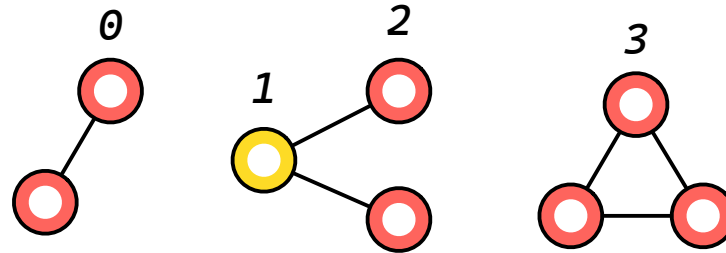
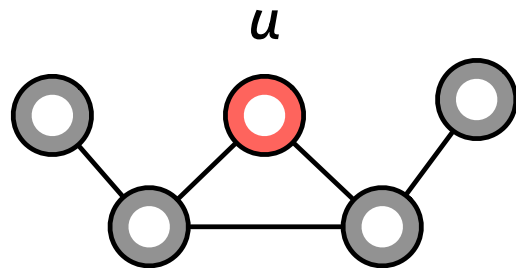
- Comparing vectors of two nodes provides a more detailed measure of local topological similarity than node degrees or clustering coefficient.

Usually, we only compute up to 4 or 5 nodes . . .

Node Features: Graphlets

More examples:

$$GFV(u) = [2, 0, 2, 1]$$

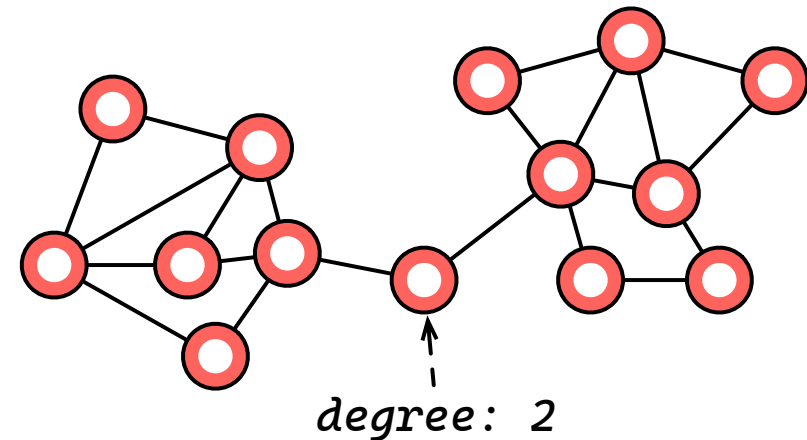


Node Centrality

Node degree counts neighbors **without capturing their importance**.
Node **centrality** takes the **node importance in a graph** into account

Different ways to model importance:

- PageRank
- Eigenvector centrality
- Betweenness centrality
- Closeness centrality
- many others...



Node Centrality: Eigenvector

Motivation

A node is important if **surrounded by important neighbors**.

We model the centrality of node v as **the sum of the centrality of neighbors**:

$$c_v = \frac{1}{\lambda} \sum_{u \in N(v)} c_u$$

λ is normalization constant
(it will turn out to be the largest eigenvalue of A)

The above equation models centrality in a **recursive manner**.
How do we solve it?

Node Centrality: Eigenvector

Optional

Math Warning!

Rewrite the recursive equation in the matrix form.

$$c_v = \frac{1}{\lambda} \sum_{u \in N(v)} c_u \longleftrightarrow \lambda \mathbf{c} = \mathbf{A} \mathbf{c}$$

λ is normalization const
(largest eigenvalue of A)

- A : Adjacency matrix
 $A_{uv} = 1$ if $u \in N(v)$
- \mathbf{c} : Centrality vector
- λ : Eigenvalue

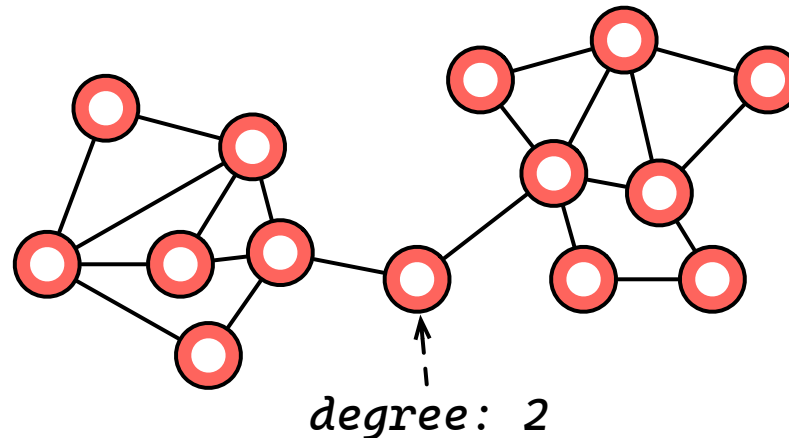
- We see that centrality \mathbf{c} is the **eigenvector of A** !
- The largest eigenvalue λ_{max} is always positive and unique (by Perron-Frobenius Theorem).
- The eigenvector \mathbf{c}_{max} corresponding to λ_{max} is used for centrality.

Node Centrality: Betweenness

Betweenness centrality:

A node is important if it lies on many shortest paths between other nodes.

$$c_v = \sum_{s \neq v \neq t} \frac{\#(\text{shortest paths between } s \text{ and } t \text{ that contain } v)}{\#(\text{shortest paths between } s \text{ and } t)}$$

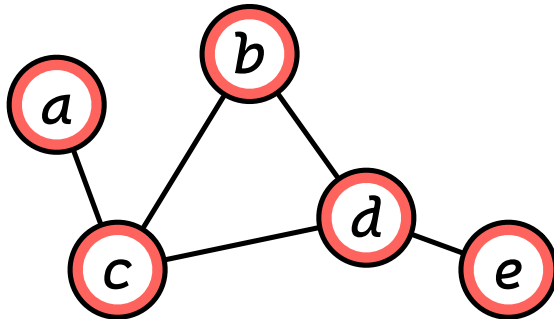


How to identify the bridge node

Node Centrality: Betweenness (cont)

Example:

$$c_v = \sum_{s \neq v \neq t} \frac{\#(\text{shortest paths between } s \text{ and } t \text{ that contain } v)}{\#(\text{shortest paths between } s \text{ and } t)}$$



$$c_a = c_b = c_e = 0$$

$$c_c = 3$$

$(a-\underline{c}-b, a-\underline{c}-d, a-\underline{c}-d-e)$

$$c_d = 3$$

$(a-c-\underline{d}-e, b-\underline{d}-e, c-\underline{d}-e)$

Computing Betweenness Centrality

Exact solution:

$O(nm)$ for unweighted graphs

$O(nm+n^2\log n)$ for weighted graphs

<https://kops.uni-konstanz.de/server/api/core/bitstreams/420590d1-3010-4eab-a585-6fa3eff46f9e/content>

Approximate solution:

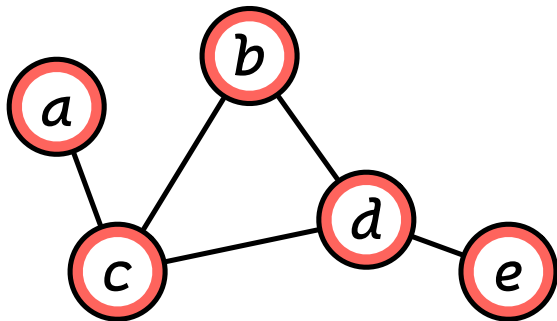
Sampling a set of shortest paths...

Node Centrality: Closeness

Closeness centrality:

A node is important if it lies on many shortest paths between other nodes.

$$c_v = \frac{1}{\sum_{u \neq v} \text{shortest path length between } u \text{ and } v}$$



$$c_a = 1/(2 + 1 + 2 + 3) = 1/8$$

(a-c-b, a-c, a-c-d, a-c-d-e)

$$c_d = 1/(2 + 1 + 1 + 1) = 1/5$$

(d-c-a, d-b, d-c, d-e)

Computing Closeness Centrality

Can you design an algorithm to compute the closeness centrality of all nodes in a graph with n nodes and m edges?

Node-Level Feature: Summary

- Importance-based features:
 - Node degree
 - Different node centrality measures
- Structure-based features:
 - Node degree
 - Clustering coefficient
 - Graphlet count vector

Node-level Feature: Summary

Importance-based features: capture the importance of a node in a graph

- ~~Node degree:~~
 - Simply counts the number of neighboring nodes
- Node centrality:
 - Model **importance of neighbors** in a graph
 - Different modeling choices: eigenvector centrality, betweenness centrality, closeness centrality

Useful for predicting influential nodes in a graph

- **Example:** predicting celebrity users in a social network

Node-level Feature: Summary

Structure-based features: Capture topological properties of local neighborhood around a node.

- **Node degree:**
 - Counts the number of neighboring nodes
- **Clustering coefficient:**
 - Measures how connected neighboring nodes are
- **Graphlet degree vector:**
 - Counts the occurrences of different graphlets

Useful for predicting a particular role a node plays in a graph:

- **Example:** Predicting protein functionality in a protein-protein interaction network.

Learning Outcome

- **Traditional ML Pipeline**
 - Hand-crafted feature + ML model
- **Hand-crafted node features for graph data**
 - Node degree, centrality, clustering coefficient, graphlets