Graph Learning

IFT6758 - Data Science

Sources:

http://snap.stanford.edu/proj/embeddings-www/

https://jian-tang.com/files/AAAI19/aaai-grltutorial-part2-gnns.pdf





Announcements

- Mid-term exam grades will be published on Gradescope today.
- Assignment 4, final presentation, group and individual reports will be published on Gradescope on Wednesday.
- Presentation format is similar to mid-term, i.e., 7 Min for presentation (all teammembers should present to get a score)

BUT:

- 3 Min questions about the presentation
- 5 Min coding questions from all team-members
- You should ONLY present the model that you will submit on December 2.

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Graphs are everywhere



Graphs are a general language for describing and modeling complex systems

Graph G = (V, E)





Graphs are everywhere



Social networks



Economic networks



Biomedical networks



• Graphs can have labels on their edges and/or nodes



Labels can also be considered weights







Road maps

Protein interation networks





• Labels don't have to be numerical, they can be textual.



• Labels don't have to be unique; it's entirely possible and sometimes useful to give multiple nodes the same label.



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• Graphs can have features (a.k.a attributes).













Graphs can be either:

- Heterogeneous composed of different types of nodes
- **Homogeneous** composed of the same type of nodes

and are either:

• Static — nodes and edges do not change, nothing is added or taken away

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• **Dynamic** — nodes and edges change, added, deleted, moved, etc.



Why are graphs useful?

- This is a very **flexible data structure** that generalizes many other data structures. For example, if there are no edges, then it becomes a set; if there are only "vertical" edges and any two nodes are connected by exactly one path, then we have a tree.
- Nodes and edges typically come from some **expert knowledge** or intuition about the problem.

e.g., Atoms in molecules, Users in a social network, Cities in a transportation system, Players in team sport, Neurons in the brain, Interacting objects in a dynamic physical system, and Pixels, bounding boxes or segmentation masks in images





Why are graphs useful?

• Most ML/CV problems can be viewed as graphs



from (Antonakos et al., CVPR, 2015)

- Graph gives a lot of flexibility and can give a very different and interesting perspective on the problem
- Neural networks can be viewed as graph where nodes are neutrons and weights are edges





Graph mining tasks

- Classical ML tasks in graphs:
 - Node classification: Predict a type of a given node
 - Link prediction: Predict whether two nodes are linked
 - Community detection: Identify densely linked clusters of nodes
 - Network similarity: How similar are two (sub)networks





Node Classification







Node Classification



Image from: Ganapathiraju et al. 2016. <u>Schizophrenia interactome with 504 novel</u> protein-protein interactions. *Nature*.





Link Prediction







Link Prediction

Content recommendation is link prediction!









Community Detection

 The field of community detection aims to identify highly connected groups of individuals or objects inside these networks, these groups are called communities.





Network Similarity





Graph Basics





Traverse a graph



- Walk: A graph traversal a closed walk is when the destination node is the same as the source node
- Trail: A walk with no repeated edges a circuit is a closed trail
- Path: A walk with no repeated nodes a cycle is a closed path



Adjacency Matrix

 The Adjacency Matrix of a graph is be made of 1s and 0s unless it is otherwise weighted or labelled. A can be built by following this rule:

$$a_{ij} = \begin{cases} 1, \text{ if } (v_i, v_j) \in E \\ 0, \text{ otherwise} \end{cases}$$

The Adjacency Matrix of a undirected graph is therefore symmetrical along its diagonal





Degree Matrix

• The Degree Matrix **D** of a graph is essentially a diagonal matrix, where each value of the diagonal is the degree of its corresponding node.





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Laplacian Matrix

• The Laplacian Matrix of a graph is the result of subtracting the Adjacency Matrix from the Degree Matrix:

L = D - A

• Each value in the Degree Matrix is subtracted by its respective value in the Adjacency Matrix as such:

Labeled graph	Degree matrix						Adjacency matrix							Laplacian matrix						
\bigcirc	(2	0	0	0	0	0)		(0	1	0	0	1	0)	1	2	-1	0	0	$^{-1}$	0)
Θ	0	3	0	0	0	0		1	0	1	0	1	0		$^{-1}$	3	-1	0	$^{-1}$	0
(4)	0	0	2	0	0	0		0	1	0	1	0	0		0	-1	2	-1	0	0
IL	0	0	0	3	0	0		0	0	1	0	1	1		0	0	-1	3	-1	-1
(3)-(2)	0	0	0	0	3	0		1	1	0	1	0	0		$^{-1}$	-1	0	-1	3	0
	0/	0	0	0	0	1/		0	0	0	1	0	0/		0	0	0	$^{-1}$	0	1/



Graph Learning





Traditional Machine Learning Pipeline

• (Supervised) Machine Learning Lifecycle





Feature Learning in Graphs

• **Goal**: Efficient task-independent feature learning for machine learning in graphs!







Representation





* slide from Thomas Kipf, University of Amsterdam

Why graph learning is hard?



Standard machine learning/deep learning approaches don't work on this data!





Why Graph learning is hard?

Modern deep learning toolbox is designed for simple sequences or grids.

CNNs for fixed-size images/grids....



RNNs or word2vec for text/sequences...





recap: Isomorphism problem

• The graph isomorphism problem is the computational problem of determining whether two finite graphs are isomorphic.



• The graph isomorphism problem is neither known to be NPcomplete nor known to be tractable



Why Graph learning is hard?

- Graphs are far **more complex** than text or visual data!
- Complex topographical structure (i.e., no spatial locality like grids)
- No fixed node ordering or reference point (i.e., the isomorphism problem)
- Often dynamic and have multimodal features.





Node Embedding





Node Embedding



 Intuition: Find embedding of nodes to d-dimensions so that "similar" nodes in the graph have embeddings that are close together.





Node Embedding (Set up)

- Assume we have a graph G:
 - V is the vertex set.
 - A is the adjacency matrix (assume binary).
 - No node features or extra information is used!


Node Embedding (Set up)

- Assume we have a graph G:
 - V is the vertex set.
 - A is the adjacency matrix (assume binary).
 - No node features or extra information is used!
- Goal is to encode nodes so that similarity in the embedding space (e.g., dot product) approximates similarity in the original network.



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Learning Node Embeddings

- 1. Define an encoder (i.e., a mapping from nodes to embeddings)
- 2. Define a node similarity function (i.e., a measure of similarity in the original network).
- 3. Optimize the parameters of the encoder so that:

similarity
$$(u, v) \approx \mathbf{z}_v^\top \mathbf{z}_u$$



Key Components

• Encoder maps each node to a low-dimensional vector.

$$\operatorname{ENC}(v) = \mathbf{z}_{v} \quad \text{embedding}$$

Similarity function specifies how relationships in vector space map to relationships in the original network.

node in the input graph

similarity
$$(u, v) \approx \mathbf{z}_v^\top \mathbf{z}_u$$

Similarity of u and v in the original network

dot product between node embeddings

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Shallow Encoding

 Simplest encoding approach: encoder is just an embeddinglookup

$$\operatorname{ENC}(v) = \mathbf{Zv}$$

 $\mathbf{Z} \in \mathbb{R}^{d imes |\mathcal{V}|}$ matrix, each column is node embedding [what we learn!]

 $\mathbf{v} \in \mathbb{I}^{|\mathcal{V}|} \quad \text{indicator vector, all zeroes except a one in column indicating node } v$



Shallow Encoding

 Simplest encoding approach: encoder is just an embeddinglookup



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From Shallow to Deep

- Limitations of shallow encoding:
 - O(|V|) parameters are needed: there no parameter sharing and every node has its own unique embedding vector.
 - Inherently "transductive": It is impossible to generate embeddings for nodes that were not seen during training.
 - Do not incorporate node features: Many graphs have features that we can and should leverage.



From Shallow to Deep

 We will now discuss "deeper" methods based on graph neural networks.

 $ENC(v) = \frac{complex function that}{depends on graph structure.}$

 In general, all of encoders can be combined with the similarity functions that depends on graph structure.





How to Define Node Similarity?

- Key distinction between "shallow" methods is how they define node similarity.
- E.g., should two nodes have similar embeddings if they....
 - are connected?
 - share neighbors?
 - have similar "structural roles"?
 - ...?



How to Define Node Similarity?

- 1. Adjacency-based similarity
- 2. Multi-hop similarity
- 3. Random walk approaches

High-level structure and material from:

• <u>Hamilton et al. 2017</u>. Representation Learning on Graphs: Methods and Applications. *IEEE Data Engineering Bulletin on Graph Systems*.



Material based on:

• Ahmed et al. 2013. Distributed Natural Large Scale Graph Factorization. WWW.





- Similarity function is just the edge weight between u and v in the original network.
- Intuition: Dot products between node embeddings approximate edge existence.





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$$\mathcal{L} = \sum_{(u,v)\in V\times V} \|\mathbf{z}_u^{\top}\mathbf{z}_v - \mathbf{A}_{u,v}\|^2$$

- Find embedding matrix that minimizes the loss
 - Option 1: Use stochastic gradient descent (SGD) as a general optimization method.
 - Highly scalable, general approach
 - Option 2: Solve matrix decomposition solvers (e.g., SVD).
 - Only works in limited cases.



Recap: SVD for Word embedding





Recap: SVD for Node embedding





Recap: Node Similarity is preserved



• Given an Adjacency matrix C, we can get a decomposition C' from SVD.

$$\hat{X}\hat{X}^{T} = (U\Sigma V^{T})(U\Sigma V^{T})^{T}$$
$$= (U\Sigma V^{T})(V\Sigma U^{T})$$
$$= U\Sigma \Sigma^{T} U^{T} \quad (\because V^{T} V = I)$$
$$= U\Sigma (U\Sigma)^{T}$$



Matrix factorization

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 Matrix factorization algorithms work by decomposing the useritem interaction matrix into the product of two lower dimensionality rectangular matrices.



• This family of methods became widely known during the Netflix prize challenge 2006 due to its effectiveness in recommender systems..





$$\mathcal{L} = \sum_{(u,v)\in V\times V} \|\mathbf{z}_u^{\top}\mathbf{z}_v - \mathbf{A}_{u,v}\|^2$$

- Drawbacks:
 - O(IVI²) runtime. (Must consider all node pairs.)
 - Can make O([EI) by only summing over non-zero edges and using regularization (e.g., <u>Ahmed et al., 2013</u>)
 - O(IVI) parameters! (One learned vector per node).
 - Only considers direct, local connections.



e.g., the blue node is obviously more similar to green compared to red node, despite none having direct connections.

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Material based on:

- Cao et al. 2015. <u>GraRep: Learning Graph Representations with Global Structural</u> Information. CIKM.
- Ou et al. 2016. Asymmetric Transitivity Preserving Graph Embedding. KDD.
- Jian Tang, Meng Qu, Mingzhe Wang, Jun Yan, Ming Zhang and Qiaozhu Mei. LINE: Large-scale Information Network Embedding . WWW'15



- Idea: Consider k-hop node neighbors.
- E.g., two or three-hop neighbors.



- Red: Target node
- Green: 1-hop neighbors
 - A (i.e., adjacency matrix)
- Blue: 2-hop neighbors
 - A²
- Purple: 3-hop neighbors
 - A³



Power of djacency matrix

• If A^k gives us the number of walks from node i to node j after k steps.





Power of djacency matrix

• If A^k gives us the number of walks from node i to node j after k steps.





Basic idea:
$$\mathcal{L} = \sum_{(u,v) \in V \times V} \|\mathbf{z}_u^\top \mathbf{z}_v - \mathbf{A}_{u,v}^k\|^2$$

• Train embeddings to predict k-hop neighbors.





• Basic idea:
$$\mathcal{L} = \sum_{(u,v) \in V imes V} \|\mathbf{z}_u^\top \mathbf{z}_v - \mathbf{A}_{u,v}^k\|^2$$

- In practice (GraRep from Cao et al, 2015):
 - Use log-transformed, probabilistic adjacency matrix:

$$\begin{split} \tilde{\mathbf{A}}_{i,j}^k &= \max\left(\log\left(\frac{(\mathbf{A}_{i,j}/d_i)}{\sum_{l \in V}(\mathbf{A}_{l,j}/d_l)^k}\right)^k - \alpha, 0\right) \\ & \text{node degree} \end{split} \quad \text{constant shift} \end{split}$$

Train multiple different hop lengths and concatenate output.

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• Another option: Measure overlap between node neighborhoods.



• Example overlap functions:

• Jaccard similarity
$$J(A,B) = \frac{|A \cap B|}{|A \cup B|} = \frac{|A \cap B|}{|A| + |B| - |A \cap B|}$$

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- S_{u,v} is the neighborhood overlap between u and v (e.g., Jaccard overlap).
- This technique is known as HOPE (Yan et al., 2016).





Summary

- Basic idea so far:
 - 1) Define pairwise node similarities.
 - 2) Optimize low-dimensional embeddings to approximate these pairwise similarities.
- Issues:
 - Expensive: Generally O(IVI²), since we need to iterate over all pairs of nodes.
 - Brittle: Must hand-design deterministic node similarity measures.
 - Massive parameter space: O(IVI) parameters





Material based on:

- Perozzi et al. 2014. DeepWalk: Online Learning of Social Representations. KDD.
- Grover et al. 2016. node2vec: Scalable Feature Learning for Networks. KDD.





Transition Matrix

0	1	0
0	0	1
1	1	0

1	0
0	1
1/2	0
	1 0 1/2

Adjacency matrix A

Transition matrix P







Random Walk



Random Walk Embeddings

probability that u $\mathbf{z}_u^\top \mathbf{z}_v \approx \begin{array}{l} \text{and } \mathbf{v} \text{ co-occur on} \\ \text{a random walk over} \end{array}$ the network





recap: Skip-gram model







(recap) word2vec

- Node2vec is similar to word2vec skip-gram model.
- The same way as a document is an ordered sequence of words, one could sample sequences of nodes from the underlying network and turn a network into a ordered sequence of nodes.





(recap) word2vec



• word2vec can embed a very specific graphs:





Random Walk Embeddings

 Estimate probability of visiting node v on a random walk starting from node u using some random walk strategy R.

2. Optimize embeddings to encode these random walk statistics.







Why Random walks?

- Expressivity: Flexible stochastic definition of node similarity that incorporates both local and higher-order neighborhood information.
- Efficiency: Do not need to consider all node pairs when training; only need to consider pairs that co-occur on random walks.




- **1.** Run short random walks starting from each node on the graph using some strategy R.
- 2. For each node u collect $N_R(u)$, the multiset^{*} of nodes visited on random walks starting from **u**.
- 3. Optimize embeddings to according to:

$$\mathcal{L} = \sum_{u \in V} \sum_{v \in N_R(u)} -\log(P(v|\mathbf{z}_u))$$

* $N_R(u)$ can have repeat elements since nodes can be visited multiple times on random walks.



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$$\mathcal{L} = \sum_{u \in V} \sum_{v \in N_R(u)} -\log(P(v|\mathbf{z}_u))$$

 Intuition: Optimize embeddings to maximize likelihood of random walk co-occurrences.





$$\mathcal{L} = \sum_{u \in V} \sum_{v \in N_R(u)} -\log(P(v|\mathbf{z}_u))$$

- Intuition: Optimize embeddings to maximize likelihood of random walk co-occurrences.
- Parameterize $P(v | z_u)$ using softmax:

$$P(v|\mathbf{z}_u) = \frac{\exp(\mathbf{z}_u^{\top}\mathbf{z}_v)}{\sum_{n \in V} \exp(\mathbf{z}_u^{\top}\mathbf{z}_n)}$$



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• Putting things together:



• Optimizing random walk embeddings = Finding embeddings z_u that minimize \mathcal{L}





• But doing this naively is too expensive!!





• But doing this naively is too expensive!!

$$\mathcal{L} = \sum_{u \in V} \sum_{v \in N_R(u)} -\log\left(\frac{\exp(\mathbf{z}_u^{\top} \mathbf{z}_v)}{\sum_{n \in V} \exp(\mathbf{z}_u^{\top} \mathbf{z}_n)}\right)$$

The normalization term from the softmax can we approximate it?





• But doing this naively is too expensive!!

$$\mathcal{L} = \sum_{u \in V} \sum_{v \in N_R(u)} -\log\left(\frac{\exp(\mathbf{z}_u^{\top} \mathbf{z}_v)}{\sum_{n \in V} \exp(\mathbf{z}_u^{\top} \mathbf{z}_n)}\right)$$

The normalization term from the softmax can we approximate it?

Negative Sampling



How we should randomly walk?

- So far we have described how to optimize embeddings given random walk statistics.
- What strategies should we use to run these random walks?
 - Simplest idea: Just run **fixed-length**, unbiased random walks starting from each node (i.e., <u>DeepWalk from Perozzi et al., 2013</u>).
 - But can we do better?





node2vec: Biased Walks

 Idea: use flexible, biased random walks that can trade off between local and global views of the network (Grover and Leskovec, 2016).





node2vec: Biased Walks

 Two classic strategies to define a neighborhood of a given node :



 $N_{BFS}(u) = \{ s_1, s_2, s_3 \}$ Local microscopic view $N_{DFS}(u) = \{ s_4, s_5, s_6 \}$ Global macroscopic view



node2vec sampling strategy

- Node2vec's sampling strategy, accepts 4 arguments:
 - **Number of walks**: Number of random walks to be generated from each node in the graph
 - Walk length: How many nodes are in each random walk
 - **P**: Return hyperparameter
 - **q**: Inout hyperaprameter ("walk away" hyperaprameter)

and also the standard skip-gram parameters (context window size, number of iterations etc.)





Biased Random Walks

- Biased 2nd-order random walks explore network neighborhoods:
 - Random walk started at u and is now at w
 - Insight: Neighbors of w can only be:



Idea: Remember where that walk came from



Biased Random Walks

• Walker is at w. Where to go next?



1/p, 1/q, 1 are unnormalized probabilities

- p, q model transition probabilities
 - p ... return parameter
 - q ... "walk away" parameter





Biased Random Walks

• Walker is at w. Where to go next?



- BFS-like walk: Low value of p
- DFS-like walk: Low value of q

Ns(u) are the nodes visited by the walker

Unnormalized transition prob.

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BFS vs. DFS



BFS: Micro-view of neighbourhood





DFS: Macro-view of neighbourhood

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Experiments: Micro vs. Macro

• Interactions of characters in a novel:





p=1, q=2

Microscopic view of the network neighbourhood

p=1, q=0.5

Macroscopic view of the network neighbourhood





Other random walk ideas

- Different kinds of biased random walks:
 - Based on node attributes (Dong et al., 2017).
 - Based on a learned weights (<u>Abu-El-Haija et al., 2017</u>)
- Alternative optimization schemes:
 - Directly optimize based on 1-hop and 2-hop random walk probabilities (as in <u>LINE from Tang et al. 2015</u>).
- Network preprocessing techniques:
 - Run random walks on modified versions of the original network (e.g., <u>Ribeiro et al. 2017's struct2vec</u>, <u>Chen et al.</u> <u>2016's HARP</u>).





Summary

- Basic idea: Embed nodes so that distances in embedding space reflect node similarities in the original network.
- Different notions of node similarity:
 - Adjacency-based (i.e., similar if connected)
 - Multi-hop similarity definitions.
 - Random walk approaches.



So what method should I use ..?

- No one method wins in all cases....
 - e.g., node2vec performs better on node classification while multi-hop methods performs better on link prediction (Goyal and Ferrara, 2017 survey).
- Random walk approaches are generally more efficient (i.e., O(|E|) vs. O(|V|²))
- In general: Must choose a node similarity that matches application!





Thursday!

Graph Neural Networks & Graph Convolutional Networks



