

## Lecture 15:

# Graph Neural Networks (GNN)



Markus Hohle  
University California, Berkeley

Machine Learning Algorithms  
MSSE 277B, 3 Units



## Lecture 1: Course Overview and Introduction to Machine Learning

Lecture 2: Bayesian Methods in Machine Learning

classic ML tools & algorithms

Lecture 3: Dimensionality Reduction: Principal Component Analysis

Lecture 4: Linear and Non-linear Regression and Classification

Lecture 5: Unsupervised Learning: K-Means, GMM, Trees

Lecture 6: Adaptive Learning and Gradient Descent Optimization Algorithms

Lecture 7: Introduction to Artificial Neural Networks - The Perceptron

ANNs/AI/Deep Learning

Lecture 8: Introduction to Artificial Neural Networks - Building Multiple Dense Layers

Lecture 9: Convolutional Neural Networks (CNNs) - Part I

Lecture 10: CNNs - Part II

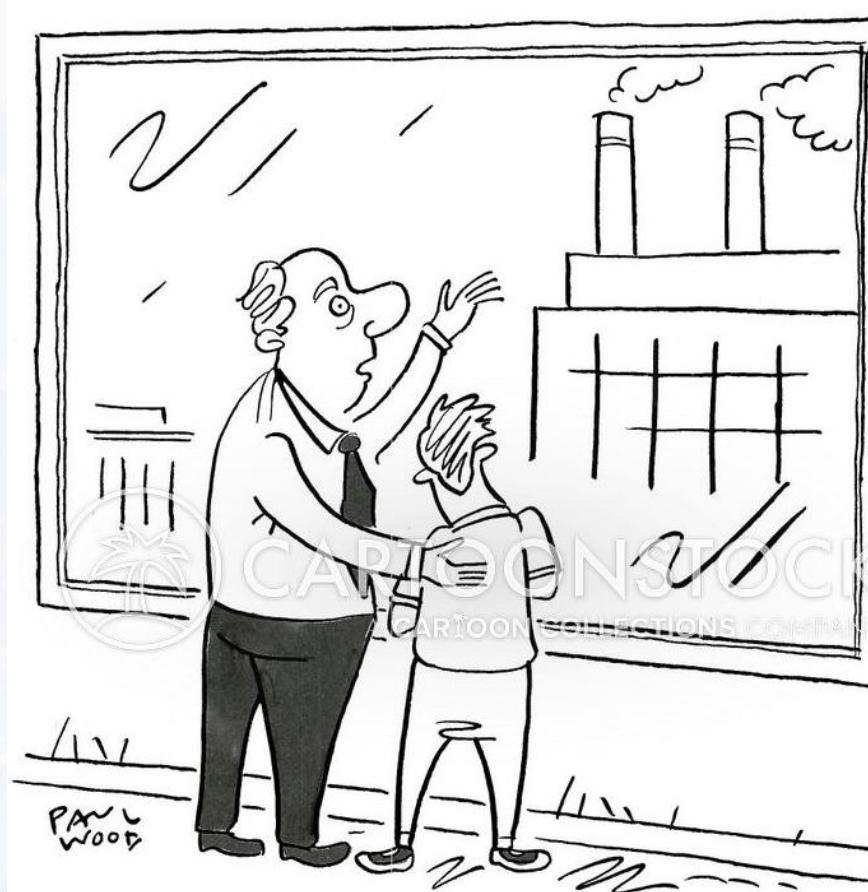
Lecture 11: Recurrent Neural Networks (RNNs) and Long Short-Term Memory (LSTMs)

Lecture 12: Combining LSTMs and CNNs

Lecture 15: Transformer

Lecture 14: Project Presentations

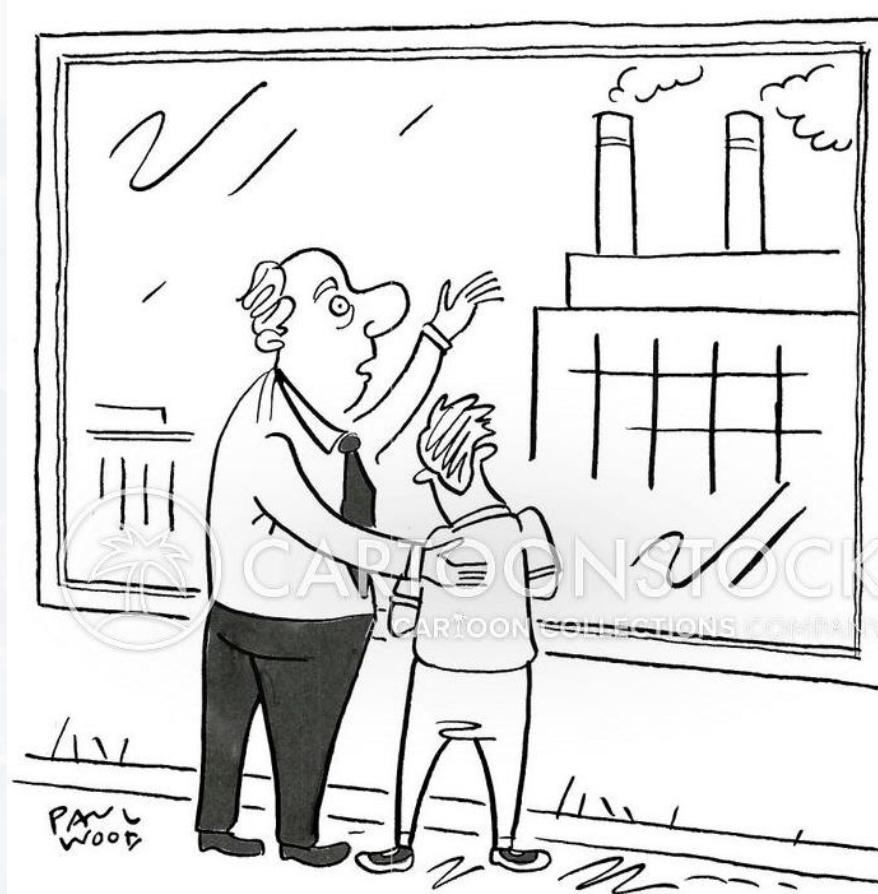
**Lecture 15: GNN**



ONE DAY SON, ALL THIS  
WILL BE RUN BY ROBOTS

## Outline

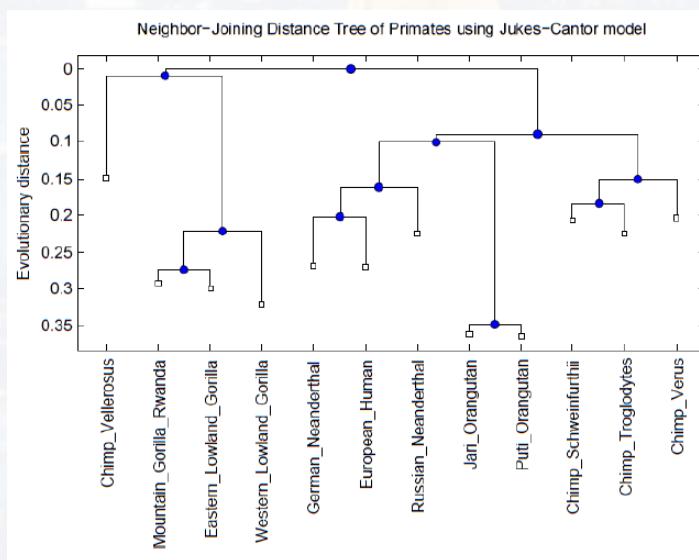
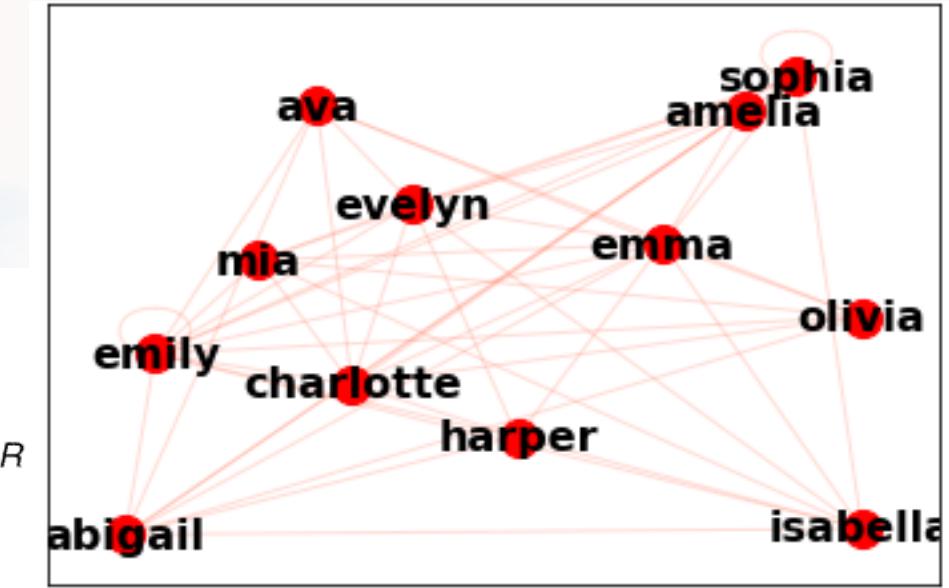
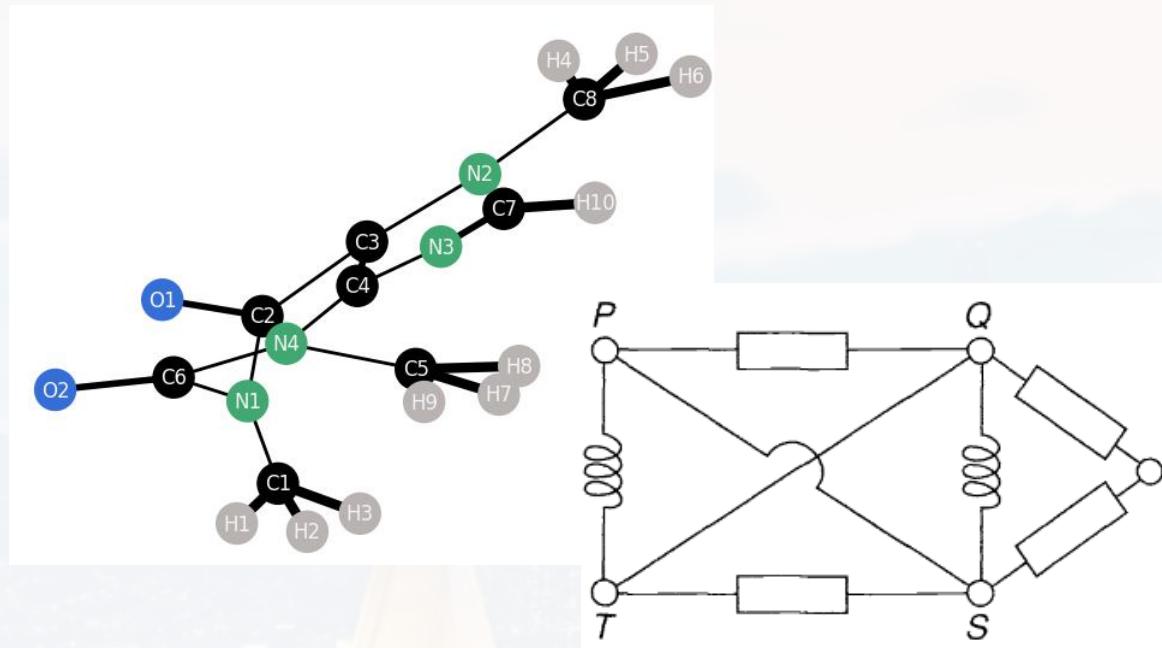
- What is a Graph
- The ANN Part
- PyTorch Example



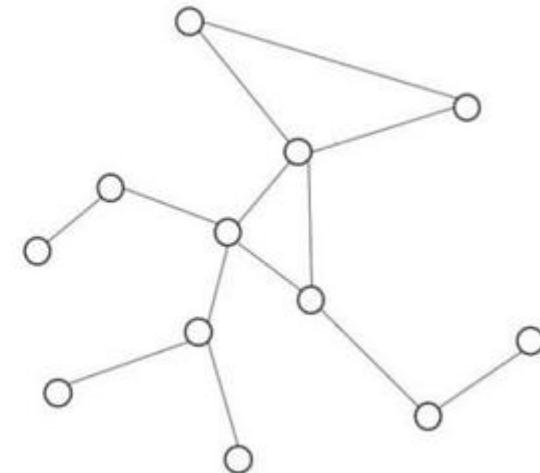
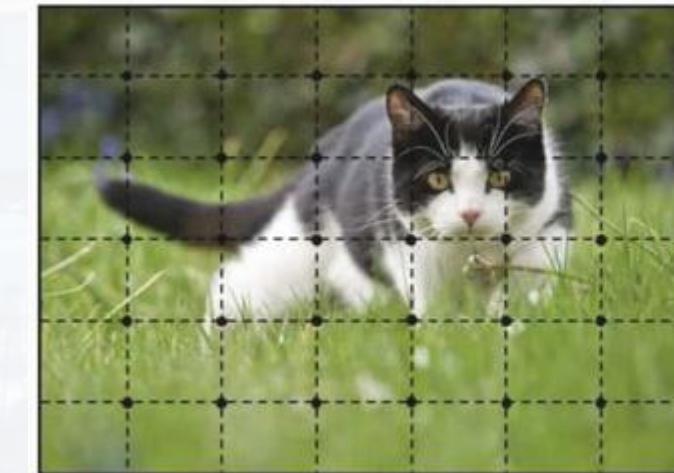
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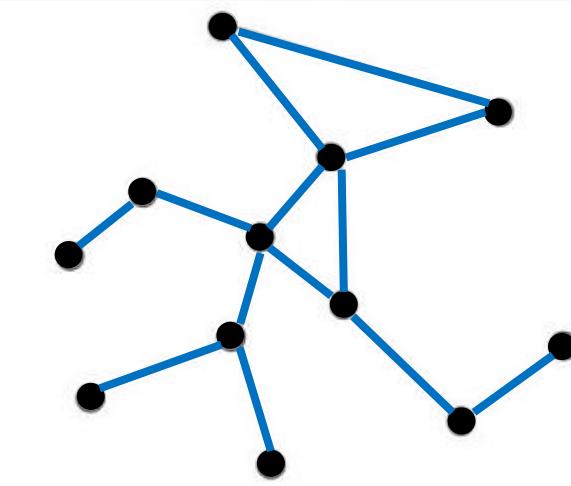
## Outline

- What is a Graph
- The ANN Part
- PyTorch Example



<https://doi.org/10.1016/j.aiopen.2021.01.001>





Graph  $G$

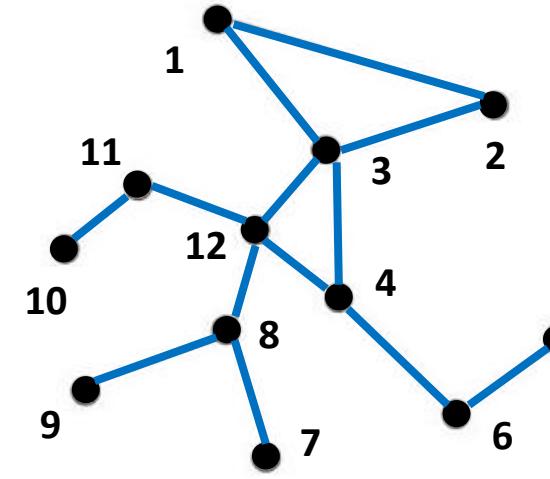
nodes  $N$  (vertices  $V$ )

edges  $E$

$G = G(N, E)$

- social networks
- street maps
- workflows/planning
- biological signal pathways
- image processing

- nodes can have **features**  
molecules: mass/ electronegativity  
people: age, income, sex, ...
- edges can have **attributes**  
molecules: bond length/strength  
people: relations (work, friend, family)



structural information: **adjacency matrix  $A$**

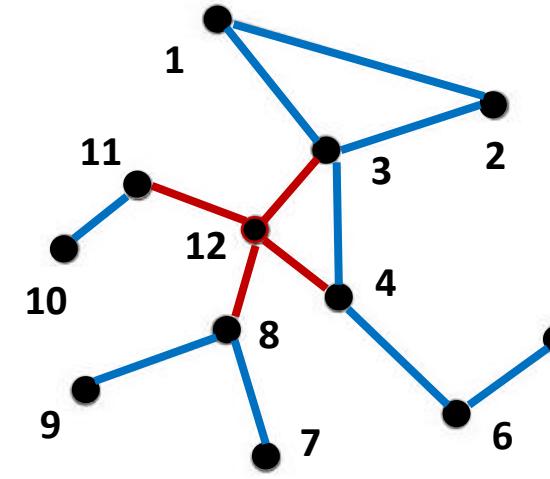
$A_{ij} = 1$  if  $(n_i, n_j) \in E$   
(nodes  $n_i$  and  $n_j$  have a common edge)

$A_{ij} = 0$  else

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Graph  $G = G(N, E)$

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node 12 has four first degree neighbors

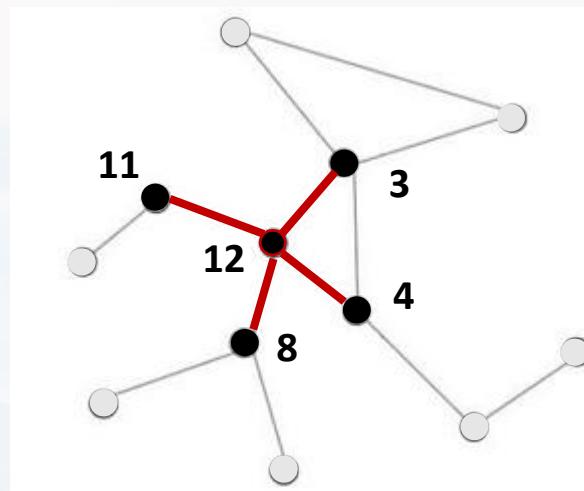
**degree  $d$**  of a node

$$d(n_i) = \sum_j A_{ij}$$

$$A = \begin{pmatrix} 0 & 1 & 1 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 1 & 0 & 1 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 1 & 1 & 0 & 1 & 0 & 0 & 0 & 0 & 0 & 0 & 1 \\ 0 & 0 & 1 & 0 & 0 & 1 & 0 & 0 & 0 & 0 & 1 \\ 0 & 0 & 0 & 0 & 0 & 1 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 1 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 1 & 1 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 1 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 1 & 0 & 1 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 1 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 1 \end{pmatrix}$$
  
 $\boxed{\begin{matrix} 0 & 0 & \textcolor{red}{1} & \textcolor{red}{1} & 0 & 0 & 0 & \textcolor{red}{1} & 0 & 0 & \textcolor{red}{1} & 0 \end{matrix}}$

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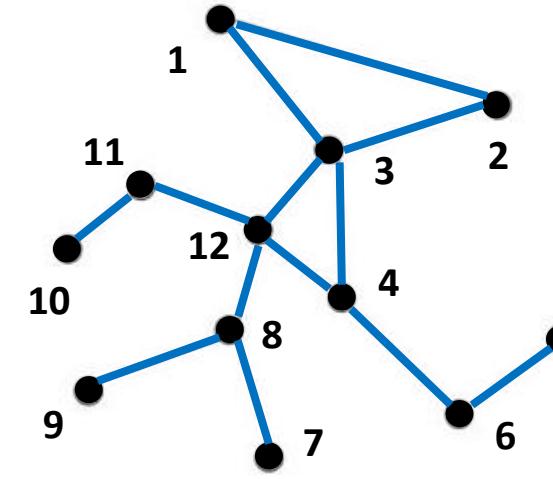
**first degree neighborhood  $\mathcal{N}$**

$$\mathcal{N}(n_i) = \{n_j \in N: (n_i, n_j) \in E\}$$

$$A = \begin{pmatrix} 0 & 1 & 1 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 1 & 0 & 1 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 1 & 1 & 0 & 1 & 0 & 0 & 0 & 0 & 0 & 0 & 1 \\ 0 & 0 & 1 & 0 & 0 & 1 & 0 & 0 & 0 & 0 & 1 \\ 0 & 0 & 0 & 0 & 0 & 1 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 1 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 1 & 1 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 1 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 1 & 0 & 1 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 1 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 1 & 0 & 1 \end{pmatrix}$$
  
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Graph  $G = G(N, E)$

nodes  $N$  (vertices  $V$ )  
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A graph can have **loops**

structural information: **adjacency matrix  $A$**

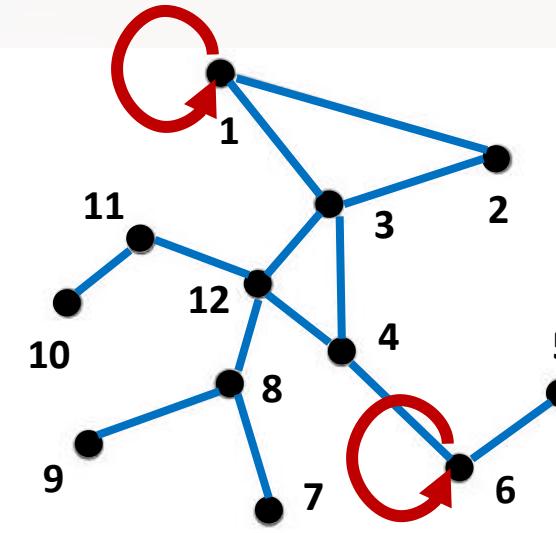
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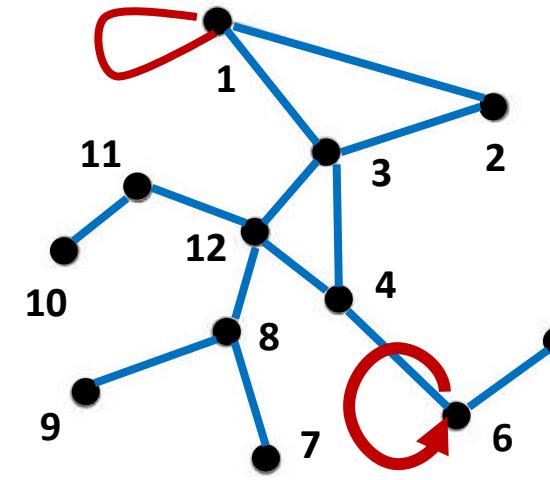
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A graph can have **loops**

### **note:**

$d(n_1) = 4$ , since loop is **undirected** and hits the node twice!

structural information: **adjacency matrix  $A$**

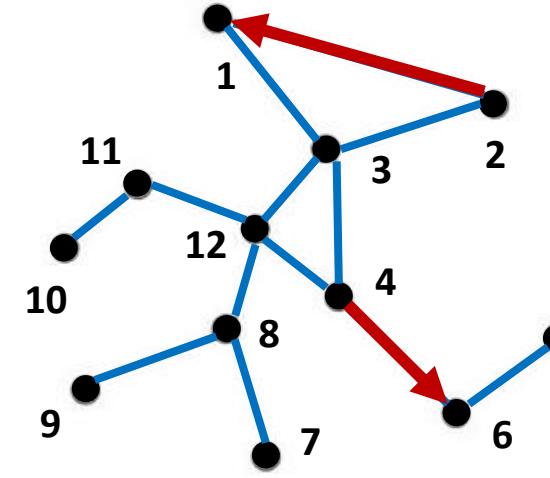
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Graph  $G = G(N, E)$

nodes  $N$  (vertices  $V$ )  
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$$A = \begin{pmatrix} 2 & 1 & 1 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 1 & 0 & 1 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 1 & 1 & 0 & 1 & 0 & 0 & 0 & 0 & 0 & 0 & 1 \\ 0 & 0 & 1 & 0 & 0 & 1 & 0 & 0 & 0 & 0 & 1 \\ 0 & 0 & 0 & 0 & 0 & 1 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 1 & 1 & 1 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 1 & 0 & 1 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 1 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 1 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 1 & 0 & 0 \\ 0 & 0 & 1 & 1 & 0 & 0 & 0 & 1 & 0 & 1 & 0 \end{pmatrix}$$



A graph can be **directed**

structural information: **adjacency matrix  $A$**

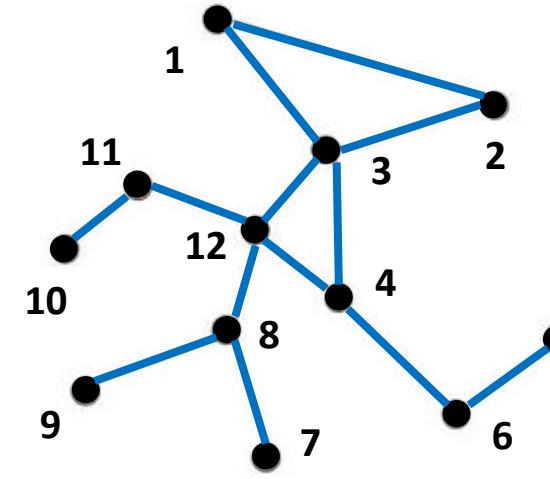
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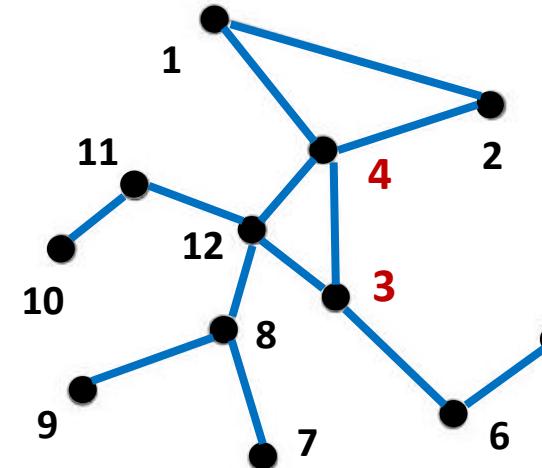
$A_{ij} = 0$  else

The order of counting the nodes is not relevant!  
**(permutation invariance)**

$$A = \begin{pmatrix} 0 & 1 & 1 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 1 & 0 & 1 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 1 & 1 & 0 & 1 & 0 & 0 & 0 & 0 & 0 & 0 & 1 \\ 0 & 0 & 1 & 0 & 0 & 1 & 0 & 0 & 0 & 0 & 1 \\ 0 & 0 & 0 & 0 & 0 & 1 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 1 & 1 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 1 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 1 & 0 & 1 \\ 0 & 0 & 0 & 0 & 0 & 0 & 1 & 0 & 1 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 1 \end{pmatrix}$$

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The order of counting the nodes is not relevant!  
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Each graph can be represented by  **$N!$**  adjacency matrices!

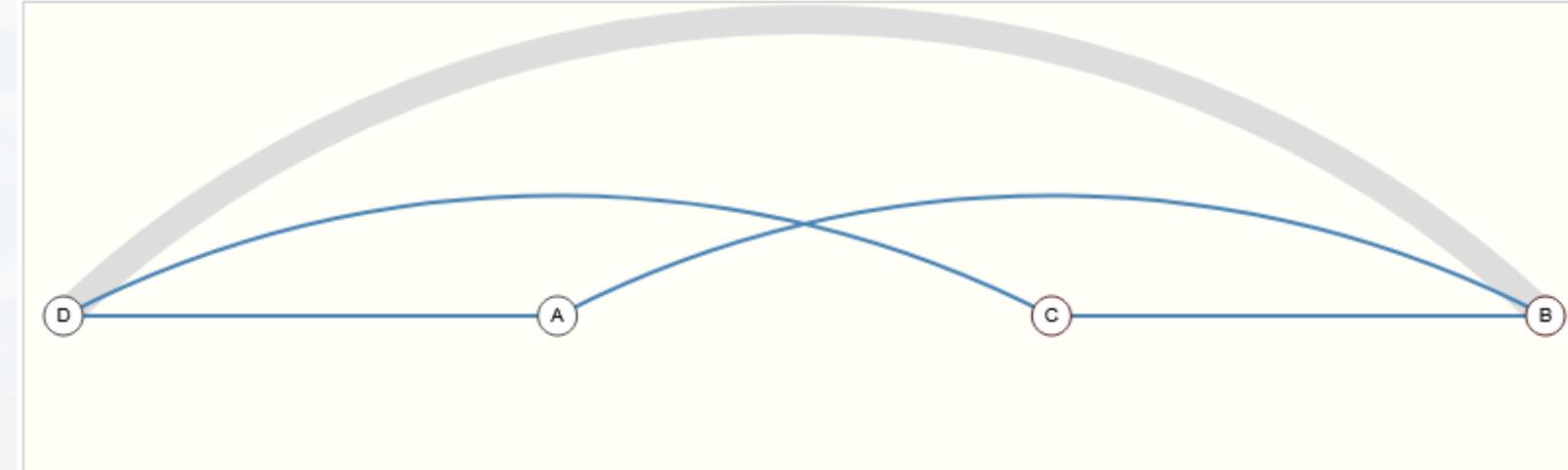
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Graph  $G = G(N, E)$

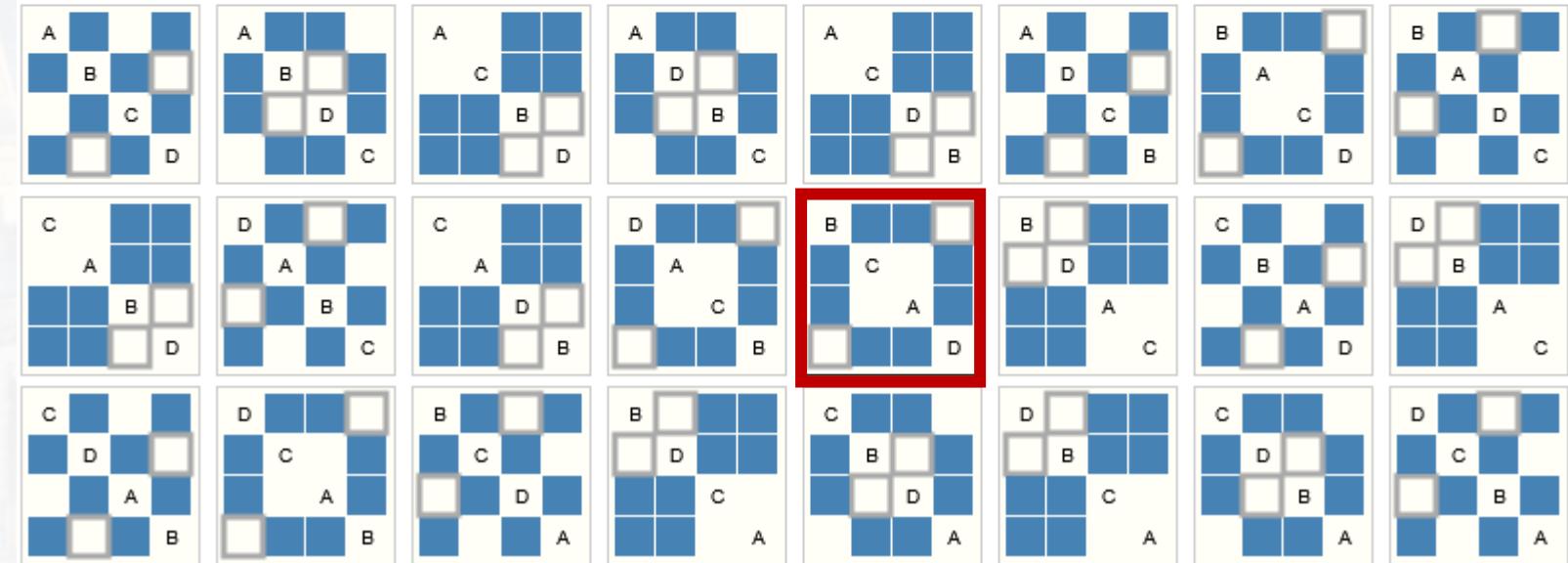
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[animation here](#)

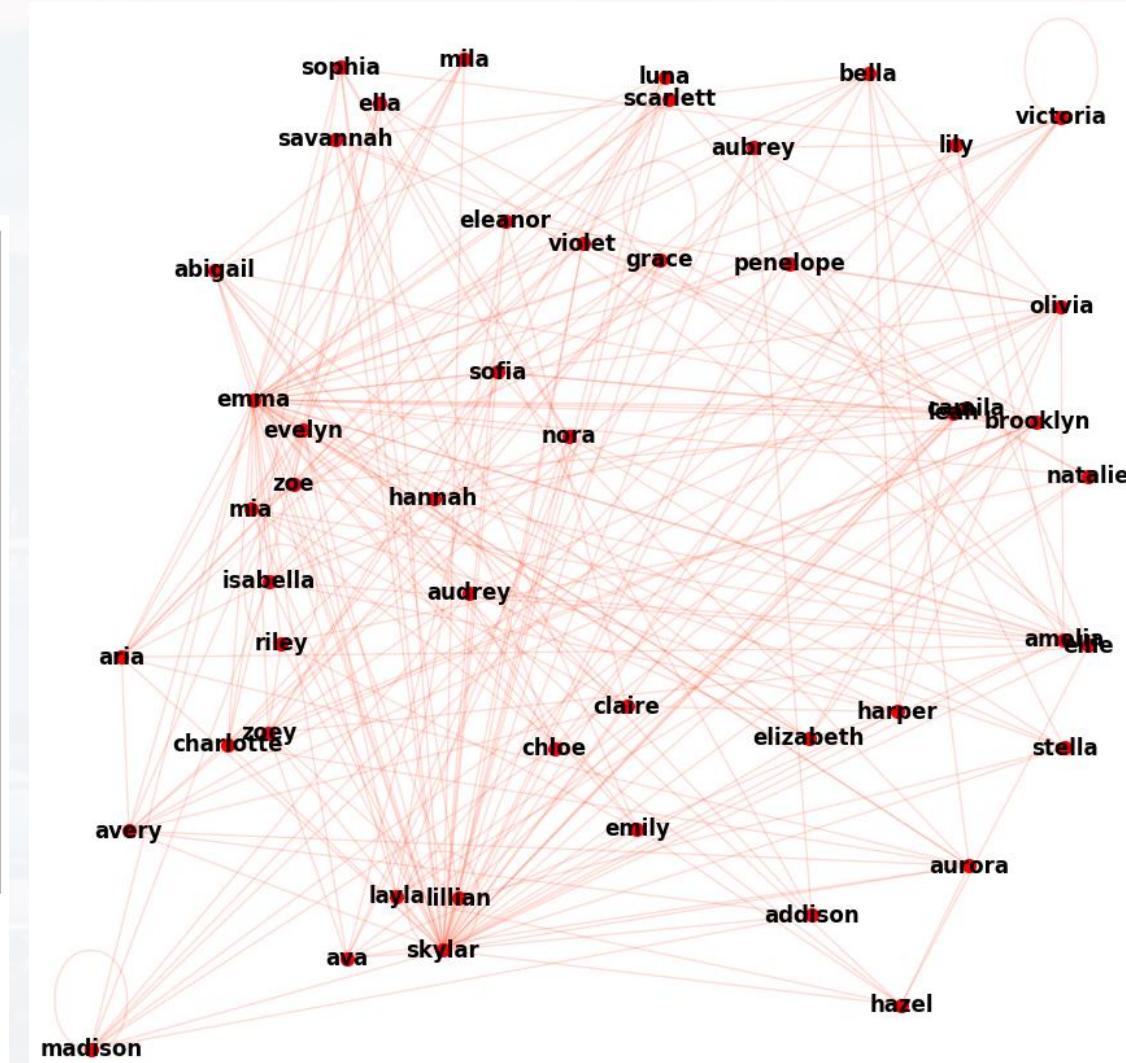
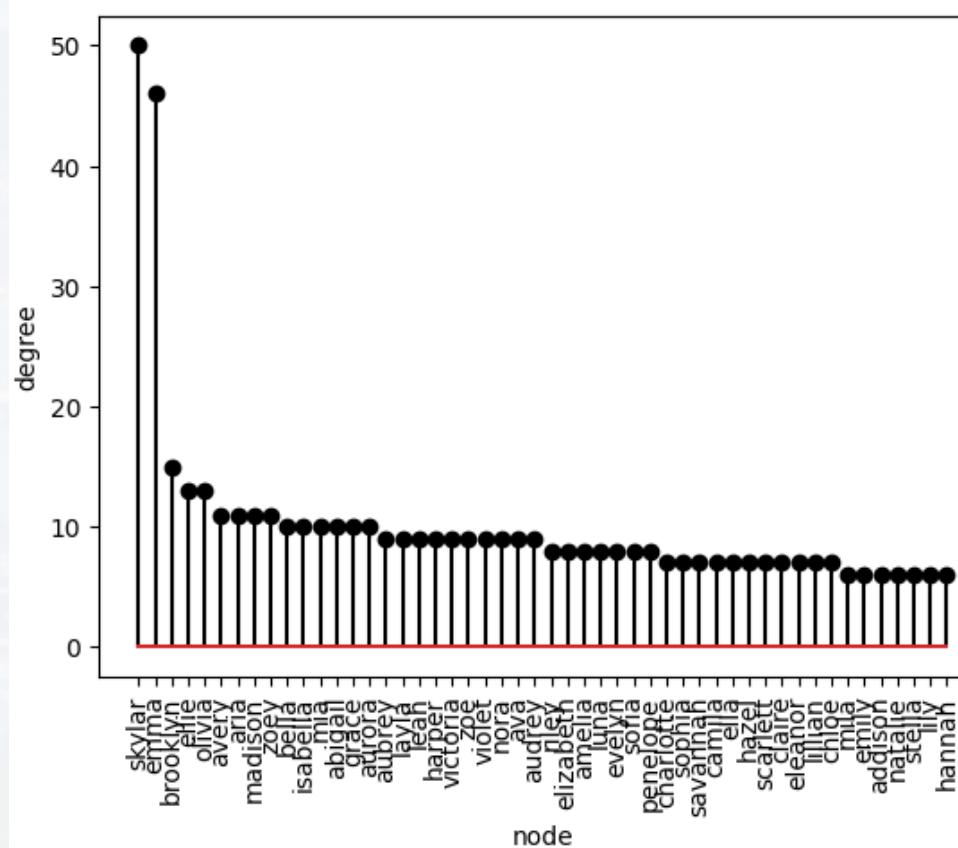




## visualizing a graph:

```
import networkx as nx #pip install networkx
```

see: Graph\_I.ipynb

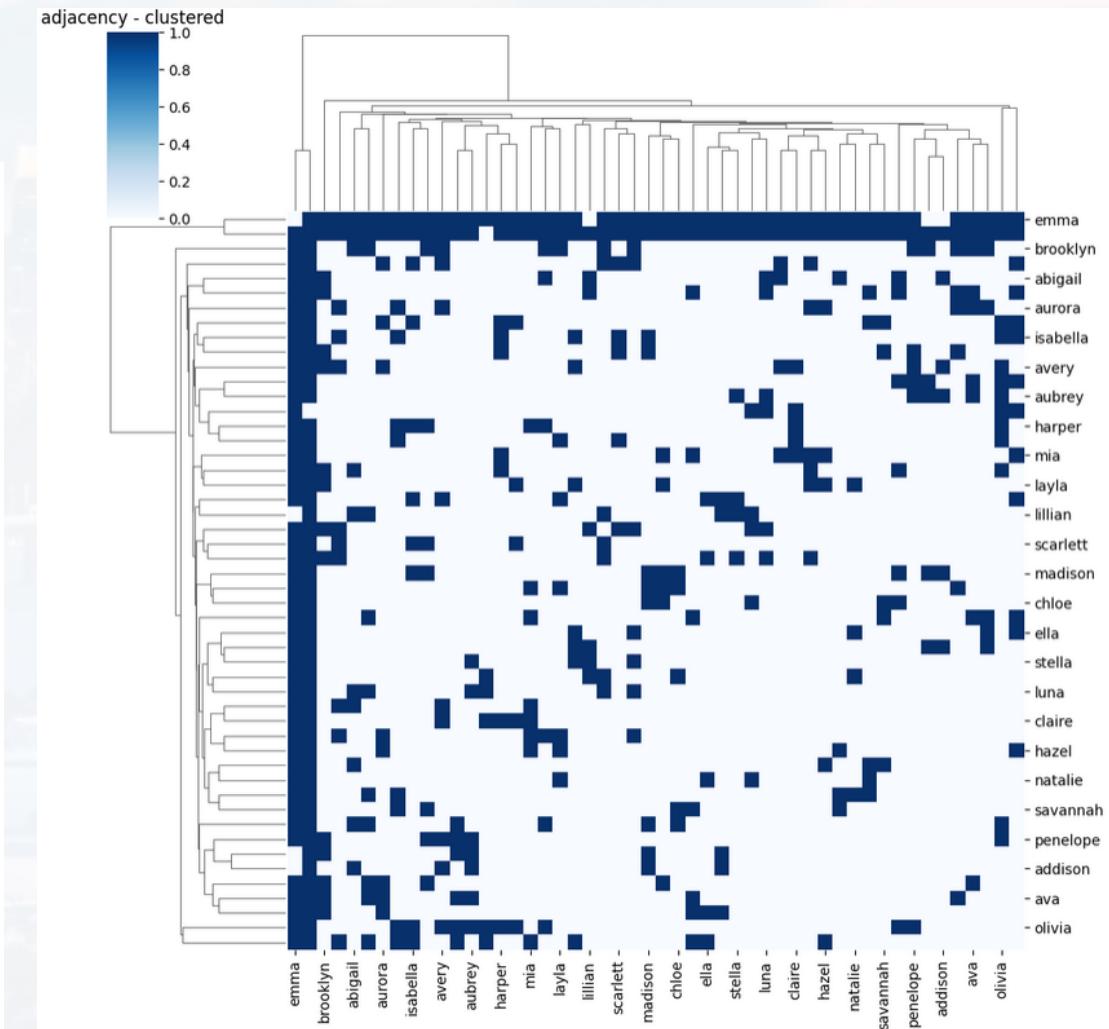
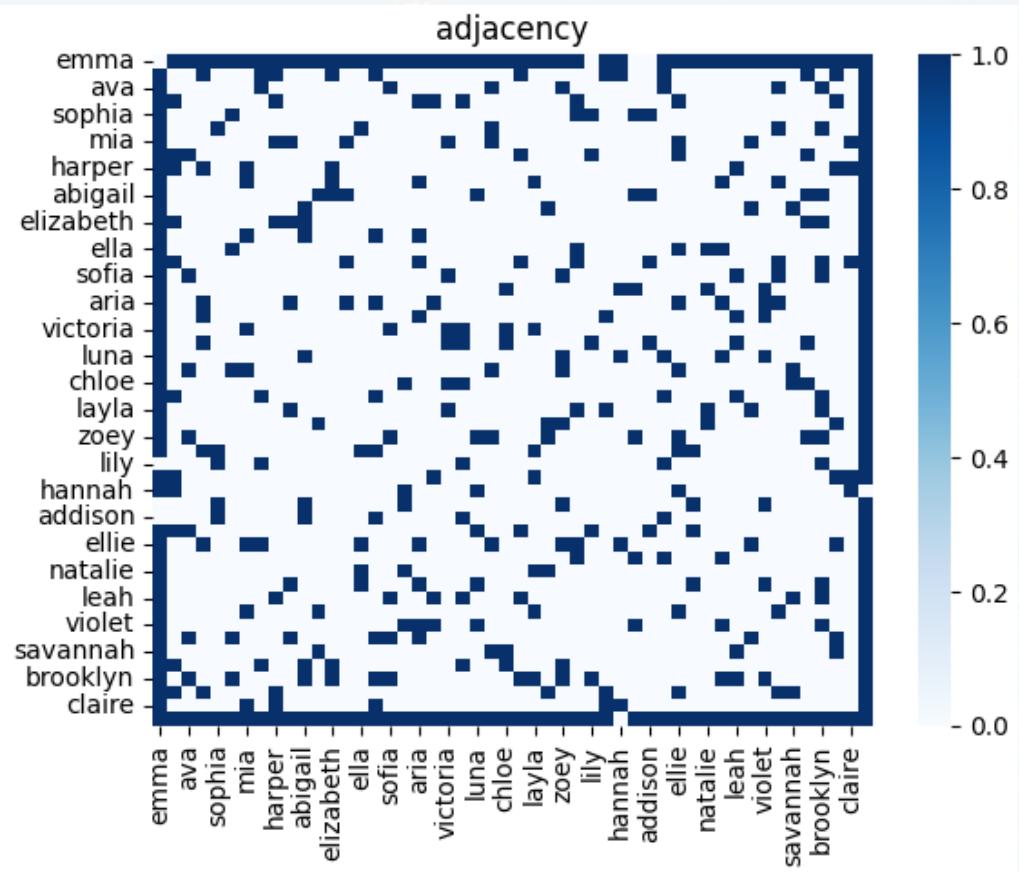




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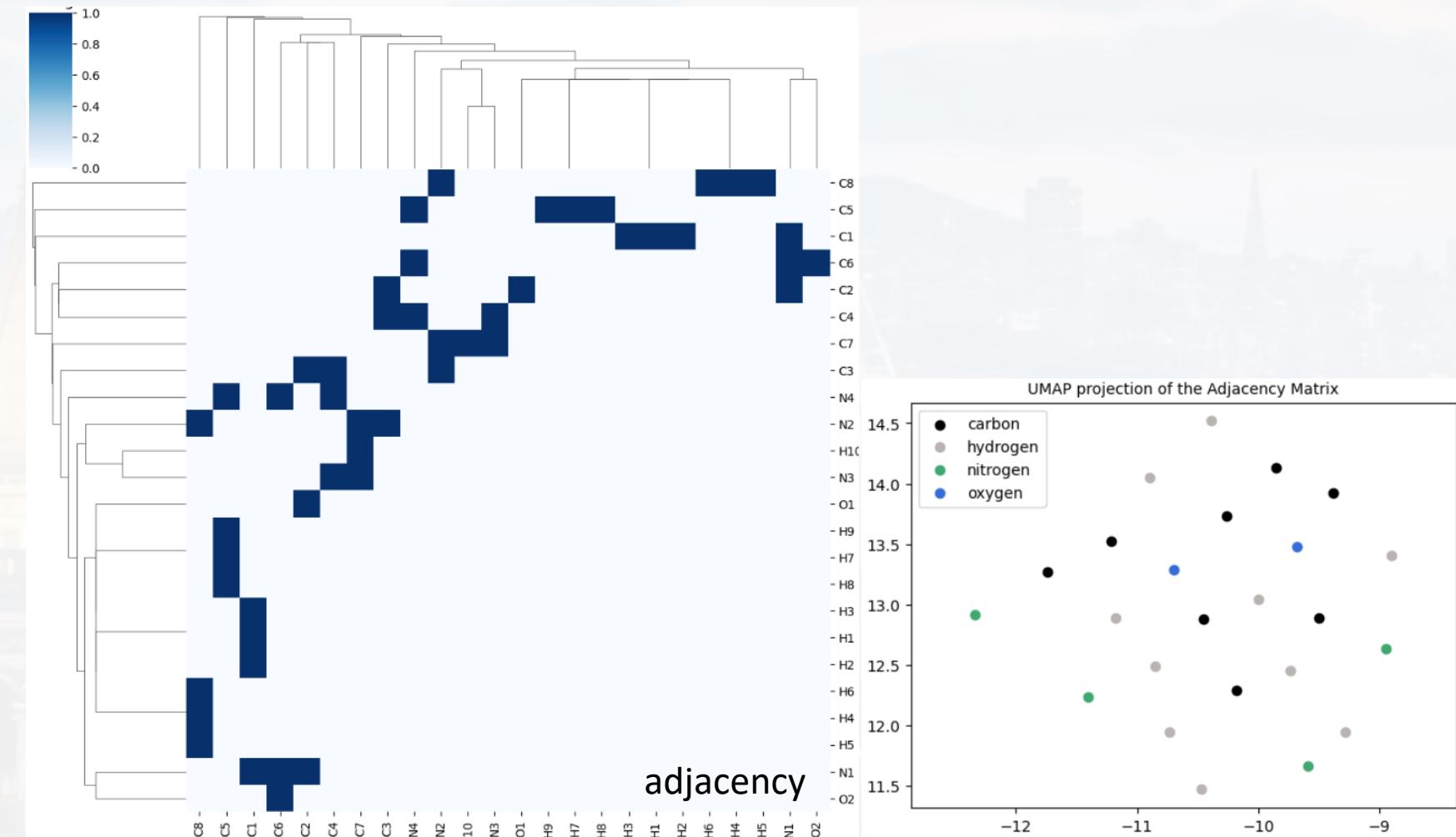


building and visualizing a **weighted** graph:

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import networkx as nx #pip install networkx
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see: [Graph\\_II.ipynb](#)

Caffein molecule





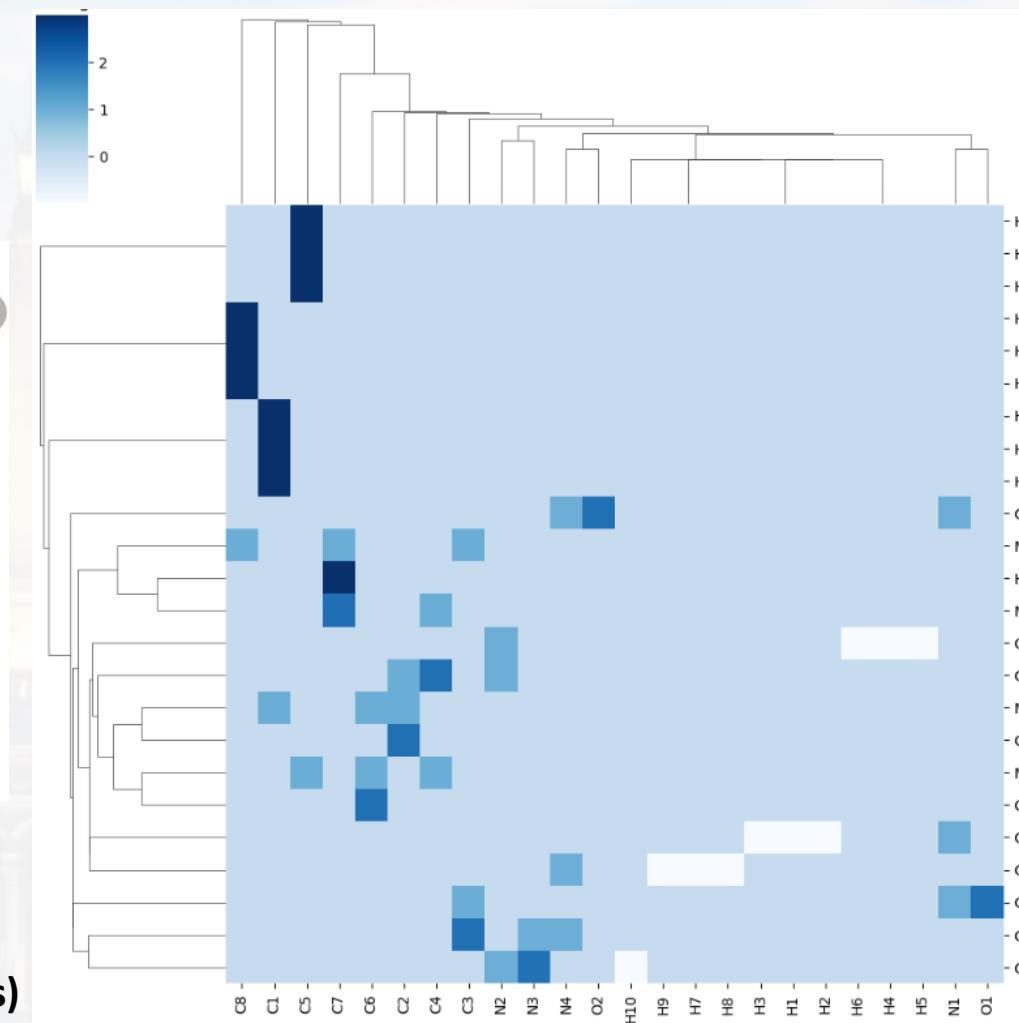
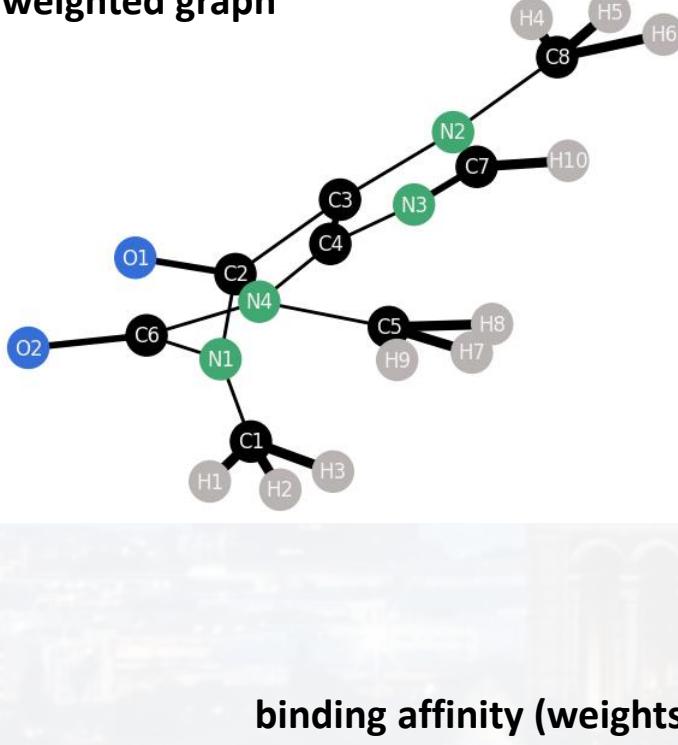
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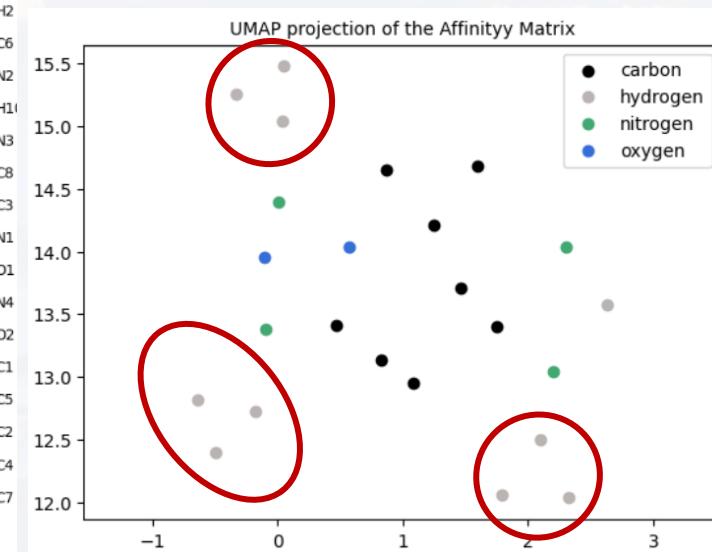
see: [Graph\\_II.ipynb](#)

Caffeine molecule

**weighted graph**



hydrogen atoms are at the edges of the molecule!





more about graphs:

$$d(n_i) = \sum_j A_{ij}$$

degree of node  $n_i$

$$\mathcal{N}(n_i) = \{n_j \in N : (n_i, n_j) \in E\}$$

neighborhood  $\mathcal{N}(n_i)$  of node  $n_i$   
for first degree neighborhood  $|\mathcal{N}(n_i)| = d(n_i)$

$$S_{com} = |\mathcal{N}(n_i) \cap \mathcal{N}(n_j)|$$

number for neighbors nodes  $n_i$  and  $n_j$  have in common.

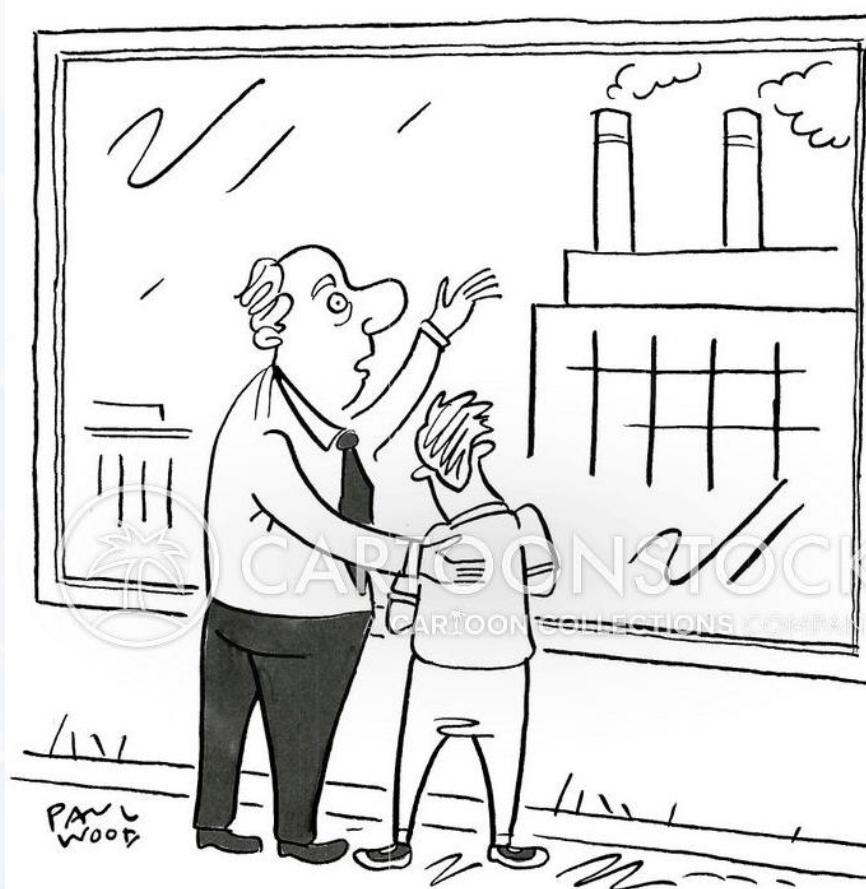
**idea:** nodes with many common neighbors are more likely to be similar or have a potential connection.

$$S_{rat} = \frac{|\mathcal{N}(n_i) \cap \mathcal{N}(n_j)|}{|\mathcal{N}(n_i) \cup \mathcal{N}(n_j)|}$$

ratio for neighbors nodes  $n_i$  and  $n_j$  have in common.

**note:**

There are more quantities (“importance”, “centrality” etc.), but they are all a function of  $A_{ij}$ .



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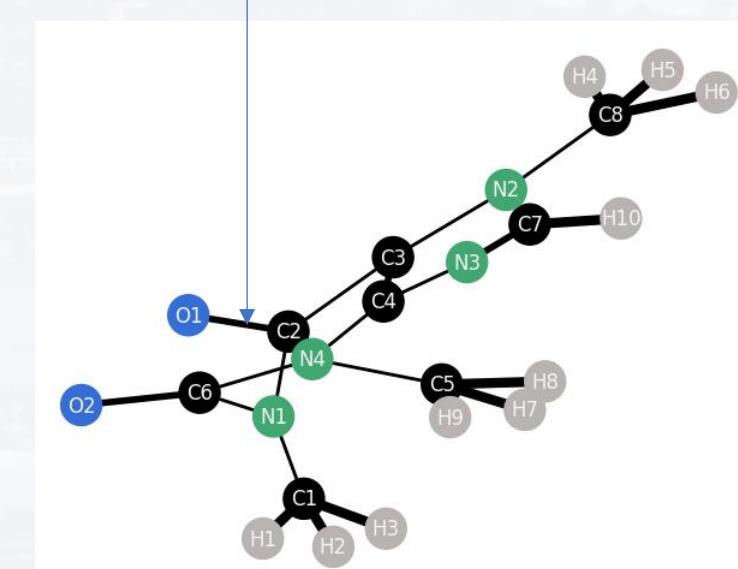
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What we can learn:

- node classification
- join nodes with similar properties to hyper nodes
- edge attributes, weights (weighted graph)
- edge prediction
- embedding (eg. 3D structure molecules)
- graph classification (is the molecule toxic y/n)
- graph regression (toxicity score)
- graph generation

weight: bond strength

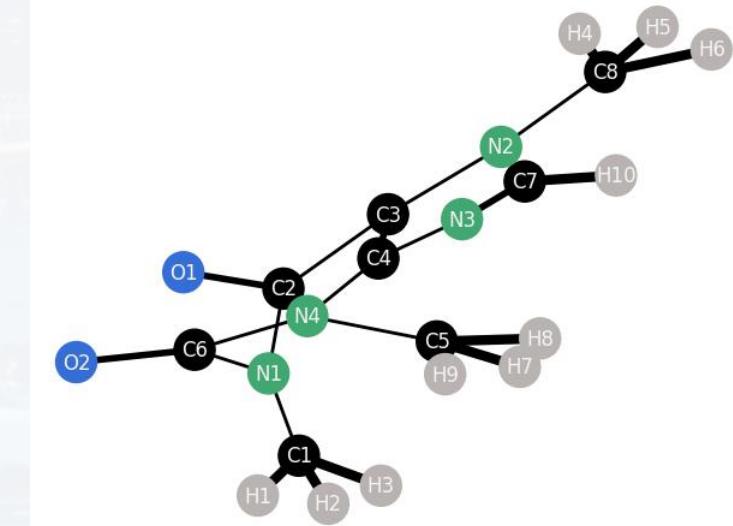




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**node (edge) level tasks**





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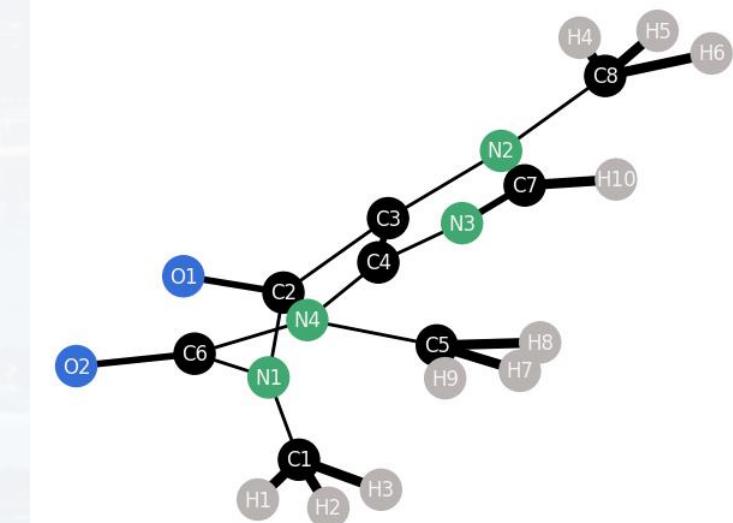
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information flow from one node to another:

**message passing**

different ways how:

- local averaging
- graph convolution (aka neighborhood aggregation)
- graph attention





What we can learn:

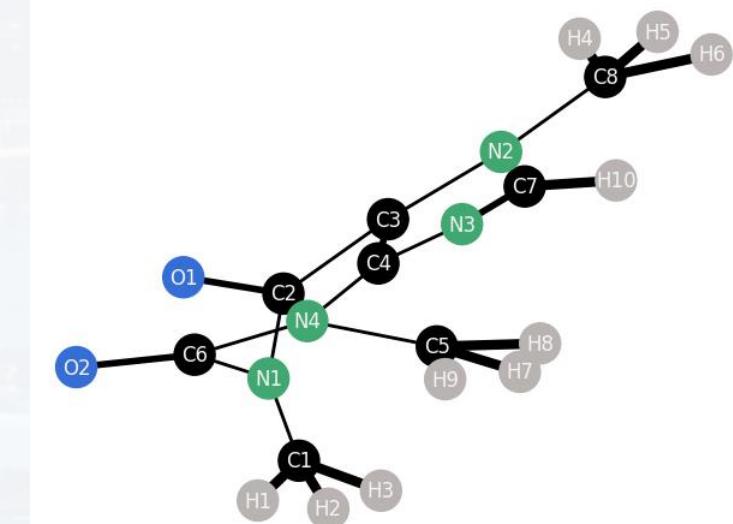
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- embedding (eg. 3D structure molecules)
- graph classification (is the molecule toxic y/n)
- graph regression (toxicity score)
- graph generation

information flow from one node to another:

**message passing**

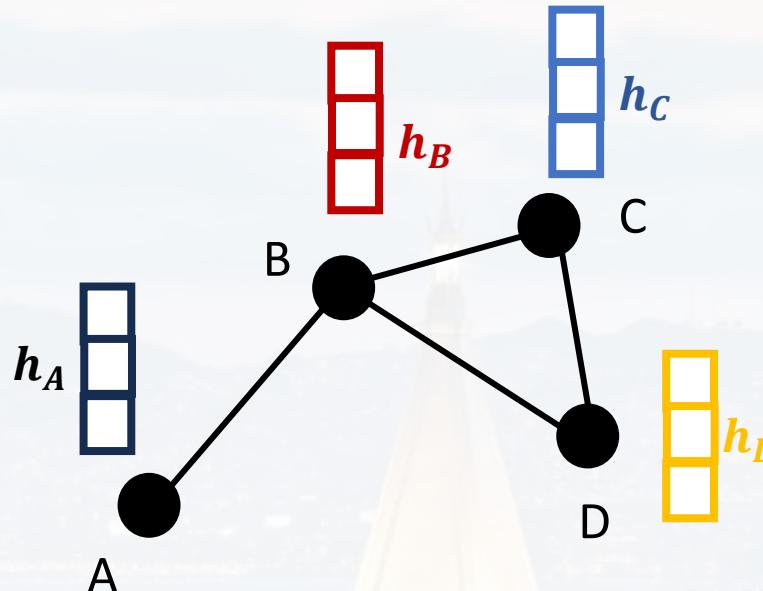
different ways how:

- local averaging
- **graph convolution** (aka neighborhood aggregation)
- **graph attention**



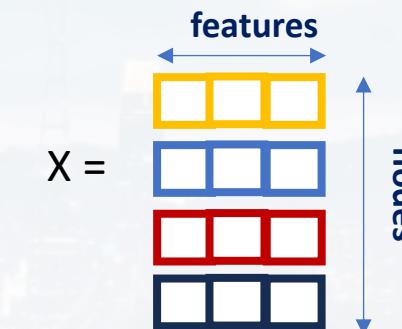


### Graph Convolution

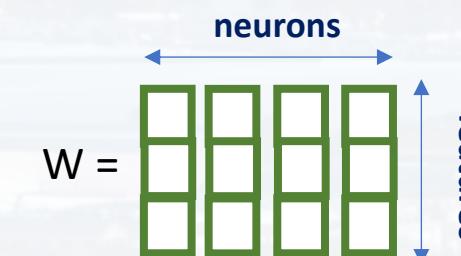


each node  $i$  has a **feature vector**  $h_i$

matrix X of shape (number of nodes, number of node features)



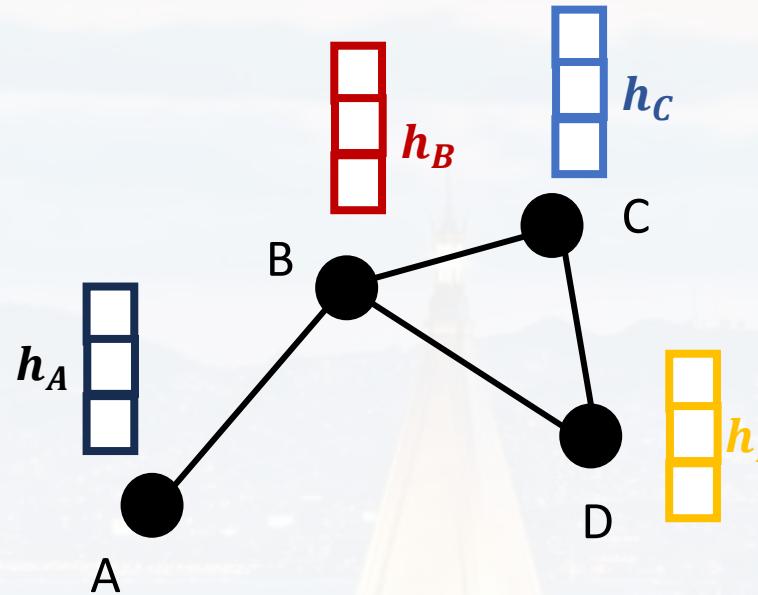
**weight matrix W** of shape  
(number of node features, number of neurons)



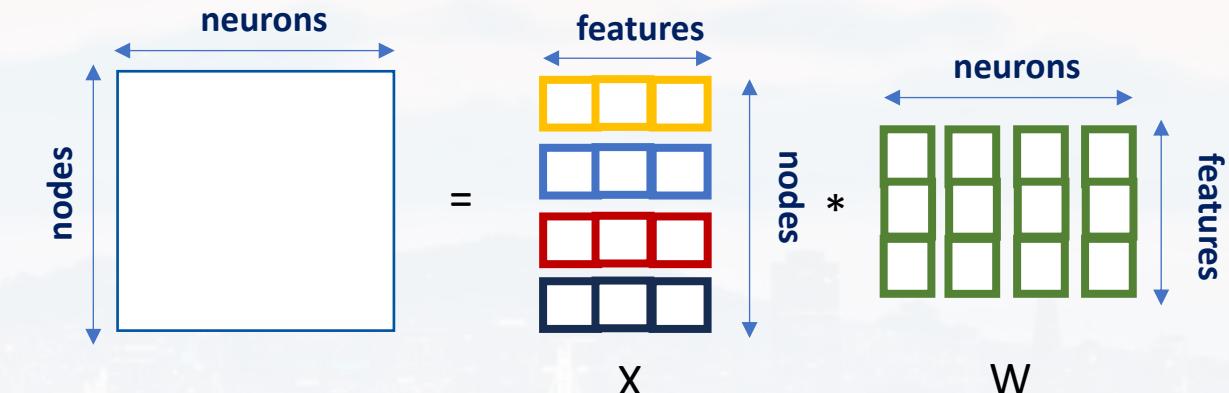
note: only **one** W for the entire graph  
W is a **learnable**



### Graph Convolution

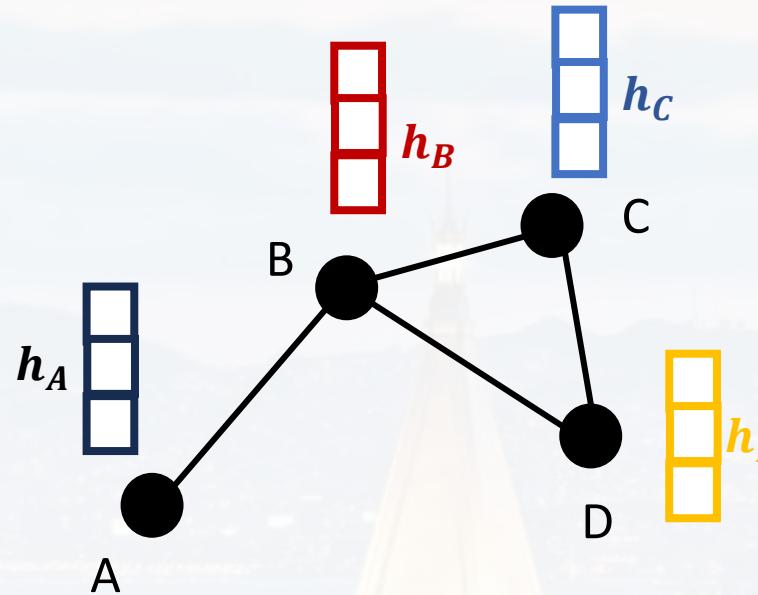


each node  $i$  has a **feature vector**  $h_i$

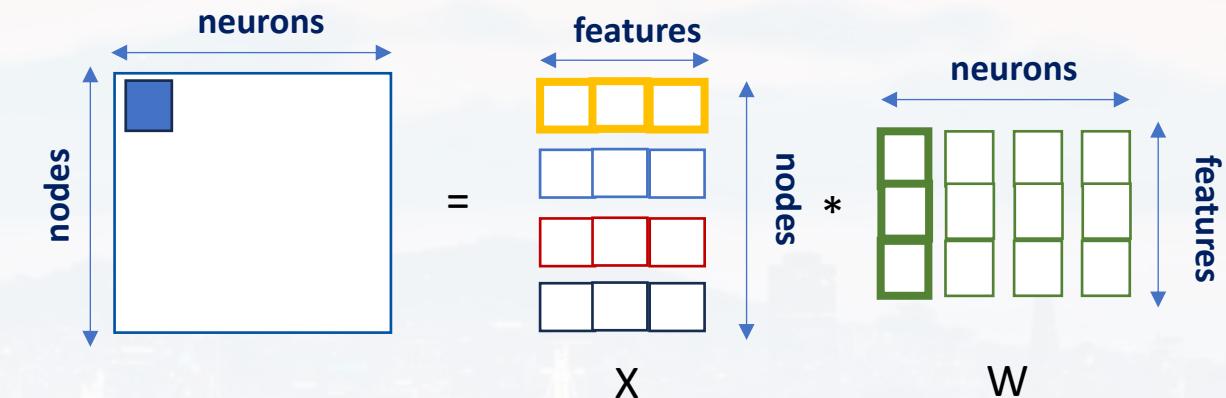




### Graph Convolution

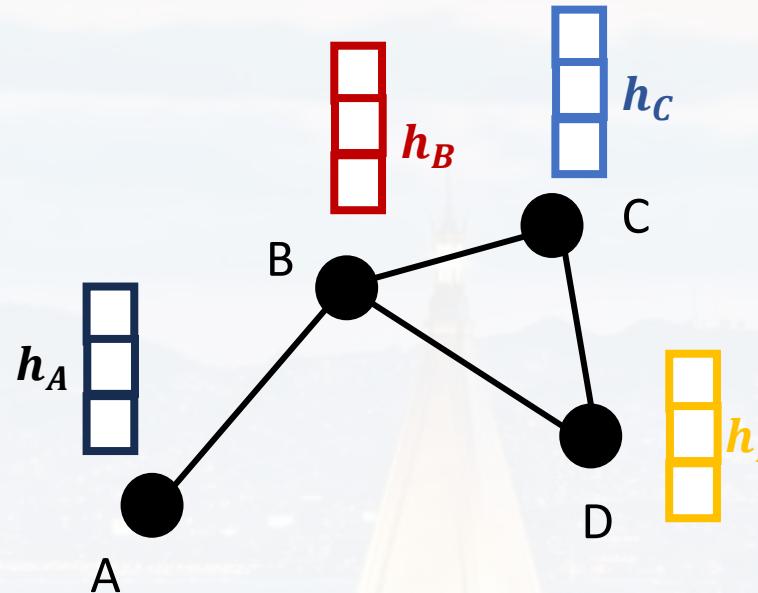


each node  $i$  has a **feature vector**  $h_i$

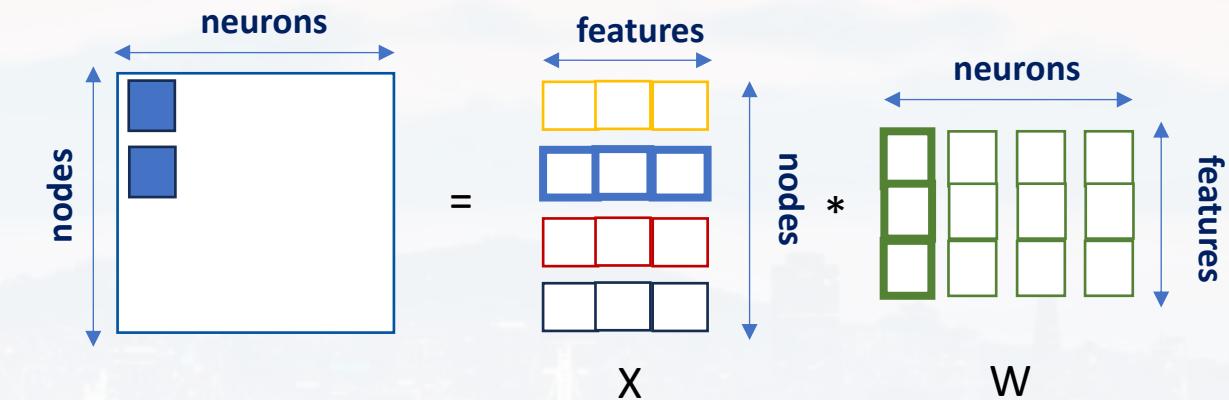




### Graph Convolution

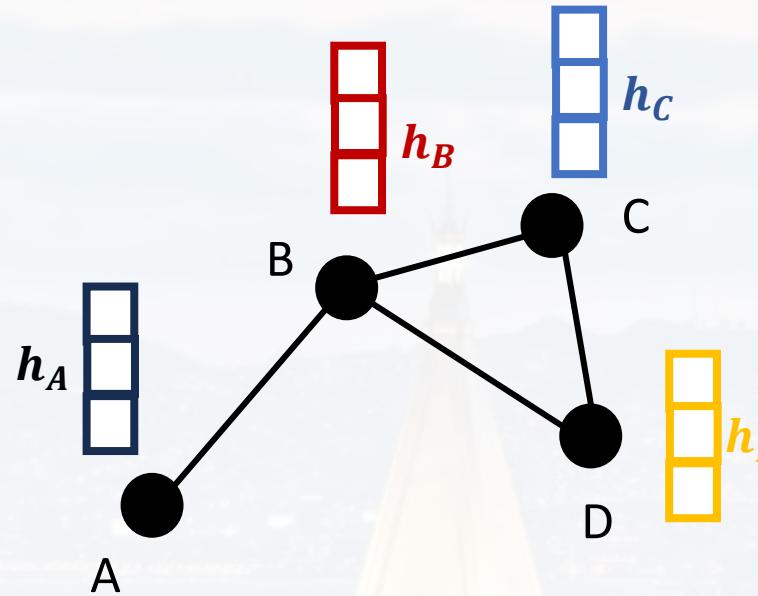


each node  $i$  has a **feature vector**  $h_i$

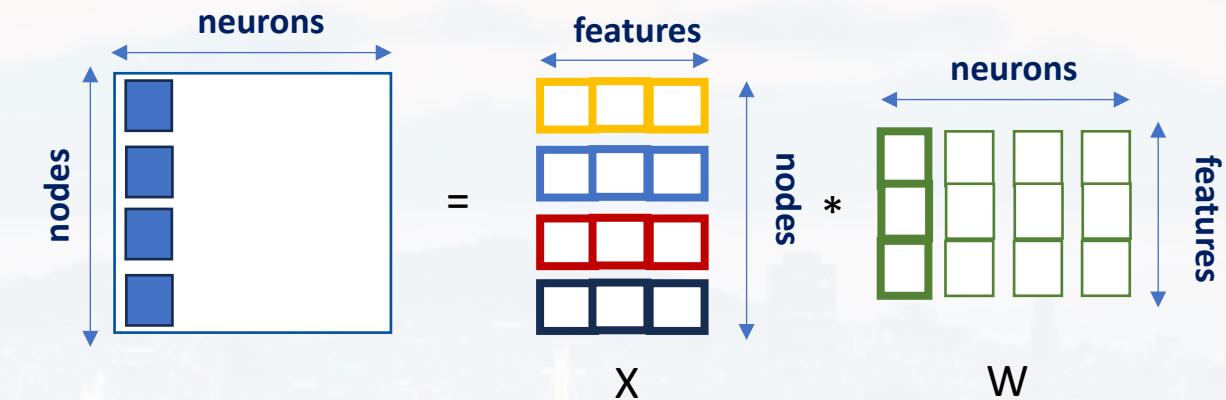




### Graph Convolution

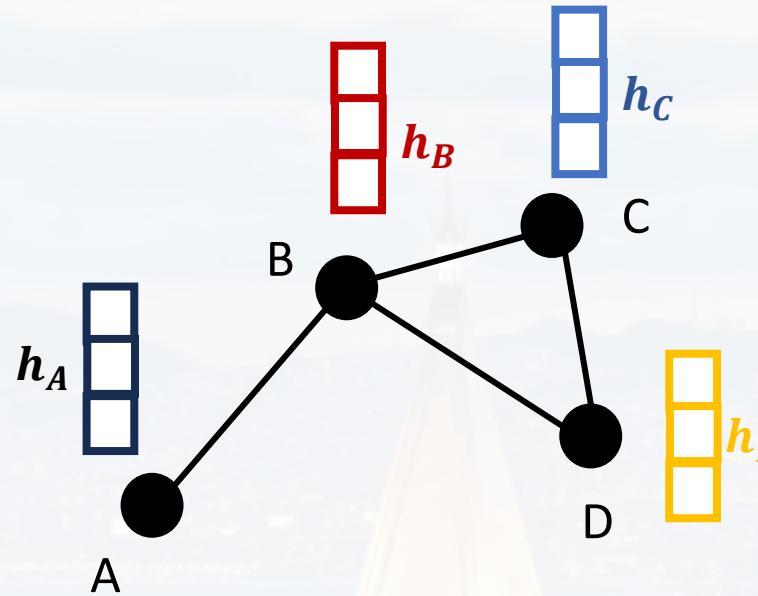


each node  $i$  has a **feature vector**  $h_i$

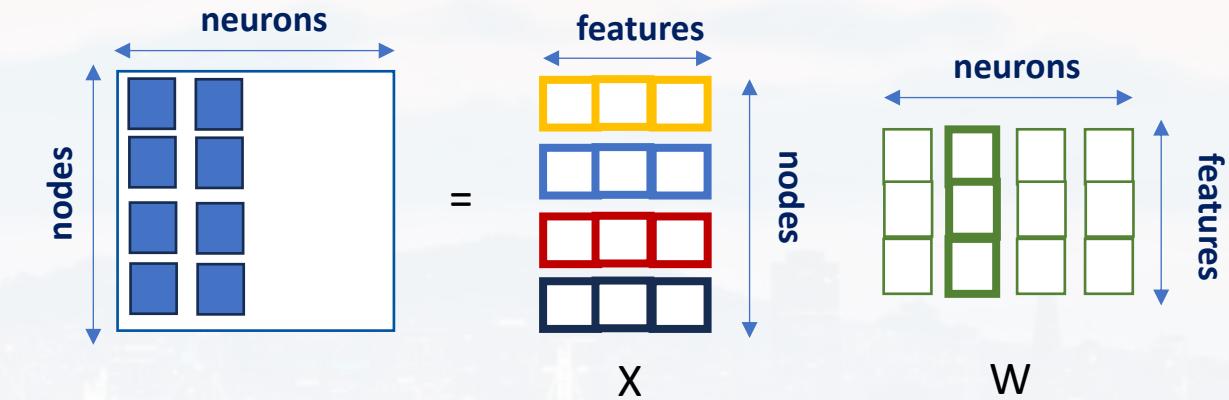




### Graph Convolution

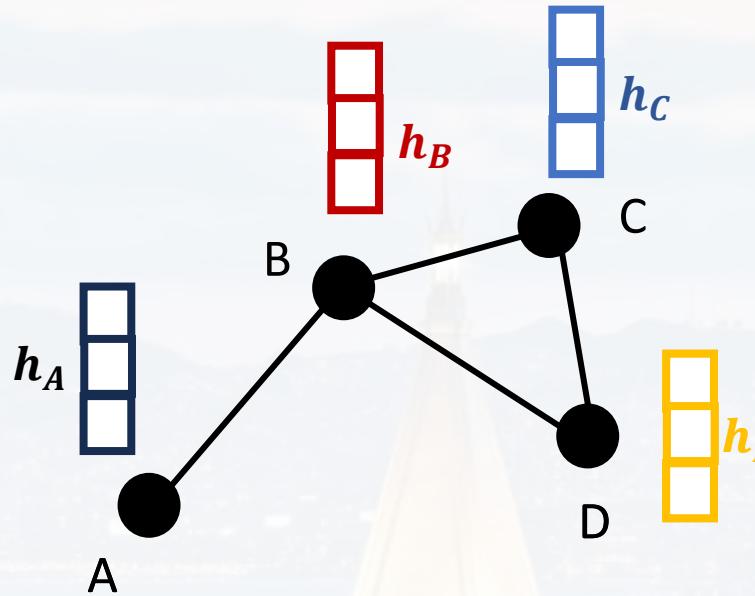


each node  $i$  has a **feature vector**  $h_i$

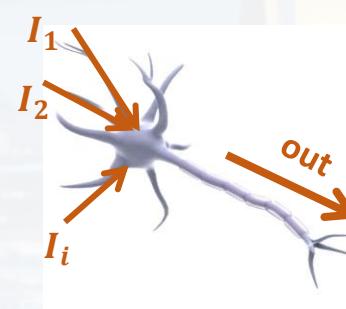
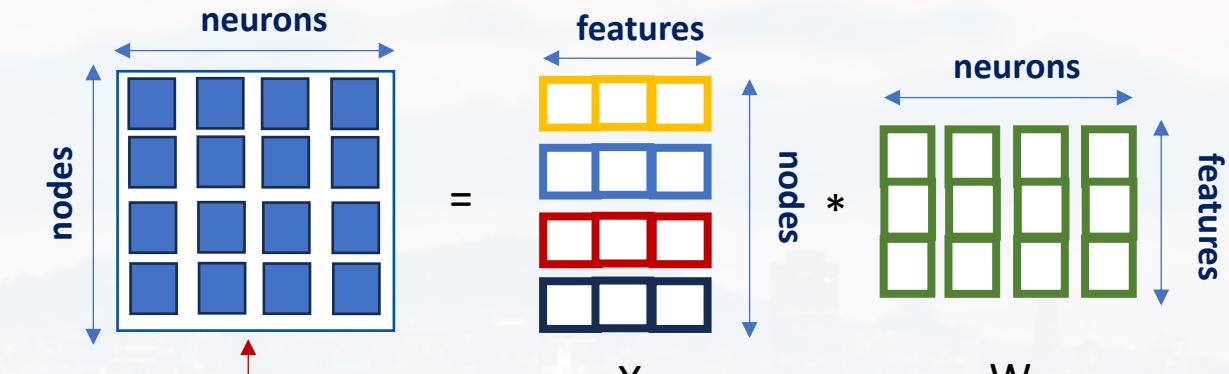




### Graph Convolution



each node  $i$  has a **feature vector**  $h_i$

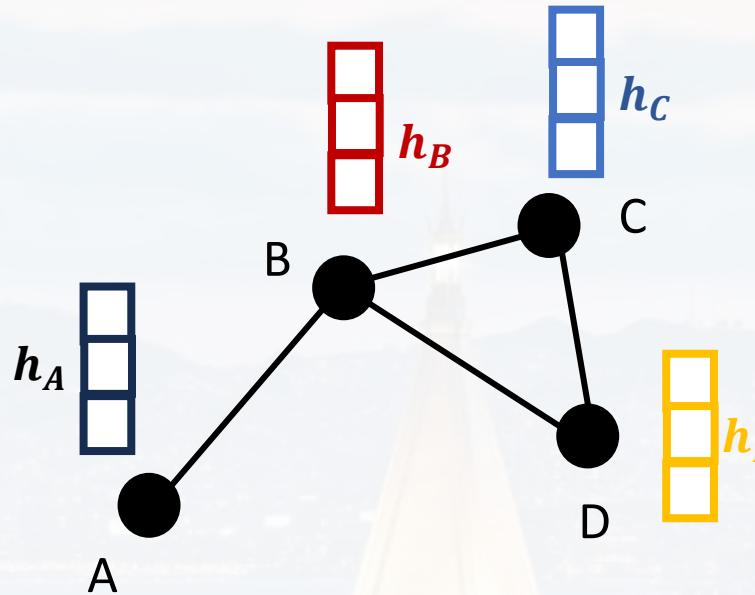


$$net = \sum_i I_i \cdot w_i + b$$

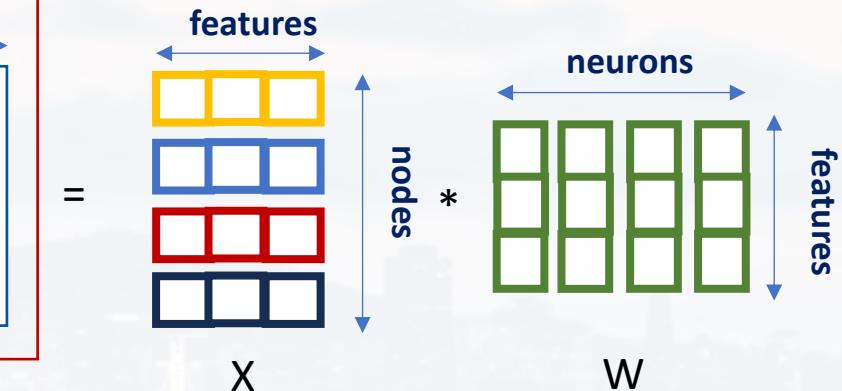
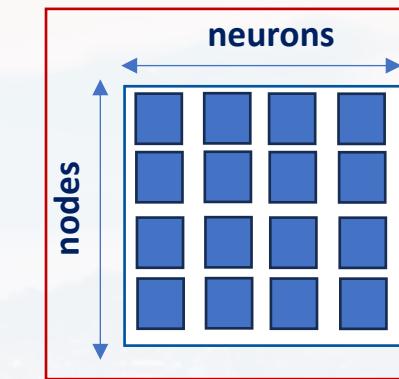
$$m_{jk} = \sum_i w_{ji} x_{ik}$$



### Graph Convolution

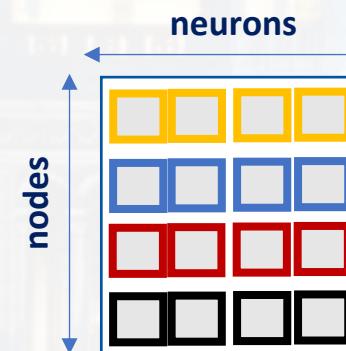


each node  $i$  has a **feature vector**  $h_i$



$$m_{jk} = \sum_i w_{ji} x_{ik}$$

depending on  $W$   
the output features  
may have different  
lengths than the  
input features

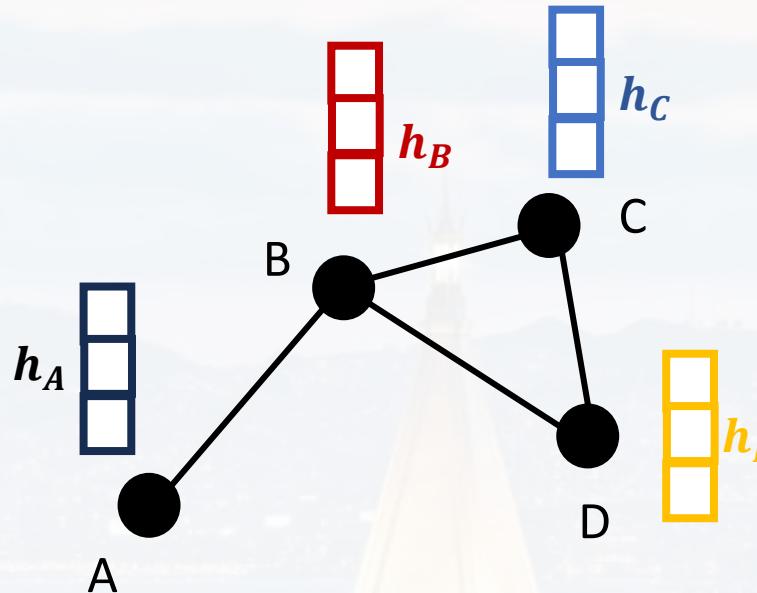


adjacency A

$$= \left[ \begin{pmatrix} 0 & 1 & 0 & 0 \\ 1 & 0 & 1 & 1 \\ 0 & 1 & 0 & 1 \\ 0 & 1 & 1 & 0 \end{pmatrix} + \begin{pmatrix} 1 & 0 & 0 & 0 \\ 0 & 1 & 0 & 0 \\ 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 1 \end{pmatrix} \right] * M$$

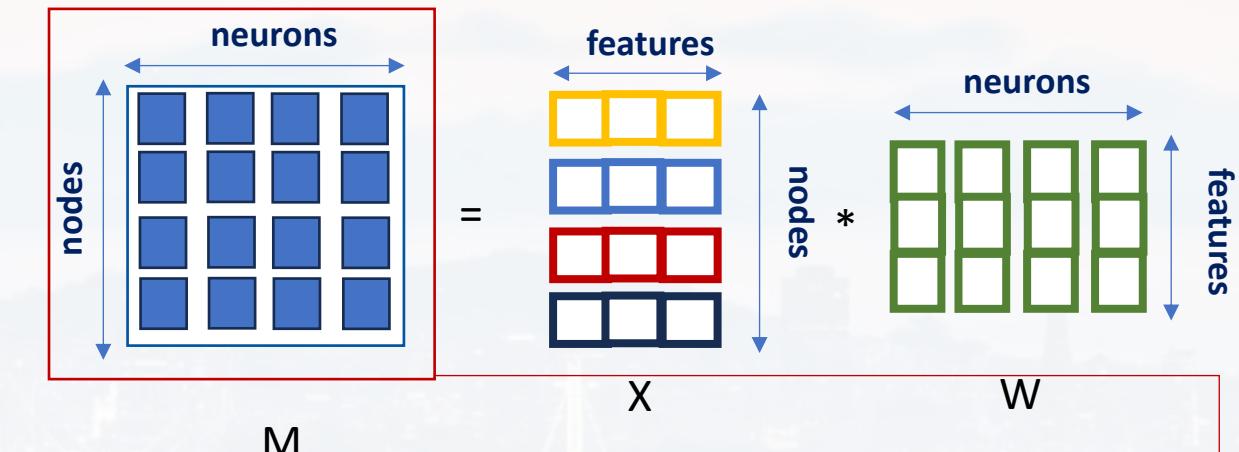


### Graph Convolution



pass through a ReLU and/or  
repeat the procedure with another  $W$   
(aka second convolution layer)

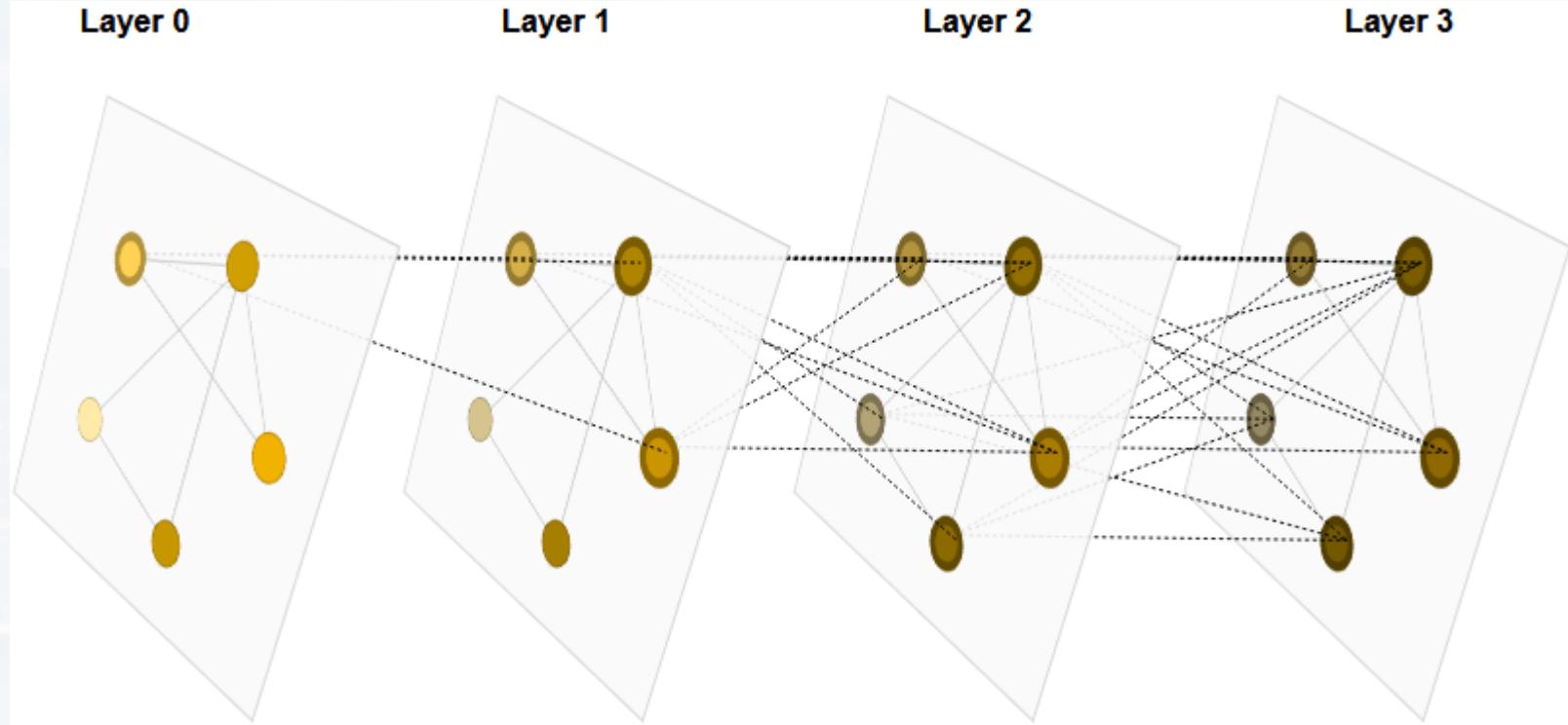
each node  $i$  has a **feature vector**  $h_i$



A diagram showing the computation of the adjacency matrix  $A$ . Below the matrix  $M$ , the equation  $A = \left[ \begin{pmatrix} 0 & 1 & 0 & 0 \\ 1 & 0 & 1 & 1 \\ 0 & 1 & 0 & 1 \\ 0 & 1 & 1 & 0 \end{pmatrix} + \begin{pmatrix} 1 & 0 & 0 & 0 \\ 0 & 1 & 0 & 0 \\ 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 1 \end{pmatrix} \right] * M$  is shown. The matrix  $M$  is highlighted with a red border. The resulting matrix  $A$  is shown below, with "nodes" on the vertical axis and "neurons" on the horizontal axis, containing 16 squares (4x4) with values 0, 1, or 2.



### Graph Convolution

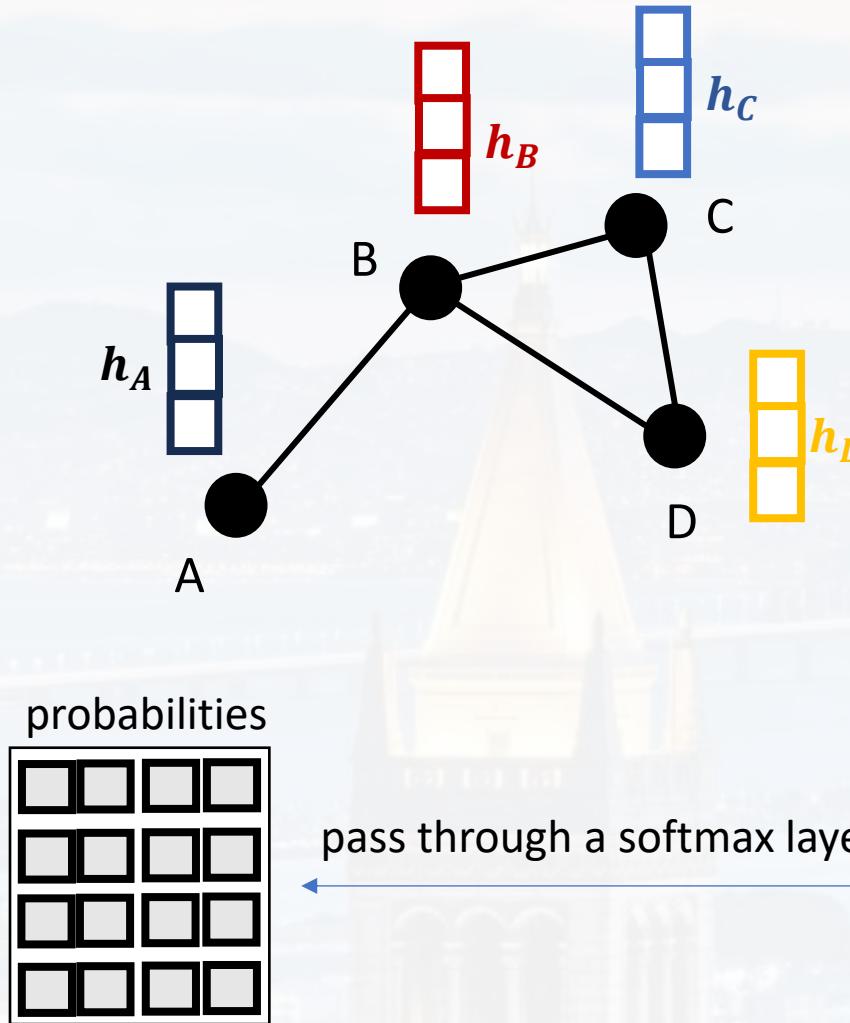


1<sup>st</sup> layer: one-hop neighborhood  
2<sup>nd</sup> layer: two-hop neighborhood  
etc

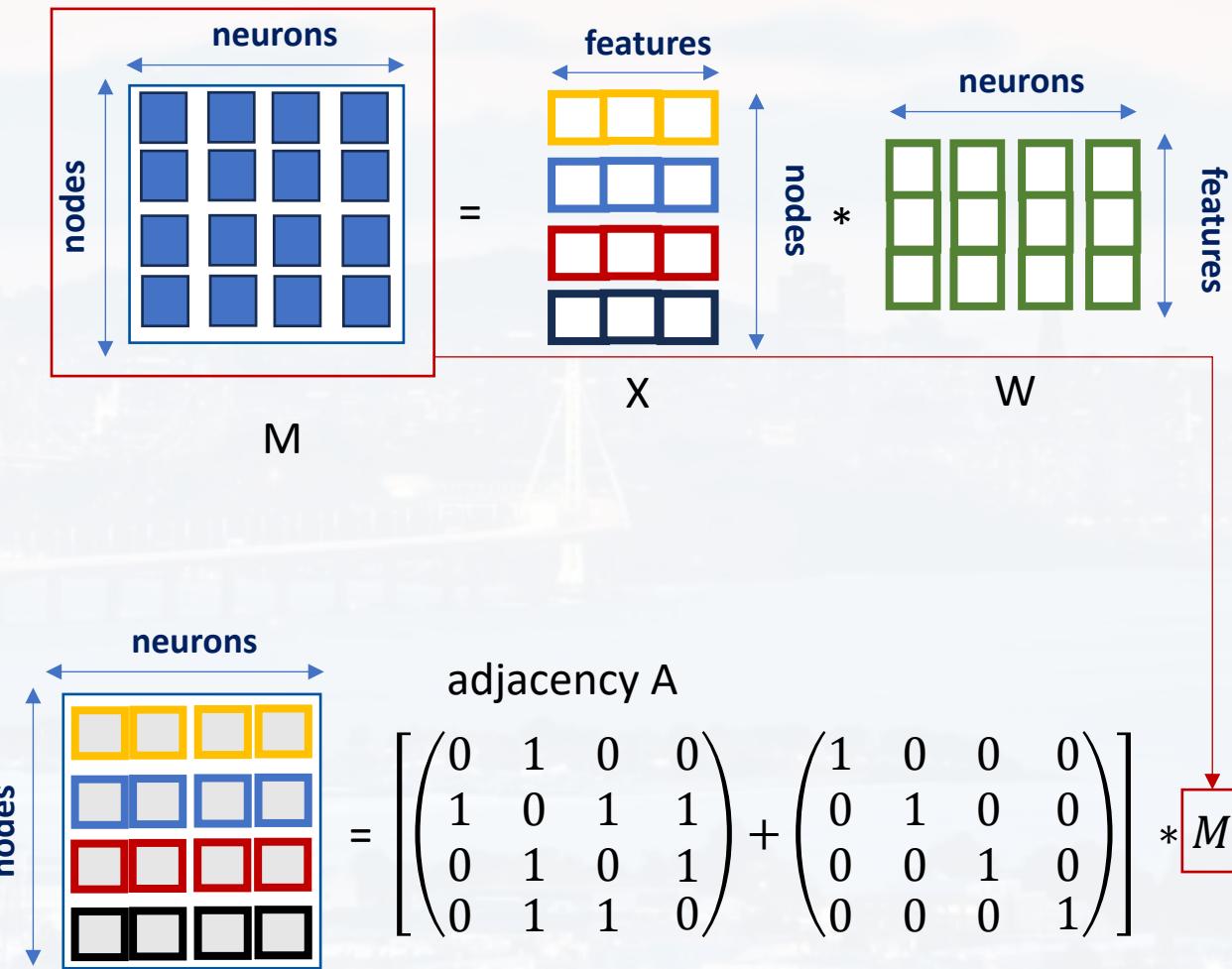
[animation here](#)



### Graph Convolution

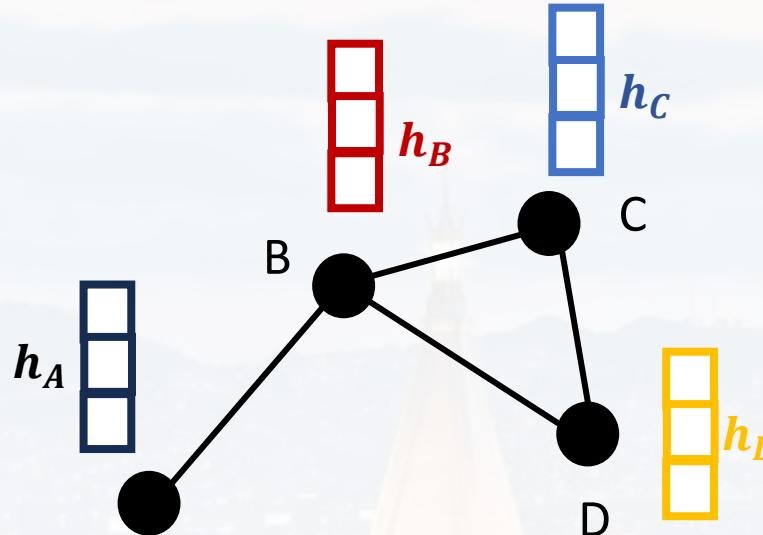


each node  $i$  has a **feature vector**  $h_i$





### Graph Convolution



### summary

- A: adjacency matrix (number of nodes x number of nodes)
- I: identity matrix (number of nodes x number of nodes)
- X: node feature matrix (number of nodes x number of features)
- W: weight matrix (number of features x number of neurons)
- $\sigma$ : any activation function
- $D^{-1/2}$ : diagonal matrix for normalization

$$H(\text{embedding}) = \sigma [ D^{-1/2} (A + I) D^{-1/2} X W ]$$

However, this would give nodes with higher degree a larger weight

→ normalizing by  $\frac{1}{\sqrt{d(n_i)}}$  and  $\frac{1}{\sqrt{d(n_j)}}$

more information [here](#)



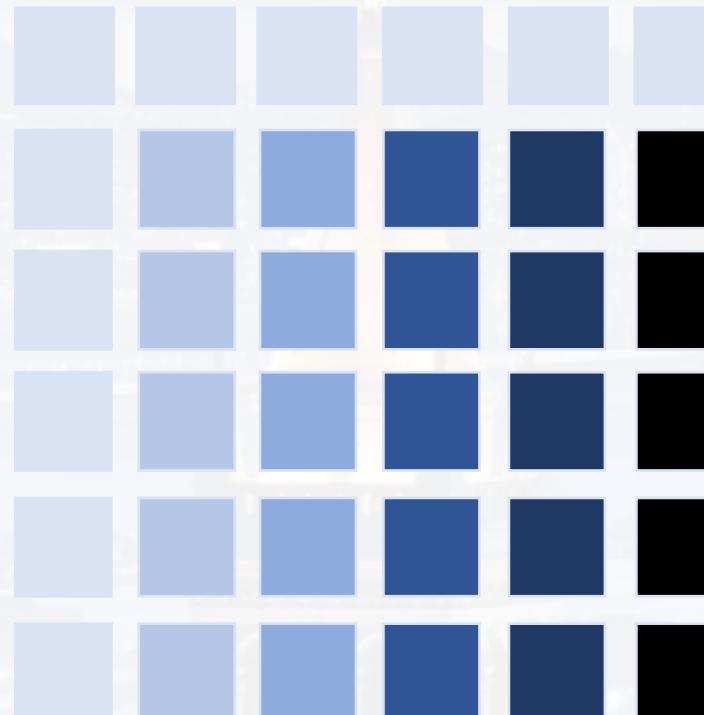
Matthew N. Bernstein



## Graph Attention

see language models (lect 15)

*“The cat jumped on the roof.”*



how the first token influences all other token

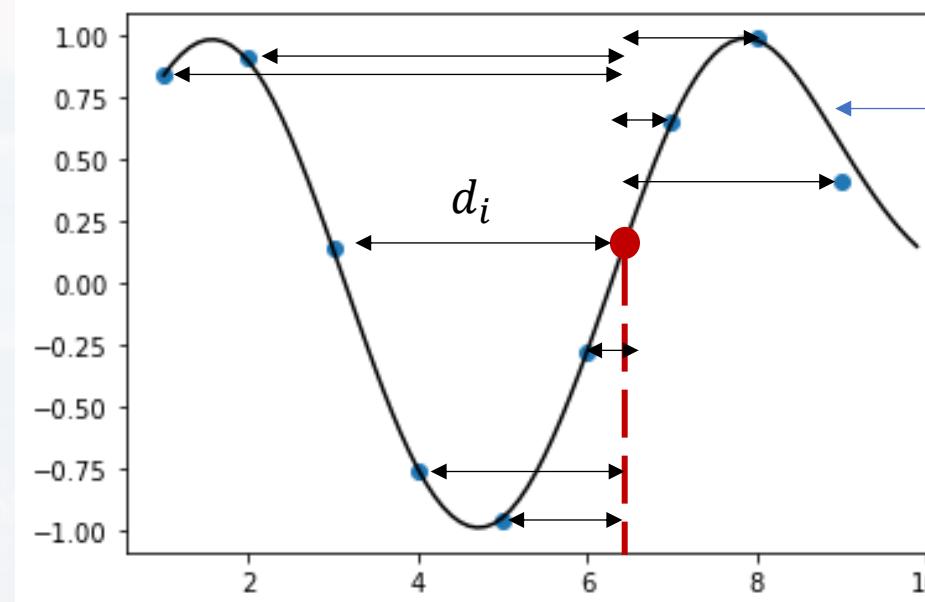


how the second token influences all other token

.... and so on



### Attention



We want to interpolate between the blue dots  
→ generating the black line  
→ **no curve fitting!**

- idea:
- select a point for which we want the interpolation for
  - calculate distance  $d_i$  to every other point
  - each data point should influence the value of the interpolated point
  - the closer, the stronger the influence → weighted mean

$$y_{int} \sim \sum_{i=1}^I w_i y_i \quad w_i \sim \frac{1}{d_i}$$



### Attention



```
D = np.tile(v, (1, len(v))) - np.tile(L.transpose(), (len(v), 1))
```

- each data point should influence the value of the interpolated point
- the closer, the stronger the influence → weighted mean

$$y_{int} \sim \sum_{i=1}^I w_i y_i$$

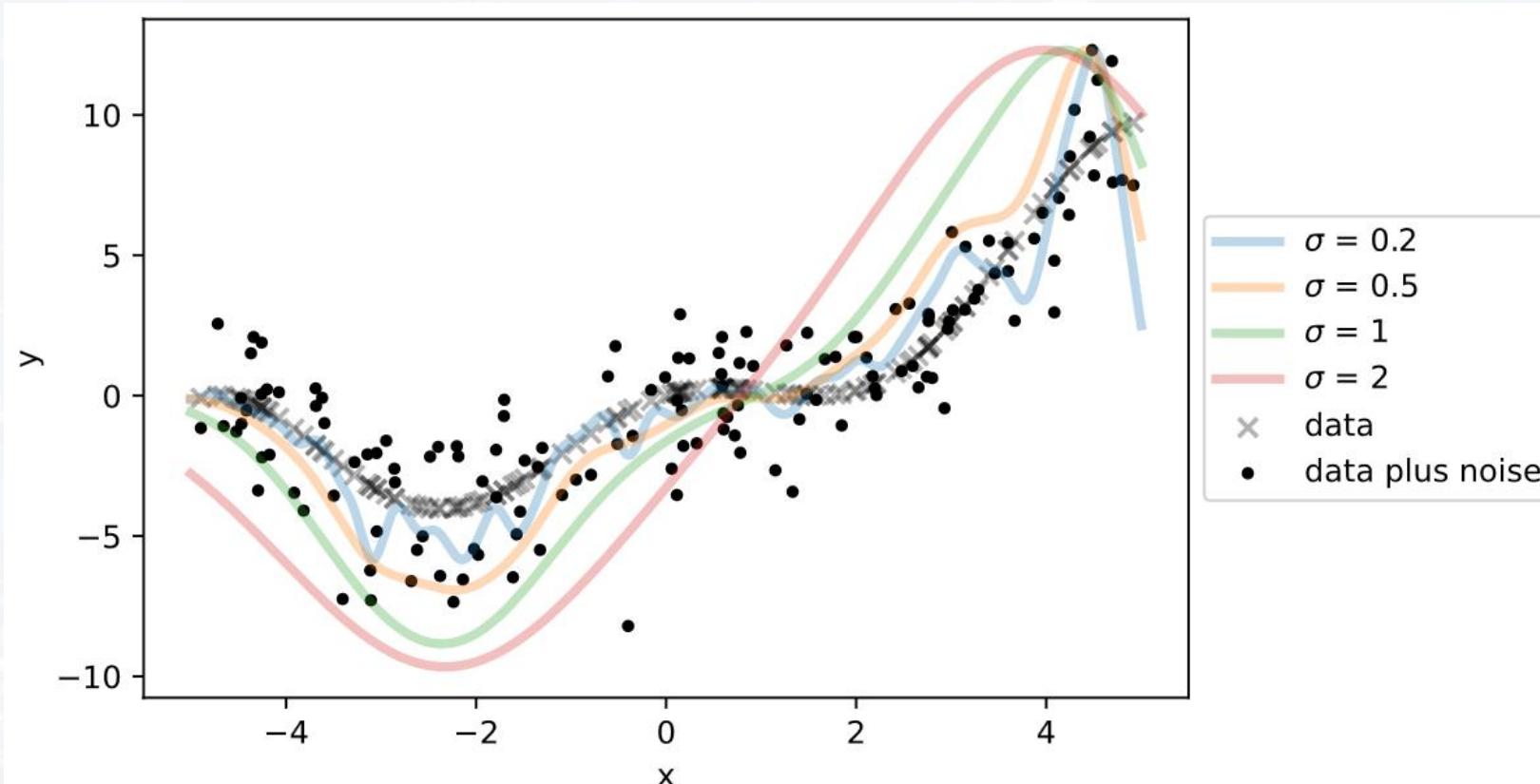
Gaussian kernel

```
w      = np.exp(-(D**2)/(sigma))
w      = w/np.sum(w + 1e-16, axis = 0)
yint  = np.dot(w.transpose(), y)
```

```
D = np.tile(V, (1, len(V))) - np.tile(L.transpose(), (len(V), 1))
```

Gaussian kernel

```
W      = np.exp(-(D**2)/(sigma))
W      = W/np.sum(W + 1e-16, axis = 0)
yint  = np.dot(W.transpose(), y)
```



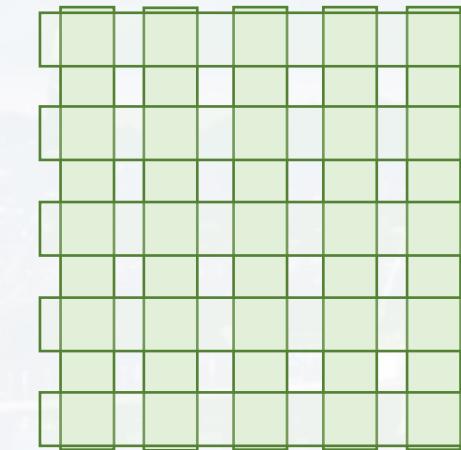
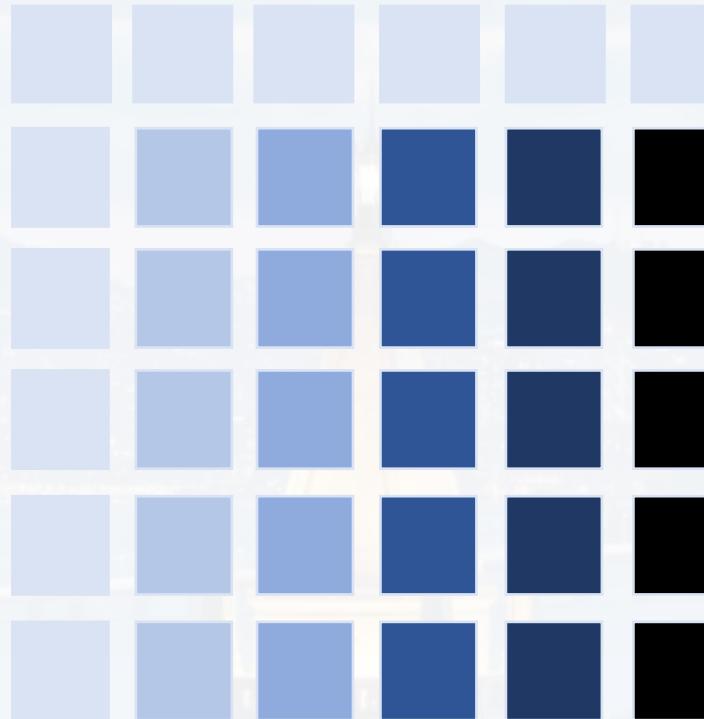
check out:

[SmoothGaussKernel.py](#)  
[SmoothExamples.py](#)



## Attention

*"The cat jumped on the roof."*



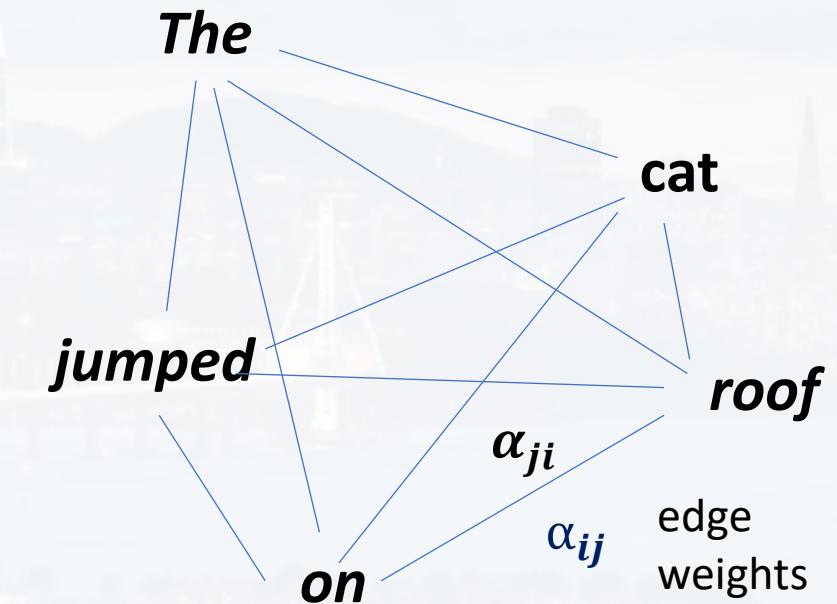
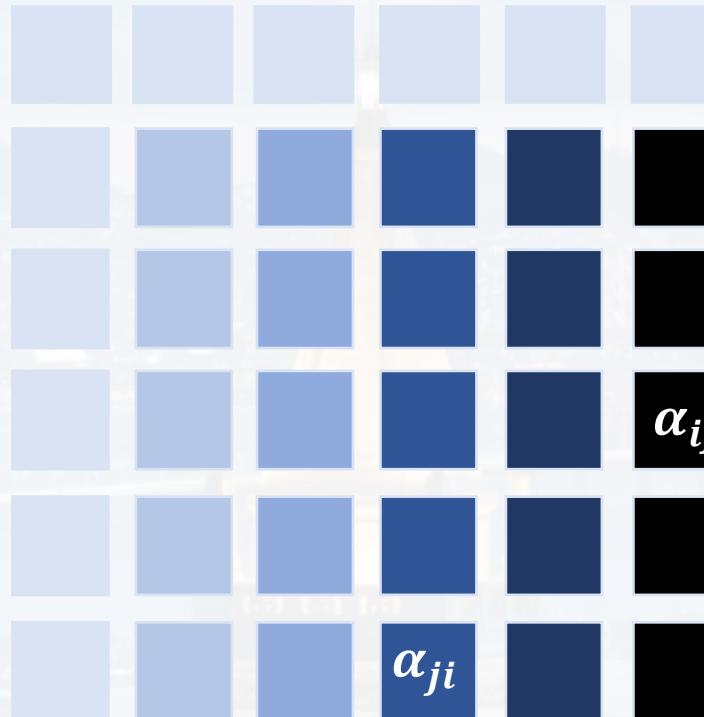
```
Gaussian kernel      W      = np.exp(-(D**2)/(sigma))  
                      W      = W/np.sum(W + 1e-16, axis = 0)  
                      yint   = np.dot(W.transpose(), y)
```

actual attention:  
these weights are learnable,  
no kernel assumed!



## Graph Attention

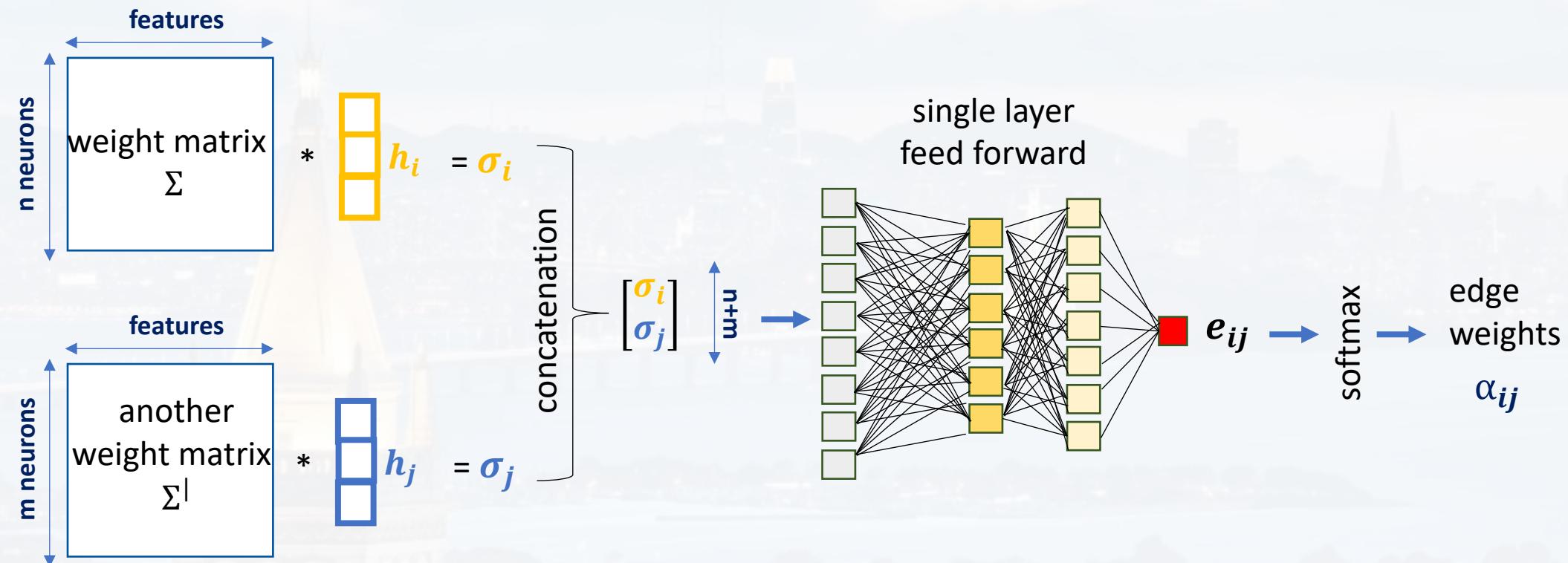
*"The cat jumped on the roof."*

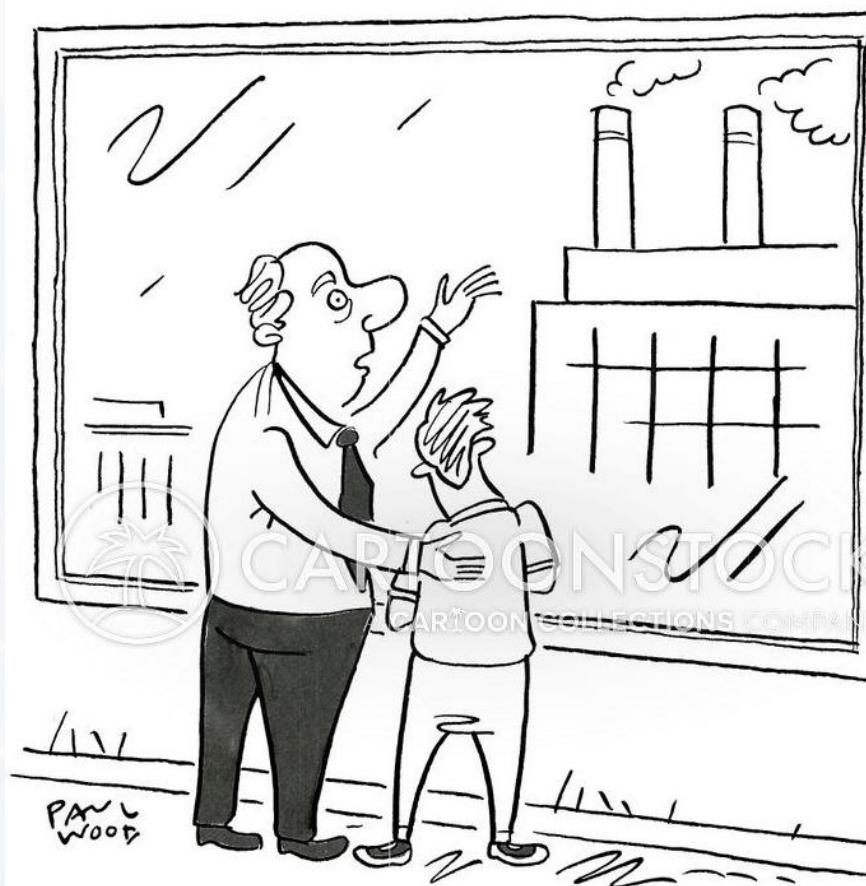




### Graph Attention

Learning the weights!  
(edge attributes)





ONE DAY SON, ALL THIS  
WILL BE RUN BY ROBOTS

## Outline

- What is a Graph
- The ANN Part
- PyTorch Example



## node classification: convolution GNN

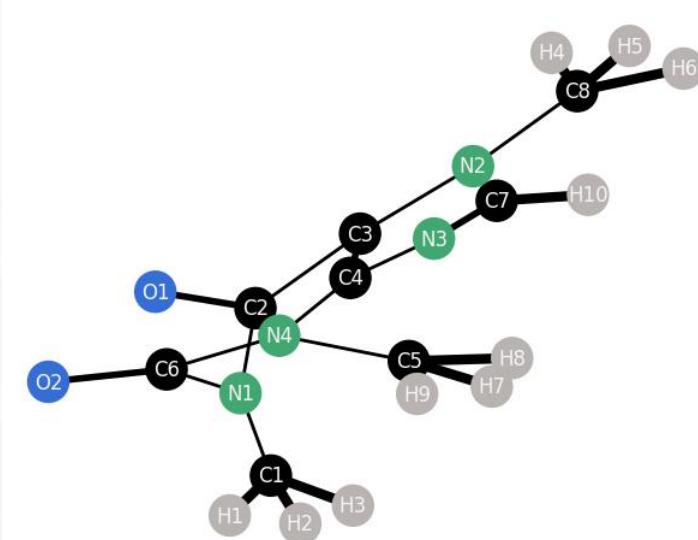
```
self.conv1 = GCNConv(n_node_features, n_neuron)  
self.conv2 = GCNConv(n_neuron, n_classes)
```

```
log_softmax(x3, dim = 1)
```

- edge weights: binding affinity

see Graph\_III.ipynb

| epoch | loss | accuracy |
|-------|------|----------|
| 0     | 1.49 | 66.67%   |
| 10    | 1.94 | 66.67%   |
| 20    | 0.17 | 79.17%   |
| 30    | 0.13 | 79.17%   |
| 40    | 0.14 | 79.17%   |
| 50    | 0.11 | 79.17%   |
| 60    | 0.11 | 79.17%   |
| 70    | 0.11 | 79.17%   |
| 80    | 0.11 | 79.17%   |
| 90    | 0.11 | 79.17%   |
| 100   | 0.11 | 79.17%   |
| 110   | 0.11 | 79.17%   |
| 120   | 0.10 | 79.17%   |
| 130   | 0.10 | 79.17%   |
| 140   | 0.10 | 79.17%   |
| 150   | 0.10 | 79.17%   |
| 160   | 0.10 | 79.17%   |



```
print(Y)  
print(Y_pred)
```

```
[0. 0. 0. 0. 0. 0. 0. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 2. 2. 2. 2. 3. 3.]  
tensor([0, 0, 0, 0, 0, 0, 1, 0, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 0, 2, 2, 0, 0, 0])
```



Thank you very much for your attention!



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WILL BE RUN BY ROBOTS