# **Statistical Network Analysis**

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Lecture 02 Foundations of Graph Theory

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- · Lecture LO2: Foundations of Graph Theory
- We introduce graph-theoretic foundations and basic mathematical notations needed throughout the course.
  - Basic definitions
  - Walks, Paths, and DAGs
  - Shortest paths, diameter, and components

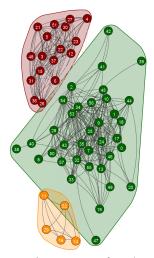
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# **Motivation**

- large volume of relational data covering
  - technical infrastructures
  - information networks
  - biological systems
  - social organizations
- network abstraction of dyadic relations
  - ► focus on topology: who links to whom
  - unified mathematical model/language
  - toolbox of algorithms and metrics

# exemplary network-analytic questions

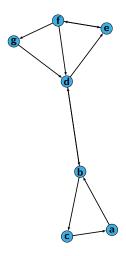
- can all students influence each other?
- which students are connected by short paths?
- can we identify natural groups?
- which student has the most central position?



**natural group structure** in student network

- The graph or network abstraction of relational data provides both a modelling
  framework as well as a common language for the study of complex systems. It enables
  the application of a toolbox of measures, algorithms, and statistics to quantitatively
  analyse systems from different disciplines. These methods allows us to answer
  questions about the topology of a graph or network, e.g.
  - 1. Are all nodes in the network connected by a path?
  - 2. Can we identify groups (or communities) of nodes that are more connected to themselves than to other nodes?
  - 3. Which nodes have the most central position in the network?
- For network models of real complex system (like, e.g., a group of students or employees, a network of documents, or users on social media) these abstract problems translate to practically relevant questions, e.g.
  - 1. Can information flow between all students? Can all students influence each other? Can an infectious disease spread to the whole population of students?
  - 2. Which groups of my employees work together closely? Which documents in this hyperlink graph address similar topics? Are there any "bottlenecks" in our communication network, i.e. are there nodes or links that are likely to be overloaded?
  - 3. Who are the most central employees in our company? How would our team be affected if a certain employee left? Whom should we target with an advertisement campaign?

# What is a network?



# graph or network

A graph or **network** is a tuple G = (V, E) where

- V is a set of vertices or nodes
- $ightharpoonup E \subseteq V imes V$  is a set of edges or **links**

 $V \times V$  denotes the Cartesian product of the node set, i.e. the set of all possible links  $(i, j) \in V \times V$ .

- we say: link (i, j) points from node i to j
- ▶ if not defined otherwise  $n := |V| \ m := |E|$
- multigraphs can have multiple links between the same nodes, i.e. E is multiset

# example network

$$V = \{a, b, c, d, e, f, g\}$$

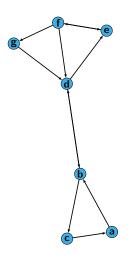
$$E = \{(a, b), (b, c), (b, d),$$

$$(c, a), (d, b), (d, e), (e, f),$$

$$(f, d), (f, e), (f, g), (g, d)\}$$

- Before we can study first network-analytic problems we formally introduce graph-theoretic foundations. In this course, we use the terms vertex and node, as well as edge and link interchangeably. This is common in the interdisciplinary network science community.
- The tuples (v, w) in the set of links E are ordered, i.e.  $(v, w) \neq (w, v)$  if  $v \neq w$ . This allows to distinguish links that have different **directionality**. We denote a link by a tuple (i, j), referring to a link that points from i to j. You sometimes find other conventions, where (i, j) refers to an edge pointing from j to i. For undirected networks  $\rightarrow$  stide 4 this does not make a difference, but it is important to clarify the notation for directed networks. Hence, we consistently use a notation where links point from the left to the right element in a tuple.
- The nodes i and i that are the endpoints of an edge (i, j) are called adjacent (from Latin "adiacere" for "border upon" or "lie near"). A link (i, j) is said to be incident on nodes i and i (from Latin "incidere" for "to fall upon").
- If not defined otherwise, we often use n to refer to the number of nodes and m to refer to the number of links. The number of different, ordered tuples between sets with n nodes is  $n^2$ , i.e. a network with n nodes can have at most  $n^2$  links.
- We can also define multigraphs where E is a multiset of links, i.e. elements in E can occur multiple times. Consequently the maximal number of links is unbounded. In this course we generally do not consider multigraphs, i.e. E is a set where elements can occur only once. For data where links are observed multiple times we can instead assign numeric edge attributes. → stide 5 on weighted graphs

# **Adjacency matrix**



▶ adjacency matrix  $\mathbf{A} \in \{0,1\}^{n \times n}$  of network G = (V, E) is a matrix with

$$A_{ij} = \left\{ egin{array}{ll} 1 & & ext{if } (i,j) \in E \\ 0 & & ext{else} \end{array} \right.$$

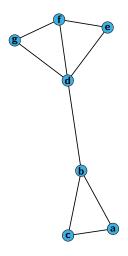
where  $A_{ij}$  refers to row i and column j

for directed networks (no self-loops)

$$|E| = m = \sum_{i,j \in V} A_{ij}$$

- Binary, square **adjacency matrices**  $\mathbf{A} \in \{0,1\}^{n \times n}$  are a simple and widely used mathematical structure to mathematically represent networks. The existence of a link (i,j) from i to j (i.e. an "adjacency") is indicated by an entry  $A_{ij} \in \{0,1\}$  in row i and column j, where 1 captures that the link is present while 0 indicates the absence of the link.
- In the example above, the adjacency matrix is not symmetric. This is due to the fact that links have a direction. For example, the link (a, b) exists, but the reverse link (b, a) does not exist. We call networks with this property directed networks. An example for a network that is naturally directed is a citation network. An article A that cites an article B does not imply that the opposite is true. In fact, except for rare cases where manuscripts were written (and published) at the same time, this cannot even happen.
- In practice, the adjacency matrices of many empirical networks are sparse matrices, i.e. there are many more 0 elements than 1 elements. This facilitates compressed representations, where only non-zero elements are actually stored.
- For the binary adjacency matrix of directed networks with no self-loops → slide 6 the sum of matrix elements corresponds to the number of links in the network. The outgoing links of node i are represented in row i of the matrix. The incoming links of node j are represented in column j of the matrix.

# **Undirected networks**



network is undirected iff

$$(i,j) \in E \Leftrightarrow (j,i) \in E$$

and directed otherwise

▶ adjacency matrices of undirected networks are **symmetric**, i.e.  $A_{ii} = A_{ii} \forall i, j \in V$ 

# adjacency matrix of undirected network

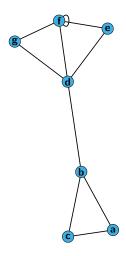
$$\mathbf{A} = \begin{bmatrix} a & b & c & d & e & f & g \\ 0 & 1 & 1 & 0 & 0 & 0 & 0 \\ 1 & 0 & 1 & 1 & 0 & 0 & 0 \\ 1 & 1 & 0 & 0 & 0 & 0 & 0 \\ 0 & 1 & 0 & 0 & 1 & 1 & 1 \\ e & 0 & 0 & 0 & 1 & 0 & 1 & 0 \\ f & g & 0 & 0 & 1 & 0 & 1 & 0 \end{bmatrix}$$

for undirected networks (no self-loops)

$$|E|=m=\frac{1}{2}\sum_{i:i\in\mathcal{V}}A_{ij}$$

- We say that a network is **undirected** iff all links exist in both directions, i.e.  $(i,j) \in E \Leftrightarrow (j,i) \in E$ . For binary adjacency matrices of undirected networks we have  $A_{ij} = A_{ji}$ , i.e. the **adjacency matrix is symmetric**. We sometimes do not explicitly differentiate between an undirected network and a directed network in which each link exists in both directions (the adjacency matrix representation of both are identical). However, we do distinguish between directed and undirected networks regarding the question what a single undirected link is. In the example above, we say that we have an undirected network with nine undirected links, rather than counting 18 directed links. This has the implication that the number of undirected links is only half of the sum of adjacency matrix entries in an undirected network.
- In the graphical representation of undirected networks we use a single undirected
  link (with no arrow heads) instead of two directed links (x, y) and (y, x). Sometimes,
  we also use mixed representations for directed networks, where an undirected link is
  a simpler notation for two directed links between the same node pair in opposite
  directions.
- Many collaboration networks are naturally undirected. If two employees A and B work together on a project, A is linked to B and B is linked to A, i.e. collaborations are symmetric.
- Most citation networks are naturally directed. Except for exceptional cases, the fact
  that article A cites article B even means that the opposite link cannot exist (since the
  directionality of links typically implies that A was published before B).

# **Self-loops**



- $\blacktriangleright$  links (i, i) are called **self-loops**
- captured in the diagonal entries of adjacency matrix A
- two different representations of self-loops:

1. 
$$(i,i) \in E \implies A_{ii} = 1$$

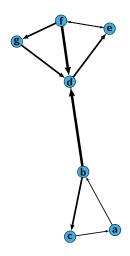
$$2. (i,i) \in E \implies A_{ii} = 2$$

# adjacency matrix of network with self-loop

$$\mathbf{A} = \begin{bmatrix} a & b & c & d & e & f & g \\ b & 1 & 1 & 0 & 0 & 0 & 0 \\ 1 & 0 & 1 & 1 & 0 & 0 & 0 \\ 1 & 1 & 0 & 0 & 0 & 0 & 0 \\ 0 & 1 & 0 & 0 & 1 & 1 & 1 \\ e & 0 & 0 & 0 & 1 & 0 & 1 & 0 \\ f & g & 0 & 0 & 1 & 0 & 1 & 0 \end{bmatrix}$$

- In the previous slide we mentioned so-called self-loops, which refers to links
   (i, i) ∈ E from node i to the node i itself. Such self-loops are captured in the
   diagonal of the adjacency matrix.
- Different from links (i,j) for  $i \neq j$  self-loops can only exist in one direction, which translates to the fact that even in directed networks there is only a single matrix entry for each self-loop. This is the reason why, in a network with self-loops, we cannot simply double the sum of matrix entries to calculate the number of edges. To account for this special characteristic, we sometimes define non-binary adjacency matrices of unweighted network, where a self-loop is represented by a 2 on the diagonal.
- Self-loops have special characteristics that can lead to complications in the definition
  of some network-analytic measures. We thus often exclude them when we analyse
  networks. Sometimes, we do however consider (or even explicitly add) them, e.g. in
  the modelling of dynamical processes. Here self-loops represent node-internal
  feedback or memory, i.e. they encode that the future state of a node is coupled to the
  previous state of that same node.

# Weighted networks



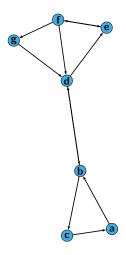
- in weighted networks links have numerical attributes  $w: E \to \mathbb{R}$  that capture strength, frequency, capacity, etc. of links
- weighted networks have a **real-valued** adjacency matrix  $\mathbf{A} \in \mathbb{R}^{n \times n}$  with

$$A_{ij} = \begin{cases} w(i,j) \text{ if } (i,j) \in E \\ 0 \text{ else} \end{cases}$$

# $\mathbf{A} = \begin{bmatrix} a & b & c & d & e & f & g \\ 0 & 1 & 0 & 0 & 0 & 0 & 0 \\ b & 0 & 0 & 2 & 3 & 0 & 0 & 0 \\ 1 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 1 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 1 & 0 & 0 & 2 & 0 & 0 & 0 \\ e & 0 & 0 & 0 & 0 & 0 & 1 & 0 \\ f & g & 0 & 0 & 0 & 2 & 0 & 0 & 0 \end{bmatrix}$

- In many networks, we want to capture additional numerical properties of links, e.g. the strength, capacity, cost or frequency of an interaction or connection. For such settings weighted networks, each edge has an additional real- or integer-valued property, the so-called link weight. Examples for properties captured by link weights include
  - the frequency or duration of contacts between actors in social networks
  - the level of trust between actors in a social network
  - the number of co-authored papers in a co-authorship network
  - the bandwidth of a network connection in a communication network
  - the number of passengers travelling on a route between two airports
  - the average cost of flights between two airports
  - the geographical distance between two stations in a train network
  - the capacity of a transmission line in a power grid
  - the trade volume in a network of financial transactions between economic actors
- We can mathematically represent a weighted network by real-valued adjacency matrices, in which non-zero entries capture the weights of links.
- Note that also networks in which all links exist in both directions (which would qualify as an undirected network) can have asymmetric adjacency matrices if the weights of links in different directions differ.

# **Node degrees**



### undirected networks

**degree**  $d(i) = d_i$  of node i is defined as

$$d_i := |\{j \in V : (i,j) \in E\}|$$

### directed networks

- indegree  $d_{in}(i)$  is the number of incoming edges, i.e.  $d_{in}(i) := |\{j \in V : (j,i) \in E\}|$
- **outdegree**  $d_{out}(i)$  is the number of outgoing edges, i.e.  $d_{out}(i) := |\{j \in V : (i, j) \in E\}|$

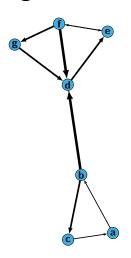
# example network

- $ightharpoonup d_{in}(f) = 1$
- $ightharpoonup d_{out}(f) = 3$

- The degree of a node i corresponds to the number of nodes to which it is directly connected. In directed networks we distinguish between indegree and outdegree. The indegree of i counts the number of predecessors, i.e. the number of nodes j for which a link (j, i) exists. The outdegree of i counts the number of sucessors, i.e. the number of nodes j for which a link (i, j) exists.
- For an undirected networks, we have  $d_{in}(i) = d_{out}(i) = d_i$  and we simply call this the degree of a node.
- Sometimes, for directed networks a **total degree**  $d_{total}(i)$  is defined as  $d_{total}(i) = d_{in}(i) + d_{out}(i)$ , i.e. the total degree counts both incoming and outgoing links.
- We can easily calculate degrees in directed and undirected networks by summing the rows/columns of their adjacency matrix. In directed networks, the outdegree of node i is the sum of entries in row i, i.e.  $d_{out}(i) = \sum_j A_{ij}$  where index j runs over the columns. The indegree of node j is the sum of entries in column i, i.e.  $d_{in}(j) = \sum_i A_{ij}$ , where index i runs over the rows. In undirected networks both yields the same value since the adjacency matrix is symmetric, i.e. we can compute the degree in either way.
- The degree sequence (or distribution) of a network is an macroscopic feature of networks, which allows us to make surprisingly strong statements about the expected properties of a network. 

  more in LOT

# Weighted node degree



# weighted degrees

for weighted networks, the **weighted in- or outdegree** of a node is the sum of incoming or outgoing link weights, i.e.

$$w_{in}(i) := \sum_{j \in V} w(j,i) = \sum_{j=1}^{n} A_{ji}$$

$$w_{out}(i) := \sum_{j \in V} w(i,j) = \sum_{j=1}^{n} A_{ij}$$

# adjacency matrix of weighted network

$$\mathbf{A} = \begin{bmatrix} a & b & c & d & e & f & g \\ 0 & 1 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 2 & 3 & 0 & 0 & 0 \\ 1 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 1 & 0 & 0 & 2 & 0 & 0 \\ e & 0 & 0 & 0 & 0 & 0 & 1 & 0 \\ f & 0 & 0 & 0 & 3 & 1 & 0 & 2 \\ g & 0 & 0 & 0 & 2 & 0 & 0 & 0 \end{bmatrix}$$

$$w_{in}(i) = \sum_{j \in V} A_{ji}, w_{out}(i) = \sum_{j \in V} A_{ij}$$

- We can extend the definition of node degrees to weighted networks by summing the weights of incoming or outgoing links.
- For a binary adjacency matrix, the weighted in-degree of a node i is the sum of entries in column i of the adjacency matrix. Conversely, the weighted out-degree of node i is the sum of entries in row i of the adjacency matrix. In undirected networks with symmetric adjacency matrices, bot are the same and we call this the weighted degree.
- Sometimes, the strength or total weighted degree of a node i in a weighted and directed network is defined as the sum w<sub>in</sub>(i) + w<sub>out</sub>(i) of the weighted in- and outdegree. However, for weighted directed networks it is more common to consider the weighted in- and out-degree separately.

# **Representing networks**

# adjacency matrix

$$\mathbf{A} = \begin{bmatrix} a & b & c & d & e & f & g \\ 0 & 1 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 1 & 1 & 0 & 0 & 0 \\ 1 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 1 & 0 & 0 & 1 & 0 & 0 \\ e & f & g & 0 & 0 & 1 & 1 & 0 & 1 \\ g & 0 & 0 & 0 & 1 & 0 & 0 & 0 \end{bmatrix}$$

# adjacency list

a,b
b,c
b,d
c,a
d,b
d,e
e,f
f,d
f,e
f,g
g,d

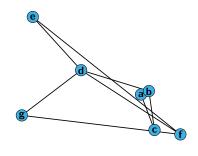
- requires  $n^2$  bits of memory
- supports algebraic operations
- suitable for small networks

- requires  $\mathcal{O}(m \log n)$  bits of memory
- ightharpoonup preferable if  $m \ll n^2$
- suitable for large networks

- We end this introductory section with a reflection on data structures that are used to
  represent networks in a computer. A widely used approach is to store a list of tuples
  that represent adjacent nodes, i.e. pairs of nodes connected by links. We call this an
  adjacency list representation. For a network with n nodes and m links we need to
  store m pairs of nodes, where each node requires log n bits.
- A second approach is to store the adjacency matrix of a network (e.g. as an array of arrays). This has the advantage that we can use standard packages to perform algebraic operations (matrix multiplication, matrix powers, eigenvector calculations, etc.) that have a natural interpretation in networks. → more in L11 . For a network with n nodes a naive binary adjacency matrix representation requires us to store n² bits, independent of the number of links that actually exist. representation.
- For fully connected networks, adjacency list and adjacency matrix representations are equally efficient in terms of memory requirements. However, for networks where the majority of node pairs are not connected, an adjacency list representation is more efficient. In particular, if the average degree of nodes is a small constant we have  $m \sim n$  and an adjacency list requires only  $\mathcal{O}(n \log n)$  bits, while a naive adjacency matrix requires  $n^2$  bits. Networks where the number of links is comparable to the number of nodes are called sparse networks. We can use **sparse matrix representations** to efficiently store matrices with many zero entries. This will be important when we apply spectral methods to large networks.
- As we shall see later, the question what is the most efficient representation of networks has interesting relations to information theory.

# **Visualizing networks**

- we need graphical representation, i.e. Euclidean representation of graph
- <u>but</u>: networks can capture arbitrary non-Euclidean topologies
- ▶ need to map nodes to positions in Euclidean space  $\mathbb{R}^d$  with  $d \in \{1, 2, 3\}$
- ightharpoonup we call  $L:V o\mathbb{R}^d$  layout of graph
- good graph layouts enable us to follow paths and recognize patterns



# example: Fruchterman-Reingold algorithm

compute stable state of multi-body simulation with

- repulsive force between all pairs of nodes
- **attractive force** between all nodes connected by an edge

→ TMJ Fruchterman, EM Reingold, 1991

# layouts used in graph drawing

- Circular layout
- Force-directed layouts
- ► Spectral layout → Lecture: ML4Nets

- The origin of the term graph is the Greek work graphos (to draw), i.e. an essential
  feature of a graph or network is that we can draw them (on paper or on a screen).
  Clearly, we can always draw nodes as circles, and connect pairs of nodes connected by
  an edge via a line. However, there are infinitely many different ways in which we can
  draw a graph or network. We are thus interested in principled methods to find a good
   or even optimal drawing of a network.
- In principle, to visualize a network, we need geometric representations of nodes and edges. However, graphs can capture arbitrary non-Euclidean topologies so mapping them to a one, two, or three-dimensional Euclidean space for the purpose of visualization is actually challenging.
- We are generally interested in mappings of nodes to coordinates such that the mapping retains as much "information" about the graph topology as possible (cf. node embedding techniques in machine learning, more in our future lecture Machine Learning for Complex Networks). In particular, a good graph layout should help us to easily follow paths along sequences of edges, and recognize clusters of nodes that are connected by many edges. For this, those nodes should be positioned close to each other. Hence we can say that for good network visualizations the notion of "similarity" captured in terms of edges between nodes should be reflected by the Euclidean distance between the geometric representations of nodes.
- There are many different algorithms that produce meaningful visual representations of networks, i.e. force-directed layouts like the Fruchterman-Reingold algorithm.

# **Practice session**

- we introduce the python-based network analysis package pathpy
- we show how to represent directed, undirected, weighted, and multi-edge networks with node/edge attributes
- we demonstrate how to read and write network data from/to files and databases
- we explain how to visualize networks with pathpy

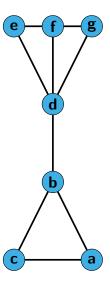
# practice session

see notebooks 02-01 - 02-03 in gitlab repository at

→ https://gitlab.informatik.uni-wuerzburg.de/ml4nets\_notebooks/2022\_wise\_sna\_notebooks

- Now that we have covered some foundations of graph theory, we move to the first practice session of our course. In the practice session, we study practical demonstrations of the theoretical concept introduced in the lecture.
- In this first session, we will show you how directed, undirected, and weighted networks are represented in the network analysis package pathpy.
- We further show how you can read/write network data from/to files or databases and how you can visualize networks based on layout algorithms.
- You can find the jupyter notebooks (and data) used in the practice sessions in an accompanying gitLab repository.

# Walks, paths and cycles



▶ sequence  $(p_0, p_1, ..., p_l)$  of nodes  $p_i \in V$  is a walk from  $p_0$  to  $p_l$  iff

$$(p_i, p_{i+1}) \in E$$
 for  $i = 0, ..., l-1$ 

- walk  $(p_0, p_1, ..., p_l)$  is a **(simple) path** iff  $p_i \neq p_i$  for  $0 \le i, j \le l$  and  $i \ne j$
- ightharpoonup walk  $(p_0, p_1, \dots, p_l)$  is a **cycle** iff
  - 1.  $p_0 = p_I$
  - 2.  $p_i \neq p_j$  for 0 < i, j < l and  $i \neq j$
- length of path, walk, or cycle is defined as  $len(p_0, ..., p_l) := l$

### example

- $\triangleright$  (a, b, a, b, d) is a walk
- $\triangleright$  (a, b, c, a) is a cycle
  - (a, b, d, g) is a path of length three from a to g

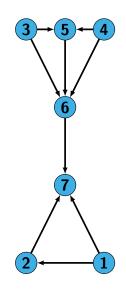
- Arguably, the main (if not only) reason why we are interested in networks is because
  they allow us to understand how the elements of a complex system can directly and
  indirectly influence each other via sequences of links. A sequence of nodes where any
  two consecutive nodes are adjacent is called a walk. Walks can contain the same node
  multiple times like, e.g., as in the example below.
- A walk (p<sub>0</sub>,..., p<sub>l</sub>) where all nodes are different is called a path from node p<sub>0</sub> to node p<sub>l</sub>. The terms "walk" and "path" are often used synonymously, in which case we call a path where all nodes are different simple path.
- A walk where only the startpoint p<sub>0</sub> and the endpoint p<sub>1</sub> are identical is called a cycle.
   A network that contains at least one cycle is called a cyclic network. If a network contains no cycle we call it acyclic network.
- We define the **length of a walk, path or cycle** as the number of traversed links (i.e. the number of traversed nodes minus one). Hence, a single edge  $(i,j) \in E$  (with  $i \neq j$ ) defines a path of length one that connects node i to node j. In communication networks, this definition of path lengths has a natural interpretation as the number of hops a message takes from the origin to the destination.
- For any two nodes, there can be many different paths by which the same pair of nodes
  is indirectly connected. In the example network, there is only a single path of exactly
  length three from node a to node g, but there is another path of length four that first
  traverses node c. This shows that a path between two nodes does not necessarily
  need to follow the shortest sequence of links.

# **Directed Acyclic Graphs**

- (directed) network with no cycle is called directed acyclic graph (DAG)
- ▶ for any DAG G = (V, E) we can find (at least) one **topological ordering of nodes**, i.e. mapping  $T : V \to \mathbb{N}$  such that T(v) < T(w) for all  $(v, w) \in E$
- we can compute topological ordering in linear time

# topological ordering algorithms

- ► Kahn's algorithm → AB Kahn, 1962
- DFS with edge classification
  - → Cormen, Leiserson, Rivest, Stein, 2001
- DAGs are important concept in scheduling, path analysis, causal inference, temporal network analysis

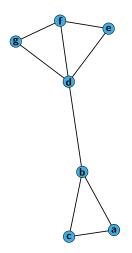


example

topological ordering of nodes in DAG

- Considering the definition of a cycle, we immediately see that any undirected network
  necessarily has cycles, since each edge (v, w) gives rise to two cycles v, w, v and
  w, v, w. For directed networks, a cycle may or may not exist depending on the
  topology of the network. We call a directed networks that has no cycle a directed
  acyclic graph, short DAG.
- Directed acyclic graphs play an important role whenever we need to model events
  that can causally influence each other. This explains why they play an important role
  in scheduling (e.g. finding transaction schedules in parallel processing), in the analysis
  of causal effects based on time series data, or in the analysis of temporal networks
  such as citation graphs.
- An important characteristic of directed acyclic graphs is that we can find at least one topological ordering of nodes, i.e. a mapping of nodes to numbers such that edges only point from smaller to larger numbers. If we consider edges as causal relationships between nodes representing events, these numbers give possible temporal orderings of events (i.e. for an event A that can influence event B, event A must occur before B). In the parallel executing of interdependent transactions, temporal orderings of directed acyclic graphs are used to find suitable executing schedules that do not violate serializability (see course on databases or operating systems).
- As you know from the introductory course on algorithms and data structures, there
  are linear-time algorithms to find topological orderings in DAGs. We can also use
  those algorithms to test whether a graph is acyclic or not.

# **Topological distance**



- distance dist(v, w) between nodes v and w is the minimum length of any path between v and w
- ▶  $\operatorname{dist}(v,w) := \infty \Leftrightarrow \nexists$  path from v to w
- ▶ path  $(p_0, ..., p_l)$  is **shortest path** iff  $len(p_0, ..., p_l) = dist(p_0, p_l)$
- for weighted network  $(p_0, \ldots, p_l)$  is cheapest path iff  $\sum_{i=1}^{l} w(p_{i-1}, p_i)$  is minimal

# example

- b dist(a, d) = 2
- $\triangleright$  shortest path: (a, b, d)

- Networks define a discrete topological space in which we can calculate a measure of
  topological distance between any pair of nodes. The topological distance between a
  node v and a node w is the minimal length of any path that connects them. We call
  the distance between the nodes the shortest path length and each path with that
  length that connects those nodes is called a shortest path.
- In the example above, there is only a single path of length two from node a to node d
  and this path is the shortest path, so the distance between those two nodes is two.
   The shortest path is not necessarily unique, i.e. different shortest paths of the same
  length can exist for a given pair of nodes. We will see that the distribution of shortest
  path lengths is an important macroscopic characteristic of complex networks.
- In many systems (e.g. communication networks) finding shortest paths is a key problem, which must be solved in order to provide routing services. Indeed, the main task of routing algorithms on the Internet is to identify optimal communication paths between computer networks. As a first approximation, best can be thought of as shortest but we can also include link costs represented as link weights in a weighted network. Here, we can extend the definition of shortest path to cheapest path, i.e. paths where the sum of edge weights from v to w is minimal.
- A number of algorithms have been proposed to compute (i) shortest paths from one node to all other nodes (single-source problem), (ii) between all node pairs (all-pairs problem), or (iii) cheapest paths in networks with positive/negative weights. I assume that you studied those algorithms in depth in your BSc courses.

# **Finding shortest paths**

# Dijstra's algorithm

ightarrow EW Dijkstra, 1959

- single-source shortest paths for graphs with positive edge weights
- repeatedly relax path length for neighbors of first node in priority queue
- ▶ worst-case time complexity  $\mathcal{O}(m + n \cdot \log n)$

# **Bellman-Ford algorithm**

ightarrow R Bellman, 1958

- single-source shortest paths for graphs with real edge weights (no negative cycles)
- repeatedly relax path length of node v for all edges (v, w)
- worst-case complexity  $\mathcal{O}(n \cdot m)$

# Floyd-Warshall algorithm

ightarrow RW Floyd, 1962

- all-pairs shortest paths for graphs with real edge weights (no negative cycles)
- test triangle inequality for all triples of nodes
- worst-case complexity  $\mathcal{O}(n^3)$



# Richard E. Bellman

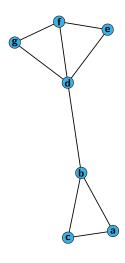
1920 - 1984

John von Neumann Theory Prize 1976

image credit: Wikipedia, Fair use

- A number of algorithms have been proposed to compute (i) shortest paths from one node to all other nodes (single-source problem), (ii) between all node pairs (all-pairs problem), or (iii) cheapest paths in networks with positive/negative weights. I assume that you studied those algorithms in depth in your BSc courses and we will thus not discuss them in detail here.
- In the practice session, we provide implementations of the three key algorithms
  mentioned above and we demonstrate how we can use them to calculate and
  reconstruct shortest and cheapest paths using pathpy and how to calculate diameter
  and average shortest path length.

# Diameter and average shortest path length



▶ **diameter** diam(G) of network G = (V, E) is length of the longest shortest path

$$\mathsf{diam}(G) := \max_{v,w \in V} \mathsf{dist}(v,w)$$

▶ average shortest path length  $\langle I \rangle$  is the average distance between nodes, i.e.

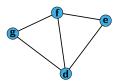
$$\langle I \rangle := \frac{1}{|V|^2} \sum_{(v,w) \in V \times V} \mathsf{dist}(v,w)$$

# example

- ightharpoonup diam(G) = 3
- ► ⟨*I*⟩ = 1.59

- The diameter is a simple but important systemic or collective property of complex networks. Why do we call this a systemic or collective feature? Because it results from the global topology of the network, rather than from a feature of any particular node or link. This is particularly true for large networks, where changing a single node or link is likely to not change the diameter.
- The diameter is defined as the length of the longest shortest path, i.e. it is the upper bound for the number of links that need to be traversed for one node to indirectly influence another node in the system. If we study the propagation of information starting from a node, the network diameter tells us how long it will take at maximum for the last node to see the information, if that information is passed along the shortest paths in the network.
- The upper bound for the shortest path length can be seen as a "worst-case estimate" that does not tell us anything about "typical shortest path lengths". We can instead characterize the distribution of path lengths by the average shortest path length. This provides a single statistical feature that captures one important aspect of the topology of a complex network. It tells us how many steps it will take on average for one node to be able to indirectly influence another node in the system. For some networks the diameter can be large, even if most shortest path lengths are actually very small. In these cases, the average shortest path length is a better characterization for the shortest path structures in a network.

# **Connected components**





- ▶ undirected network G = (V, E) is **connected** if dist $(v, w) < \infty$  for all  $v, w \in V$
- **connected components** of G = (V, E) are maximally connected subgraphs G' = (V', E') with  $V' \subseteq V$  and  $E' \subseteq E$
- size of connected component G' = (V', E') is |V'|
- ▶ largest connected component G' is called giant connected component iff

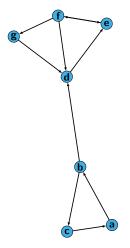
$$rac{|V'|}{|V|}pprox 1$$

# example

- connected component {a, b, c}
- largest connected component
  {d, e, f, g}

- One of the most important characteristics of a network is whether all nodes can
  actually influence each other, i.e. whether all nodes are connected via a path. If this is
  the case (i.e. when all distances are finite) we say that the network is connected.
- For networks which are not connected, we are often interested in the connected
  components, i.e. a partition into the largest subsets of nodes for which all pairs of
  nodes are connected by a path. As we will see in the practice session, we can
  efficiently calculate the connected components of a graph using Tarjan's algorithm.
- A largest connected component of a network is any connected component that
  contains a maximum number of nodes. We say that the largest connected component
  is a giant connected component if it contains almost all of the nodes (i.e. it is much
  larger than the second-largest component).
- The exact definition of a giant connected component depends on the context: For theoretical studies of random graph models with a variable number of nodes n we often call the largest connected component G' a giant connected component if  $\frac{|V'|}{|V|} \to 1$  for  $n \to \infty$ .  $\to$  see L07
- For finite-size empirical networks, we often use a reasonable threshold (like e.g. a fraction of 0.95 or 0.99), i.e. we accept that there is a small fraction of disconnected nodes. In these cases, we often disregard those nodes and study the largest connected component as a model of the system in question. In some network analysis packages "giant connected component" is used as a synonym for "largest connected component".

# **Connected components in directed graphs**



- we distinguish between strongly and weakly connected directed networks
- directed network is weakly connected iff corresponding undirected network is connected
- ▶ directed network G = (V, E) is **strongly connected** iff  $dist(v, w) < \infty \quad \forall v, w \in V$
- ▶ strongly connected components of G = (V, E) are maximal strongly connected subgraphs G' = (V', E') with  $V' \subseteq V$  and  $E' \subseteq E$

# example

- weakly but not strongly connected
- strongly connected components {a, b, c}, {d, e, f, g}

- The definitions of connectedness and connected components in undirected networks can be extended to directed networks in a natural way. In an undirected network, we can traverse any path  $(p_0, \ldots, p_l)$  in both directions, which implies that for any pair of nodes we have dist(v, w) = dist(w, v). A trivial consequence is that  $dist(v, w) < \infty \Leftrightarrow dist(w, v) < \infty$ .
- For directed networks a path p<sub>0</sub>,..., p<sub>l</sub> may not be a path if we reverse the order of nodes. Hence, we generally have dist(v, w) ≠ dist(w, v) and we can have the dist(v, w) < ∞ while dist(w, v) = ∞, i.e. v is connected to w via path, but w is not connected to v via a path. For directed networks, we thus distinguish between strongly and weakly connected networks.</li>
- A directed network is called weakly connected if the undirected network that we obtain by replacing every directed link (v, w) ∈ E with a corresponding undirected link is connected. In a connected network, for each pair of nodes v, w we have that there is either a path from v to w or a path from w to v (or both).
- A directed network is called **strongly connected** if all pairs of nodes are connected by paths, i.e. for every pair  $v, w \in V$  we have  $dist(v, w) < \infty$ . This is the directed equivalent of a connected undirected network, where all nodes are connected to each other via a path. Every strongly connected network is necessarily weakly connected.
- Similar as in undirected networks, we can define the strongly connected components
  of a directed network as the maximal strongly connected subgraphs.

# **Practice session**

- we implement three key algorithms to calculate shortest paths
- we use Tarjan's algorithm to calculate maximally connected subgraphs
- we calculate diameter, component sizes and average path length of empirical networks



# practice session

see notebooks 02-04 and 02-05 in gitlab repository at

→ https://gitlab.informatik.uni-wuerzburg.de/ml4nets\_notebooks/2022 wise sna notebooks

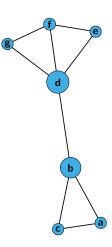
- In the second practice session, we show how to implement two basic shortest path
  algorithms in pathpy, Dijstra's algorithm for the single-source-shortest-path problem
  as well as the Floyd-Warshall algorithm for the all-pairs-shortest-path problem.
- We then use Tarjan's algorithm to calculate connected components in directed and undirected networks.
- We finally use those algorithms to compute the diameter, component sizes, and average shortest path lengths of empirical networks.

# In summary

- we introduced fundamental definitions and concepts of graph theory
- we showed how to construct, read, write, and visualize networks with pathpy
- we repeated foundational algorithms to compute shortest paths and connected components

# open questions

- how can we detect cluster patterns in large networks?
- how can we assess the importance of nodes in a network?



- In today's lecture we have introduced foundational concepts of network analysis and graph theory, such as directed, undirected, and weighted networks, node degrees, walks and paths, the adjacency matrix and the meaning of its powers as well as connected components. These concepts are the basis for advanced methods that we will introduce in the following weeks.
- We have further introduced the notion of community structures. This is an example for an important network-analytic problem that we will address in the coming week.

# **Self-study questions**

- Give examples for complex systems that can naturally be represented by directed, undirected, or weighted networks.
- 2. Give an example for a walk, path, and cycle in a network.
- 3. Give an example for an acyclic network. How can we detect whether a network is acyclic?
- Summarize the advantages and disadvantages of adjacency matrix and adjacency list representations.
- 5. Investigate how the number of edge crossings be used to evaluate graph layouts?
- 6. Define the average shortest path length of a network.
- 7. What is the computational complexity of calculating the *k*-th power of a network with *n* nodes.
- 8. Given an example for a weakly connected network where all nodes are in different strongly connected components.

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