

### Networks in R

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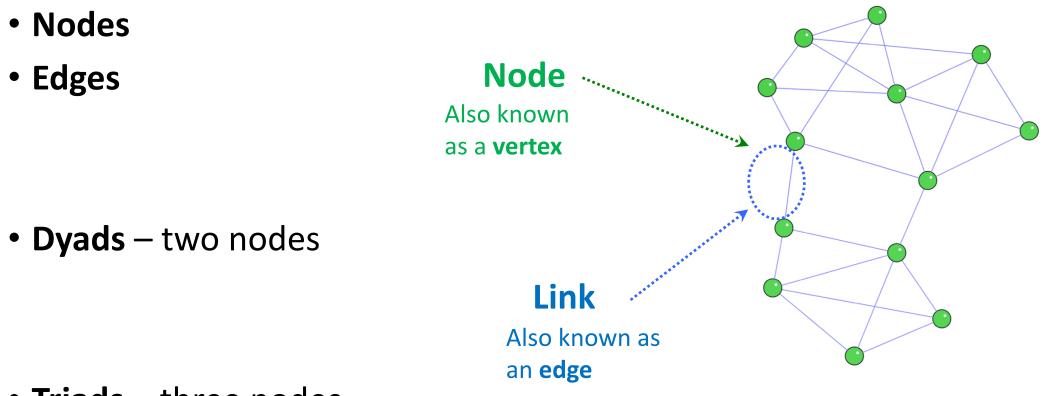




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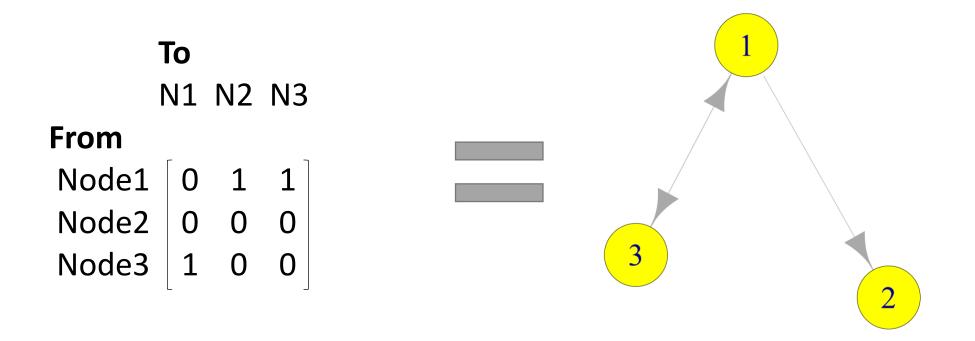
#### What is a network? Some key terms



• Triads – three nodes

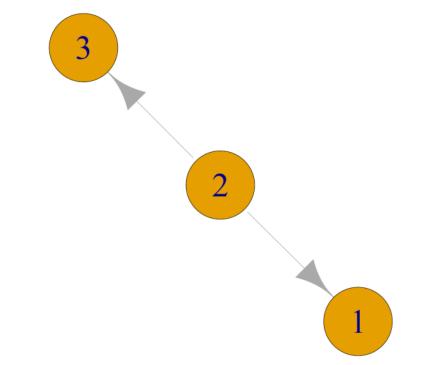
### A network is basically an adjacency matrix

 An adjacency matrix is a square matrix that describes a network, where each element of the matrix indicates whether a link exists between two nodes (and potentially also includes information about the weight associated with that link)



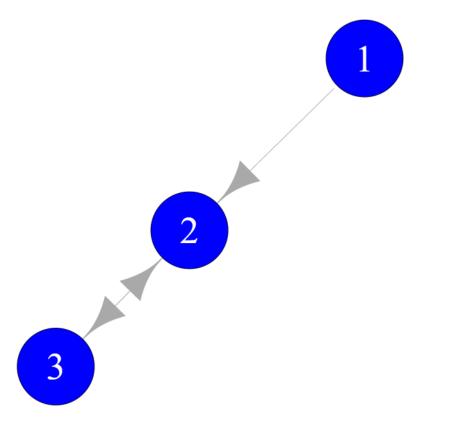
## Can you draw what this network would look like?

ToN1N2N3FromN100N210N300



How about this adjacency matrix?

To N1 N2 N3 From N1  $\begin{bmatrix} 0 & 1 & 0 \\ N2 & 0 & 1 \\ N3 & 0 & 1 & 0 \end{bmatrix}$ 

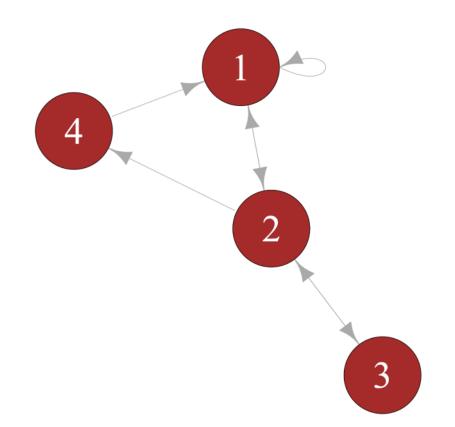


#### Network example 3

To N1 N2 N3 N4

From

N1	<b>1</b>	1	0	0
N2	1	0	1	1
N2 N3 N4	0	1	0	0
N4	1	0	0	0



#### Directed or undirected network?

То				То	То		
1	N1	N2	N3	N1 N2 N3			
From				From			
N1	0	1	0	N1 0 1 0			
N2				N2 1 0 1			
N3	0	1	0	N3 0 1 0			

#### Symmetric!

# Examples of general questions to ask with networks

- What nodes and links have particularly important roles in the network?
- Are there clusters of nodes?
- How "complex" is the network?
- What would strengthen or weaken the network?

Today we will do most of our work in the igraph package in R

- igraph reference manual
  - <u>http://igraph.org/c/doc/igraph-introduction.html</u>
  - Definitely a reference more detail than you might want to read through for a first try
- Katya Ognyanova has prepared some user friendly introductions
  - Network visualization: <a href="http://kateto.net/network-visualization">http://kateto.net/network-visualization</a>
  - igraph in R: <u>http://kateto.net/networks-r-igraph</u>

#### Let's move to the markdown! kelseyandersen.github.io/NetworksPlantPathology/

Networks in Plant Pathology	Welcome Introduction to R - Networks in R - Additional Resources -		
Networks with igraph Network data types Describing networks Bonus: Does my network deviate from random?	Networks in R         Epidemic Networks         Microbiome Networks         Microbiome Networks         Microbiome Networks         String Networks with igraph         First, make sure you have loaded package igraph.         #install.packages("igraph")         library(igraph)		
	<pre>## The following objects are masked from 'package:dplyr': ## ## as_data_frame, groups, union</pre>		
	<pre>## The following objects are masked from 'package:purrr': ## ## compose, simplify</pre>		