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# Drawing Networks

## Using qgraph

Network Analysis 2019

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Get the latest version of R from [www.r-project.org](http://www.r-project.org) and the latest version of **qgraph** from CRAN:

```
install.packages("qgraph", dep=TRUE)
```

Make sure you can load qgraph:

```
library("qgraph")
```

And that you have version 1.6.3:

```
packageDescription('qgraph')$Version  
  
## [1] "1.6.3"
```

If this fails, make sure you have the latest (3.6.1) version of R and that all depended/imported/suggested packages are installed (see CRAN).

# The qgraph () function

- The main function in **qgraph** is `qgraph()`
  - Most other functions are either wrapping functions using `qgraph()` or functions used in `qgraph()`
- The `qgraph()` function requires only one argument (`input`)
- A lot of other arguments can be specified, but these are all optional

## Usage:

```
qgraph( input, ... )
```

## Weights matrices

- The `input` argument is the input. This can be a *weights matrix*
- A weights matrix is a square  $n$  by  $n$  matrix in which each element indicates the relationship between two variables
- Any relationship can be used as long as:
  - A 0 indicates no relationship
  - Absolute negative values are similar in strength to positive values
- We will first look at unweighted graphs, in which case the weights matrix is the same as an *adjacency matrix*
  - A 1 indicates a connection
  - A 0 indicates a connection
  - Rows indicate the node of origin
  - Columns indicate the node of destination
  - By default the diagonal is omitted
  - By default, a symmetrical weights matrix is interpreted as an undirected graph

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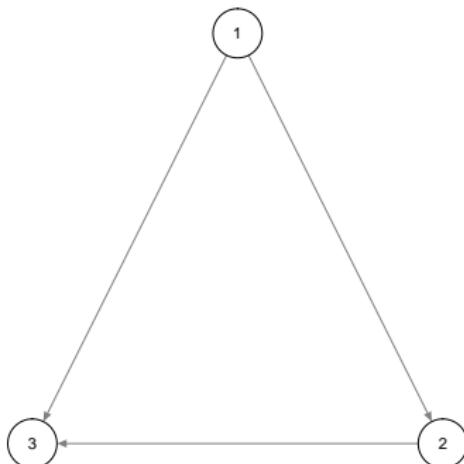
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# Weights matrices

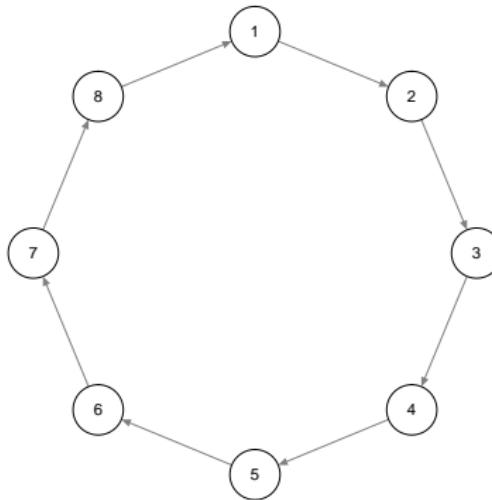
**qgraph**(input)

```
input <- matrix(c(  
  0, 1, 1,  
  0, 0, 1,  
  0, 0, 0), 3, 3, byrow=TRUE)  
print(input)  
  
##      [,1] [,2] [,3]  
## [1,]     0    1    1  
## [2,]     0    0    1  
## [3,]     0    0    0
```



# Weights matrices

Exercise: Create this graph



The layout should be right automatically, only use one argument in qgraph ()

# Weights matrices

To make this graph, we need this matrix:

```
##      [,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8]
## [1,]     0    1    0    0    0    0    0    0
## [2,]     0    0    1    0    0    0    0    0
## [3,]     0    0    0    1    0    0    0    0
## [4,]     0    0    0    0    1    0    0    0
## [5,]     0    0    0    0    0    0    1    0
## [6,]     0    0    0    0    0    0    0    1
## [7,]     0    0    0    0    0    0    0    1
## [8,]     1    0    0    0    0    0    0    0
```

# Weights matrices

These matrices become quite large, so manually defining the matrix is not effective. So some tricks are needed to make the matrix:

```
input <- matrix(0, 8, 8)
input[1,2] <- 1
input[2,3] <- 1
input[3,4] <- 1
input[4,5] <- 1
input[5,6] <- 1
input[6,7] <- 1
input[7,8] <- 1
input[8,1] <- 1
```

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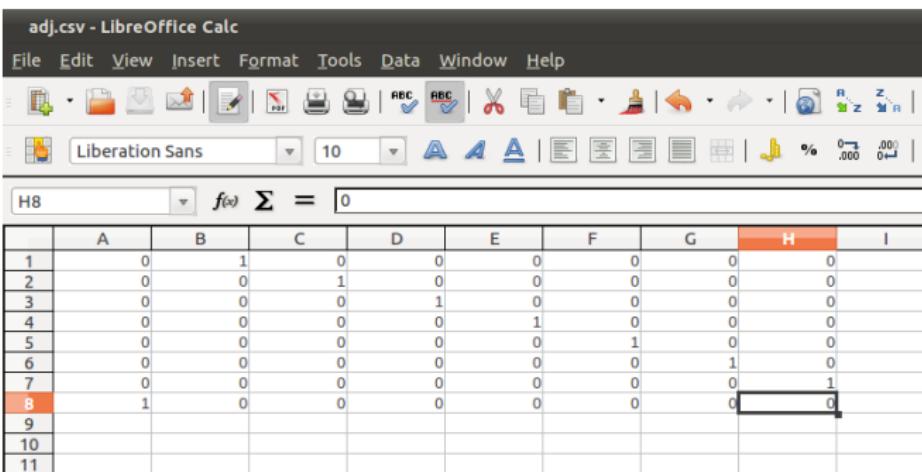
# Weights matrices

```
print(input)
```

```
##      [,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8]
## [1,]     0    1    0    0    0    0    0    0
## [2,]     0    0    1    0    0    0    0    0
## [3,]     0    0    0    1    0    0    0    0
## [4,]     0    0    0    0    1    0    0    0
## [5,]     0    0    0    0    0    0    1    0
## [6,]     0    0    0    0    0    0    0    1
## [7,]     0    0    0    0    0    0    0    1
## [8,]     1    0    0    0    0    0    0    0
```

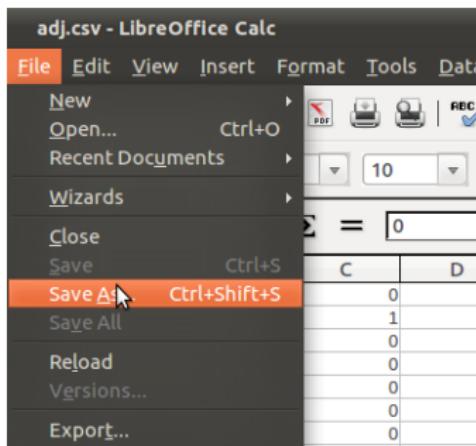
## Weights matrices

Or read the matrix from a text file! First make the matrix in a spreadsheet program (here LibreOffice)



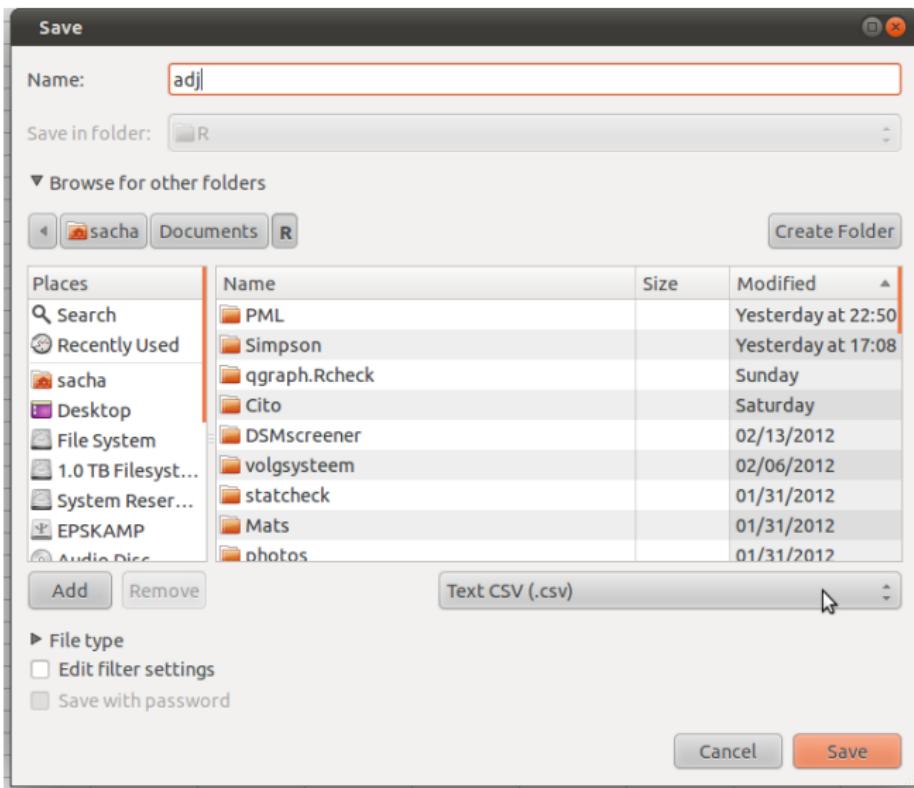
# Weights matrices

Next save as or export



# Weights matrices

Save as CSV (comma delimited text file) or tab delimited:



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# Weights matrices

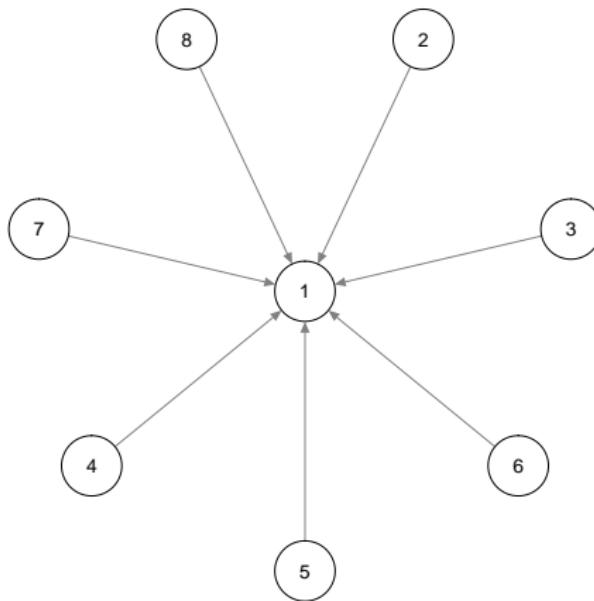
Read in R (for tab delimited use `read.table()`):

```
input <- read.csv("adj.csv", header=FALSE)  
print(input)
```

```
##      V1  V2  V3  V4  V5  V6  V7  V8  
## 1    0   1   0   0   0   0   0   0  
## 2    0   0   1   0   0   0   0   0  
## 3    0   0   0   1   0   0   0   0  
## 4    0   0   0   0   1   0   0   0  
## 5    0   0   0   0   0   0   1   0  
## 6    0   0   0   0   0   0   0   1  
## 7    0   0   0   0   0   0   0   1  
## 8    1   0   0   0   0   0   0   0
```

# Weights matrices

Exercise: Create this graph



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# Weights matrices

```
##      [,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8]
## [1,]     0   0   0   0   0   0   0   0
## [2,]     1   0   0   0   0   0   0   0
## [3,]     1   0   0   0   0   0   0   0
## [4,]     1   0   0   0   0   0   0   0
## [5,]     1   0   0   0   0   0   0   0
## [6,]     1   0   0   0   0   0   0   0
## [7,]     1   0   0   0   0   0   0   0
## [8,]     1   0   0   0   0   0   0   0
```

# Weighted graphs

- Specify edge weights to make a graph weighted
  - In a weights matrix: simply specify other values than only zeros and ones
- An edge weight of 0 indicates no connection
- Positive and negative edge weights must be comparable in strength

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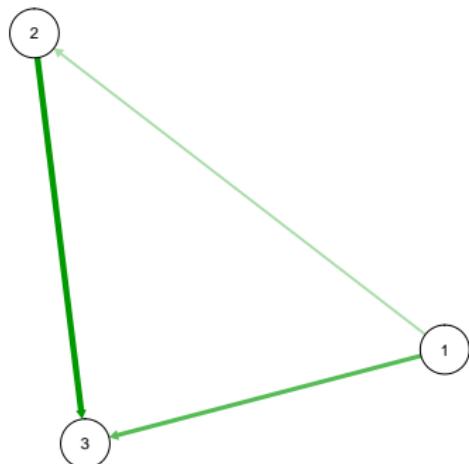
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# Weighted graphs

**qgraph**(input)

```
input<-matrix(c(  
  0,1,2,  
  0,0,3,  
  0,0,0),3,3,byrow=TRUE)  
print(input)
```

```
##      [,1] [,2] [,3]  
## [1,]     0    1    2  
## [2,]     0    0    3  
## [3,]     0    0    0
```



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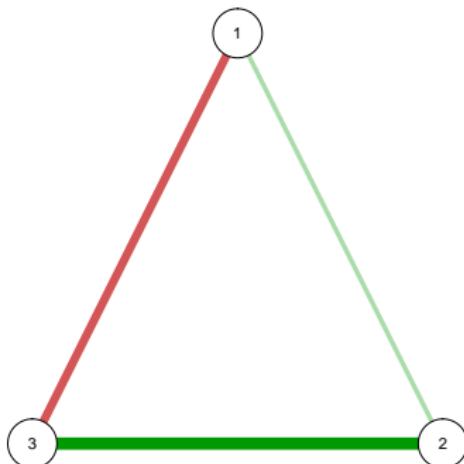
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# Weighted graphs

**qgraph**(input)

```
input<-matrix(c(  
  0,1,-2,  
  1,0,3,  
 -2,3,0),3,3,byrow=TRUE)  
print(input)
```

```
##      [,1] [,2] [,3]  
## [1,]     0    1   -2  
## [2,]     1    0    3  
## [3,]    -2    3    0
```



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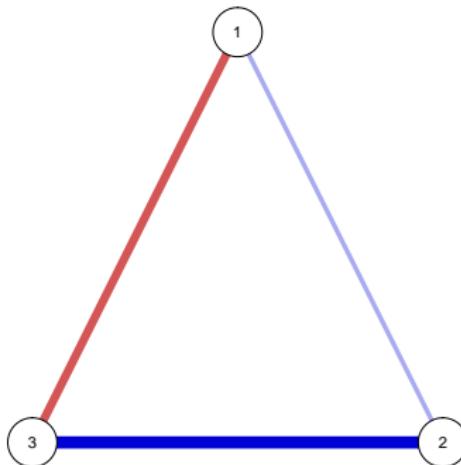
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# Weighted graphs

```
qgraph(input,  
       theme = "colorblind")  
  
input<-matrix(c(  
  0,1,-2,  
  1,0,3,  
 -2,3,0),3,3,byrow=TRUE)  
print(input)  
  
##      [,1] [,2] [,3]  
## [1,]     0    1   -2  
## [2,]     1    0    3  
## [3,]    -2    3    0
```



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## Interpreting qgraph

- Under the default coloring scheme, positive edge weights (here correlations) are shown as green (colorblind theme: blue) edges and negative edge weights as red.
- An edge weight of 0 is omitted. The wider and more colorful an edge the stronger the edge weight.

To interpret **qgraph** networks, three values need to be known:

**Minimum** Edges with absolute weights under this value are omitted

**Cut** If specified, splits scaling of width and color

**Maximum** If set, edge width and color scale such that an edge with this value would be the widest and most colorful

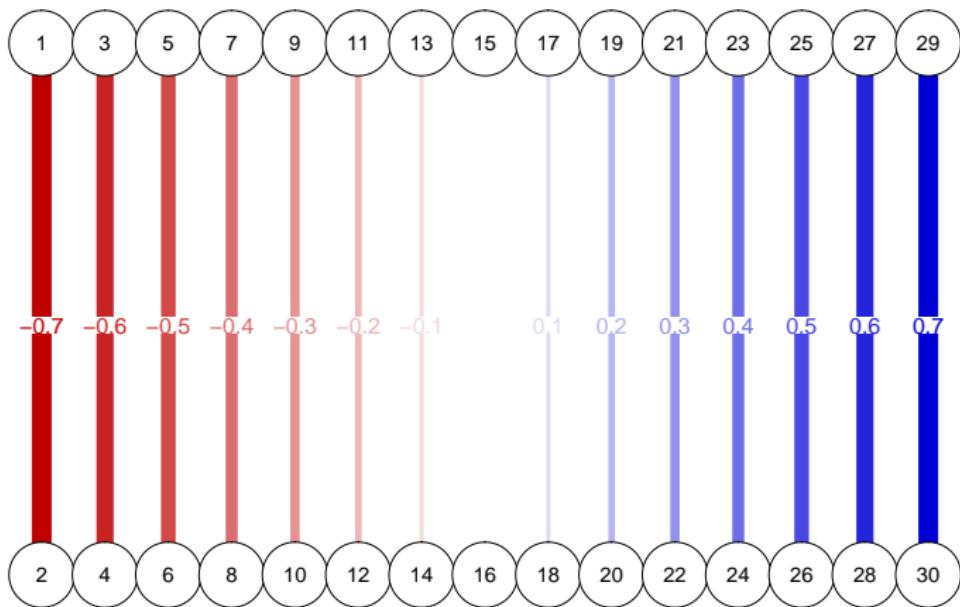
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No minimum, maximum or cut

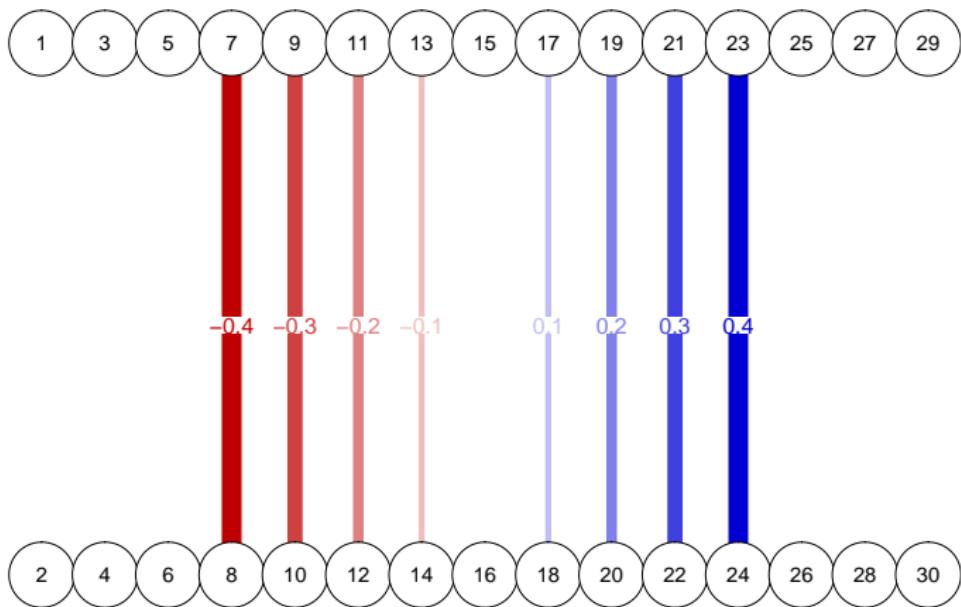
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No minimum, maximum or cut

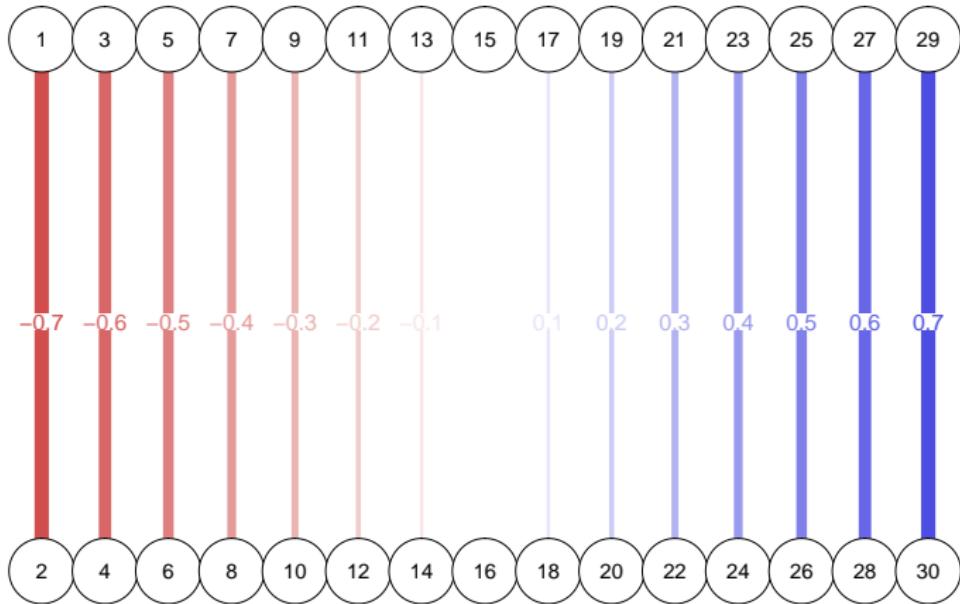
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## Maximum 1

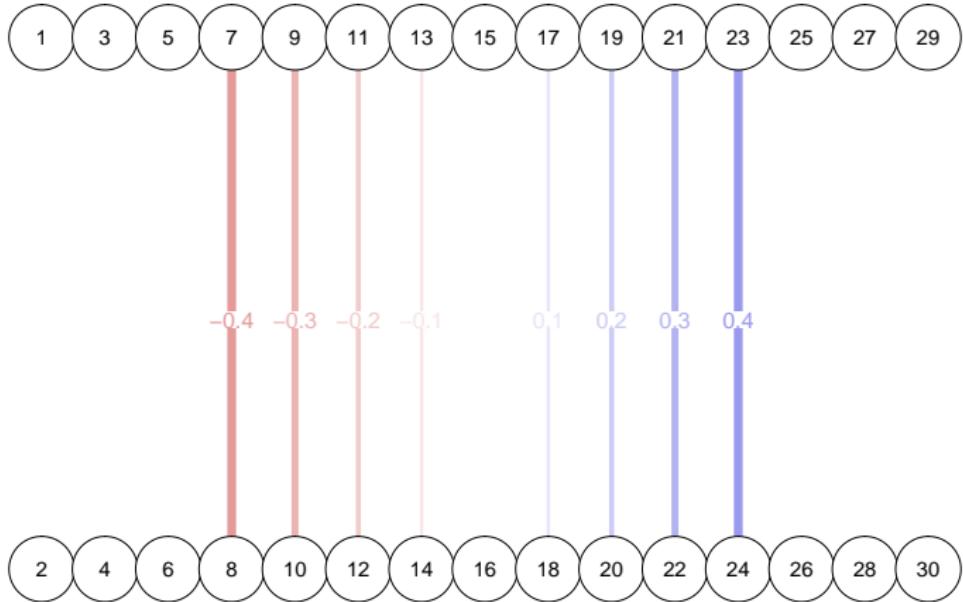
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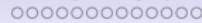


Maximum 1

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Maximum must be set to make graphs comparable!

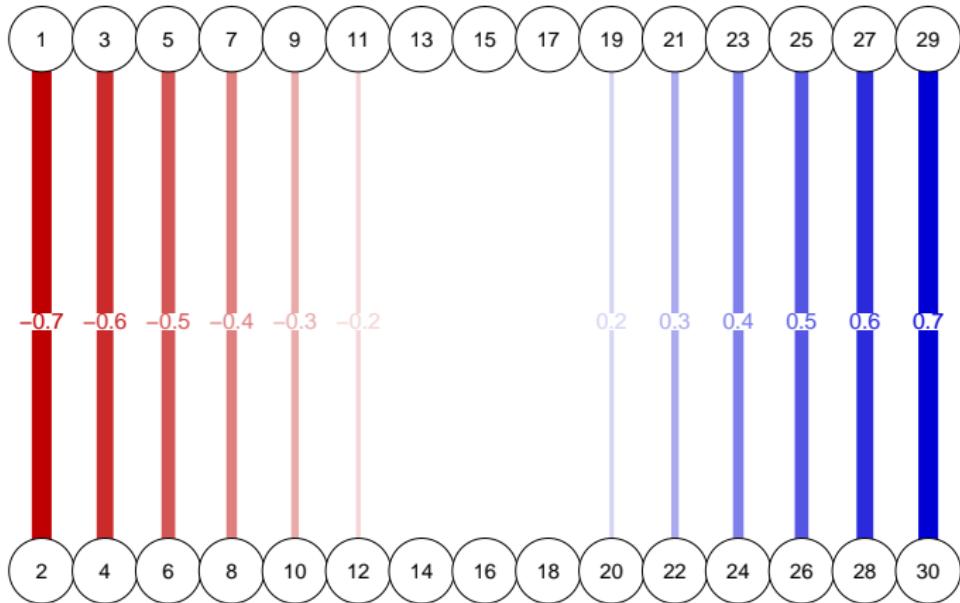
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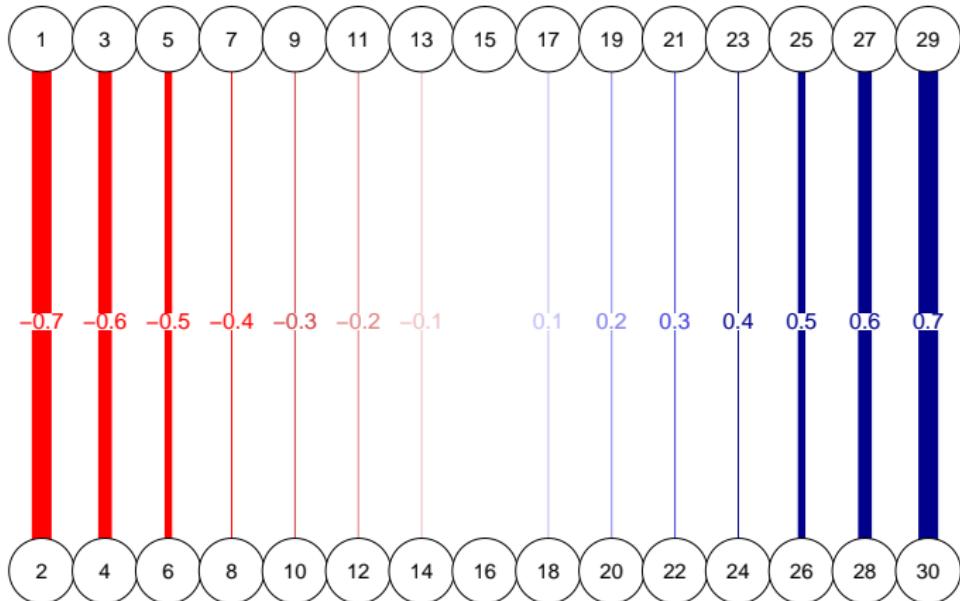
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Cut 0.4

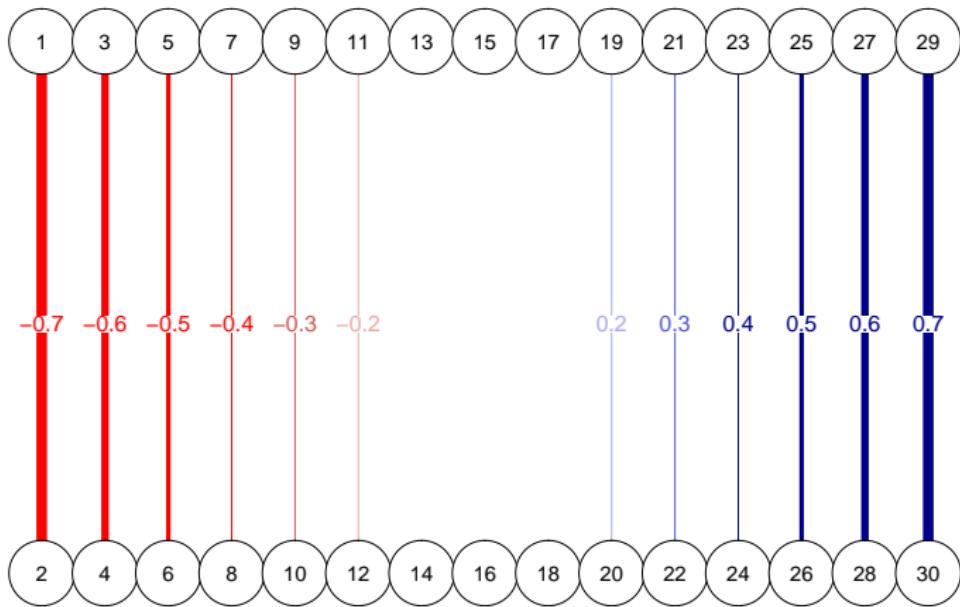
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Minimum 0.1

Cut 0.4

Maximum 1

# Layout modes

- The placement of the nodes is specified with the `layout` argument in `qgraph()`
- This can be a  $n$  by 2 matrix indicating the  $x$  and  $y$  position of each node
- `layout` can also be given a character indicating one of the two default layouts
  - If `layout="circular"` the nodes are placed in circles per group (if the `groups` list is specified)
  - If `layout="spring"` the Fruchterman Reingold algorithm is used for the placement

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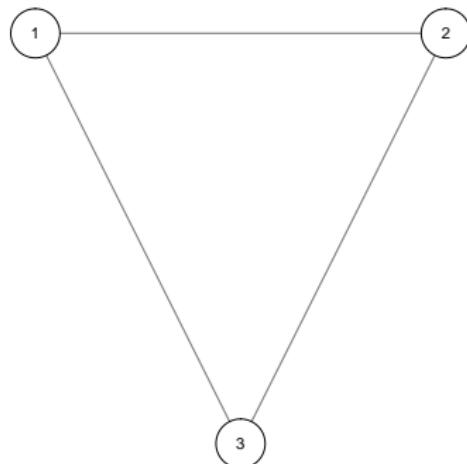
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# Layout matrix

```
qgraph(input, layout=L)
input <- matrix(1, 3, 3)
L <- matrix(c(
  0, 1,
  1, 1,
  0.5, 0),
  ncol=2, byrow=TRUE)
print(L)

##          [,1]  [,2]
## [1,]    0.0    1
## [2,]    1.0    1
## [3,]    0.5    0
```



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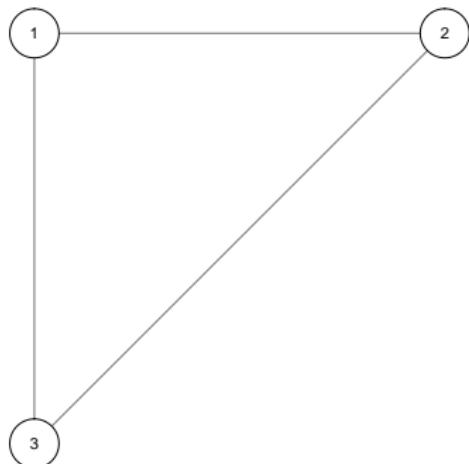
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# Layout matrix

**qgraph**(input, layout=L)

```
L <- matrix(c(  
  0, 1,  
  1, 1,  
  0, 0), ncol=2, byrow=TRUE)  
print(L)  
  
##          [,1] [,2]  
## [1,]      0     1  
## [2,]      1     1  
## [3,]      0     0
```



## Layout matrix

- With the layout matrix the layout of the network can be specified
- The scale is not relevant
- `qgraph()` returns a list containing everything needed to make the graph
- This can be used to force another graph based on the layout of the first

```
Q <- qgraph(input)
qgraph(input2, layout=Q$layout)
```

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## output

**qgraph** graphs **cannot** be manually rescaled, and hence the **RStudio** Export function **cannot** be used to save **qgraph** graphs.

For the best result, save graphs in a PDF device!

Note that if a legend is used, the plot is made square by making the width 1.4 times the height of a plot

# Export to PDF

```
# Open a pdf device:  
pdf("nameoffile.pdf")  
  
# Plot stuff:  
qgraph(1)  
  
# Close pdf device:  
dev.off()  
  
## pdf  
## 2
```

(If you get faulty output, make sure to run `dev.off()` enough times until R returns Null Device)

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# Export to PNG

```
# Open a pdf device:  
png("nameoffile.png")
```

```
# Plot stuff:  
qgraph(1)
```

```
# Close pdf device:  
dev.off()
```

```
## pdf  
## 2
```

(If you get faulty output, make sure to run `dev.off()` enough times until R returns Null Device)

## Important qgraph arguments

`minimum` Omits edge weights with absolute values under this argument

`maximum` Sets the strongest edge to scale to

`cut` Splits scaling of color and width

`vsize` Sets the size of nodes

`esize` Sets the size of edges

`asize` Sets the size of arrows

`filetype` Type of file to save the plot to

`filename` Name of the file to save the plot to

`groups` Colors nodes and adds a legend

`nodeNames` Adds a legend with specific names per node

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# qgraph

The developmental version of **qgraph** can be found on GitHub (<https://github.com/SachaEpskamp/qgraph>) and can be installed using **devtools**

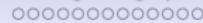
```
library("devtools")
install_github("qgraph", "sachaepskamp")
```

If you have any ideas on concepts to implement in **qgraph** or encounter any bugs please post them on GitHub!

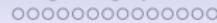
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# Practical time!