

RGraph2js: Usage from an R session

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1 Introduction

RGraph2js provides a powerful HTML visualizer to navigate and manipulate graphs/networks. This package has been designed to display results from in-house algorithms on biological networks [1], where it is required to associate a plot for each node [2]. The package is not limited to this specific usage since it is a general tool to visualize various types of networks. *RGraph2js* is highly customizable and offers a user-friendly interface.

Included features are:

- Interactive visualization tool (pan, zoom)
- Customizable appearance
- Customizable graph layout
- Different node connection types support
- Tooltips support
- Node dragging
- Export as a Scalable Vector Graphics (SVG¹) image
- Barplots and starplots displayable inside the nodes
- Compatibility with most platforms and browsers
- The generated interactive graph can be easily shared

¹https://en.wikipedia.org/wiki/Scalable_Vector_Graphics

RGraph2js takes the description of a graph/network as input and generates an HTML page the user can open in any recent web browser with SVG (Scalable Vector Graphics) rendering support to visualize it and interact with it.

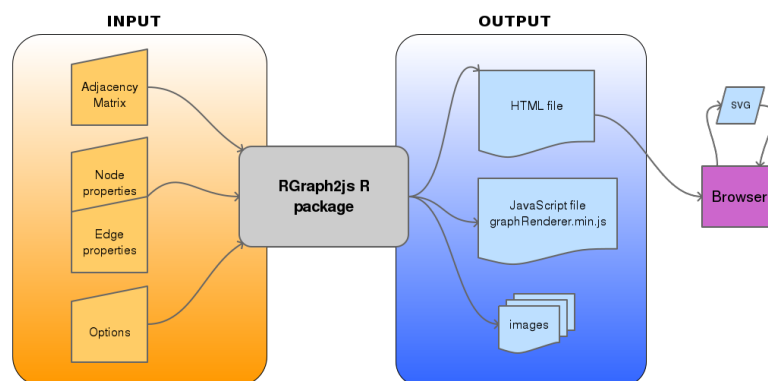


Figure 1: Overview

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1.1 Technology

The *D3js*[3] (Data-Driven Documents) JavaScript library is used to render graphs/networks. *Raphael*[5] is another JavaScript library used to render specific in-nodes plots like starplots. *JQuery*[4] and *JQueryUI*[6] are used for the graphical interface and the user interactions. *qTip2*[7], a *JQuery* plugin, is used to render advanced tooltips. A SVG (Scalable Vector Graphics) capable browser is required since both *D3js* and *Raphael* generate SVG code.

Comment: An Internet connection is required in order to use external third-party JavaScript libraries, further information is given in the next section

1.2 External third-party libraries

D3js, *JQuery*, *JQueryUI*, *qTip2* and *Raphael* are used via *CDNJS*, the links are:

<http://cdnjs.cloudflare.com/ajax/libs/jquery/1.11.0/jquery.min.js>

<http://cdnjs.cloudflare.com/ajax/libs/jqueryui/1.10.3/jquery-ui.min.js>

<http://cdnjs.cloudflare.com/ajax/libs/qttip2/2.2.0/basic/jquery.qtip.min.js>

<http://cdnjs.cloudflare.com/ajax/libs/qttip2/2.2.0/basic/imagesloaded.pkg.min.js>

<http://cdnjs.cloudflare.com/ajax/libs/d3/3.5.6/d3.min.js>

<http://cdnjs.cloudflare.com/ajax/libs/raphael/2.1.4/raphael-min.js>

<http://cdnjs.cloudflare.com/ajax/libs/jqueryui/1.10.3/css/base/minified/jquery-ui.min.css>

<http://cdnjs.cloudflare.com/ajax/libs/qttip2/2.2.0/basic/jquery.qtip.min.css>

The above URLs are declared in the function `RGraph2js:::getDefaultToolParameters()`.

1.3 Input

The graph/network is defined with a signed and weighted adjacency matrix or with the following R objects from the *graph* package:

- `graphAM`
- `graphBAM`
- `graphNEL`
- `clusterGraph`

Considering the matrix `a35`:

```
> v <- c(0, 4, 1,
+       1, 0, 0,
+       -1, 0, 0,
+       0, -2, 0,
+       0, 1, 0)
> a35 <- matrix(v, 3, 5)
> colnames(a35) <- LETTERS[1:5]
> rownames(a35) <- LETTERS[1:3]
```

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... its graphical representation would be as follows:

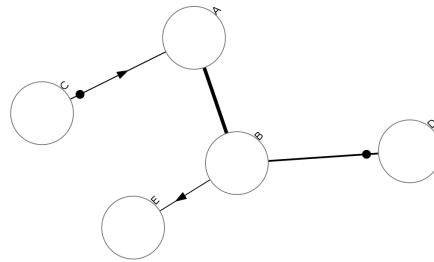


Figure 2: [Graph representation](#)

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	A	B	C	D	E
A	0	1	-1	0	0
B	4	0	0	-2	1
C	1	0	0	0	0

Table 1: 3x5 Signed Weighted Adjacency Matrix

Reading the adjacency matrix by rows, gives the following links/edges:

Line 1: [A → B], [A -• C]

Line 2: [B → A], [B -• D], [B → E]

Line 3: [C → A]

In the adjacency matrix, a value of

- 0 means "no connection"
- 1 "→" "arrow, directional connection"
- -1 "-|" "-•" "dot, directional connection"

Comment: Any bidirectional connection of the same type implies an undirected link marked as "-"

[A → B] and [B → A] ⇒ [A - B]

Comment: Any loop connection, when a node connects with itself, will not be graphically represented

Comment: Edges weights can be directly specified in the adjacency matrix as real numbers

1.4 Output

The result files will be made available in a temporary folder or in a specified folder of your choice. The folder will contain:

- A folder for the images
- The main HTML file
- A JavaScript library

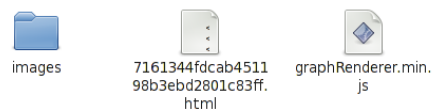


Figure 3: Output folder content

2 Examples

2.1 Simple Example

This example will show the basics, we will generate a simple network given an adjacency matrix.

Define the adjacency matrix `a1515`:

```
> library(RGraph2js)
> v <- c(1,1,1,0,0,0,0,0,0,0,0,0,0,0,0,
+       1,1,0,1,1,1,0,0,0,0,0,0,0,0,0,
+       1,0,0,0,0,0,0,0,0,0,0,0,0,0,0,
+       0,1,0,1,1,0,0,0,0,0,1,0,0,0,0,
+       0,1,0,1,1,0,0,0,1,0,0,0,0,0,0,
+       0,1,0,0,0,0,1,1,0,0,0,0,0,0,0,
+       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,
+       0,0,0,0,1,0,0,0,1,1,0,0,0,0,0,
+       0,0,0,0,0,0,0,0,1,0,0,0,0,0,0,
+       0,0,0,1,0,0,0,0,0,0,0,1,1,1,0,
+       0,0,0,0,0,0,0,0,0,0,1,0,0,0,0,
+       0,0,0,0,0,0,0,0,0,0,1,0,0,0,0,
+       0,0,0,0,0,0,0,0,0,0,1,0,0,0,1,
+       0,0,0,0,0,0,0,0,0,0,0,0,0,1,1
+     )
> a1515 <- matrix(v, 15, 15)
> colnames(a1515) <- LETTERS[1:15]
> rownames(a1515) <- LETTERS[1:15]
```

	A	B	C	D	E	F	G	H	I	J	K	L	M	N	O
A	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0
B	1	1	0	1	1	1	0	0	0	0	0	0	0	0	0
C	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
D	0	1	0	1	1	0	0	0	0	0	1	0	0	0	0
E	0	1	0	1	1	0	0	0	1	0	0	0	0	0	0
F	0	1	0	0	0	0	1	1	0	0	0	0	0	0	0
G	0	0	0	0	0	1	1	0	0	0	0	0	0	0	0
H	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0
I	0	0	0	0	1	0	0	0	1	1	0	0	0	0	0
J	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0
K	0	0	0	1	0	0	0	0	0	0	0	1	1	1	0
L	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0
M	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0
N	0	0	0	0	0	0	0	0	0	0	1	0	0	0	1
O	0	0	0	0	0	0	0	0	0	0	0	0	0	1	1

Table 2: 15x15 Adjacency matrix

Define the output destination folder `outputDir` and generate the graph with the function `graph2js()`

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```
> outputDir <- file.path(tempdir(), "RGraph2js_simpleExample")  
> g <- graph2js(a1515, outputDir=outputDir)
```

Open the `outputDir` in your browser and click on the html file. You should be able to see something similar to this:

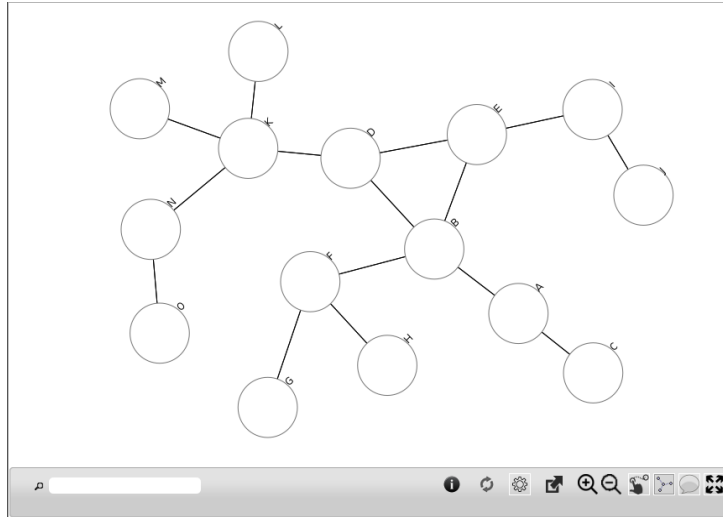


Figure 4: [Simple example](#)

RGraph2js: Usage from an R session

2.2 Visual appearance

In the previous example, we only specified the adjacency matrix. This example will show how to customize the visual appearance of both nodes and links.

The properties of edges (links) can be specified globally or for each edge. `edgesGlobal` below is applied to all edges, where `edgesProp` is only applied to the edges D to E, D to B and B to E.

```
> edgesGlobal <- list(width=2, color="#0000ff")
> edgesProp <- data.frame(from=c("D", "D", "B"),
+                          to=c("E", "B", "E"),
+                          width=c(5, 5, 5))
```

We can also define `edgesProp` by starting with its default value returned by the function `getEdgesDataFrame()`. The first column is an automatically generated unique identifier, followed by the columns `from`, `to` and `type` which are automatically set according to the given adjacency matrix `a1515`. `color`, `width`, `link` and `tooltip` have default values we can customize.

- `color` : edge color formatted as hex RGB
- `width` : edge tickness
- `link` : URL associated with the edge
- `tooltip` : tooltip content with HTML support

```
> getEdgesDataFrame(a1515)
      from to type width  color link tooltip
44849e10a384ca97bce1cfd10bfbba49  A  A  --    1 #000000
7729a246cea8bb6f94f6babde4580443  A  B  --    1 #000000
3582f418dccb4116d85bf4ebbc8a43dc  A  C  --    1 #000000
5f3376f6c060c458f0bc83dde6011158  B  A  --    1 #000000
39ad8dba0347ec71d5010c01b91161b9  B  B  --    1 #000000
8435b69634225e372b5389b2a146aef8  B  D  --    1 #000000
399b9dcabd8470e1989d539694cd2aab  B  E  --    1 #000000
bad5ddd9aacc6cd33e51ad4cde7e3991  B  F  --    1 #000000
d2b15200cf583e212151f01d8e740438  C  A  --    1 #000000
1a7cf664997bae2b944e1fb8ece4abc8  D  B  --    1 #000000
62435036701687344dd7825f2394fa49  D  D  --    1 #000000
05f671eada96e9ccebdcf011c913c613  D  E  --    1 #000000
f3bf8a0333bff715b99e18115adde968  D  K  --    1 #000000
5b049077fe3f9d2b1a06ee88d8b0499a  E  B  --    1 #000000
34a81d610eed20331ea7e096ac1e2de  E  D  --    1 #000000
71f31c02b22bb9212b41f702d1c8f876  E  E  --    1 #000000
5cde14c8755685a7c65147fb3702019e  E  I  --    1 #000000
f43411fdb5c7e9a9430af4f2755ff3dd  F  B  --    1 #000000
dd23b527644ed7642f181cce72cf392f  F  G  --    1 #000000
6a0a2e5e59a224255ad4b434cfbaf1fb  F  H  --    1 #000000
6744176738581d44180b8640d435f534  G  F  --    1 #000000
f77f3018d1792ebe0bcacf81118219379  G  G  --    1 #000000
b1b7ca5d0f5f33195e39c8cbc98f801a  H  F  --    1 #000000
e06c9b4d801061c9d6b215218be309ba  I  E  --    1 #000000
32030f5488046b2337498431b81b323d  I  I  --    1 #000000
```


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```
72879760c86d63ad8dbf5a49c0168fc4 I J -- 1 #000000
85766046803131f1923feceeeea944dc J I -- 1 #000000
a2ab2975600c198d7f163adc009237a7 K D -- 1 #000000
7450b393b94b7dfdf2c3b7dc147a111 K L -- 1 #000000
3af9e129a12b06895c61c607851b5478 K M -- 1 #000000
d73b8adba780a7497023f42ff74d1534 K N -- 1 #000000
dbcedb68807cb9cbf465da7a91aa8d83 L K -- 1 #000000
687ab195fbe14395b5ac83354e28f219 M K -- 1 #000000
9689338efd576a8c53c80f92767d1d6a N K -- 1 #000000
d05e499428d1837a3fa08369b31cc0c3 N O -- 1 #000000
93798bf5ed37d6f329832e7d00d7f287 O N -- 1 #000000
4ecf57c0fcc911300358f7502e906824 O O -- 1 #000000
```

Similarly, node properties can be global or specific.

```
> nodesGlobal <- list(color="#ebebeb")
> nodesProp <- data.frame(shape=c("triangle", "lozenge", "rect"),
+                          color=c("#ff0000", "#0000ff", "#ffff00"))
> rownames(nodesProp) <- c("C", "E", "G")
```

Since `nodesProp` holds node specific properties, row names are mandatory. We can call the `getNodeDataFrame()` to define `nodesProp`. The returned data frame contains default values for each node.

- `color` : color of the node in hex RGB format
- `shape` : the shape to use ("rect", "circle", "lozenge", "triangle")
- `link` : URL associated with the node
- `tooltip` : tooltip content with HTML support

```
> getNodeDataFrame(A=a1515, nGlobal=nodesGlobal, nProp=nodesProp)
```

	width	color	shape	link	tooltip
A	1	#ebebeb	circle		
B	1	#ebebeb	circle		
C	1	#ff0000	triangle		
D	1	#ebebeb	circle		
E	1	#0000ff	lozenge		
F	1	#ebebeb	circle		
G	1	#ffff00	rect		
H	1	#ebebeb	circle		
I	1	#ebebeb	circle		
J	1	#ebebeb	circle		
K	1	#ebebeb	circle		
L	1	#ebebeb	circle		
M	1	#ebebeb	circle		
N	1	#ebebeb	circle		
O	1	#ebebeb	circle		

Call the `graph2js()` function as before and specify both nodes and edges properties.

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```
> outputDir <- file.path(tempdir(), "RGraph2js_visualAppearance")
> g <- graph2js(a1515,
+             nodesGlobal=nodesGlobal, edgesGlobal=edgesGlobal,
+             nodesProp=nodesProp, edgesProp=edgesProp,
+             outputDir=outputDir, file="index.html")
```

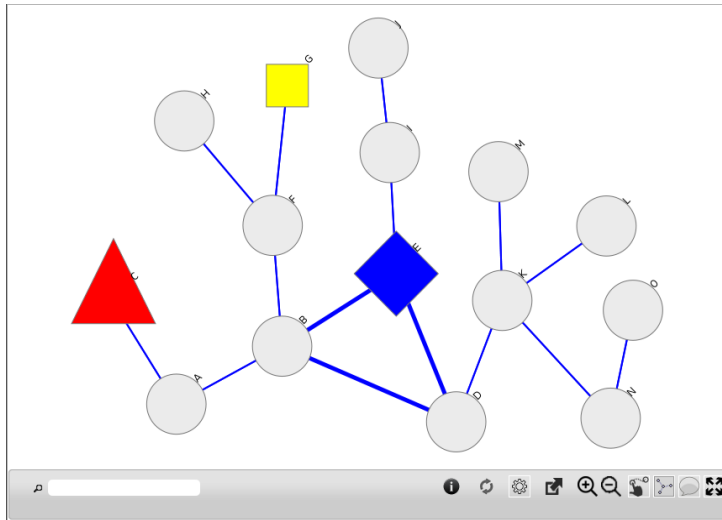


Figure 5: Appearance customized

Going further, several options can be changed via the `opts` parameter of the function `graph2js()`. `opts` defaults to the value returned by the function `getDefaultOptions()`.

Please check out the manual for further details.

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2.3 Fixed node positons

We start from a simple adjacency matrix:

```
> v <- c(0, 0, 1,
+       1, 0, 0,
+       0, 0, 0,
+       0, -1, 0,
+       0, 1, 0)
> a35 <- matrix(v, 3, 5)
> colnames(a35) <- LETTERS[1:5]
> rownames(a35) <- LETTERS[1:3]
```

Then, we specify node coordinates via the node properties. `x` and `y` represent the Cartesian coordinates, and `fixed` means they are immutable.

```
> r <- 100
> sector <- 2*pi/5
> n.prop <- data.frame(
+   x=c(r*cos(1*sector), r*cos(2*sector), r*cos(3*sector),
+       r*cos(4*sector), r*cos(5*sector)),
+   y=c(r*sin(1*sector), r*sin(2*sector), r*sin(3*sector),
+       r*sin(4*sector), r*sin(5*sector)),
+   fixed=c(TRUE, TRUE, TRUE, TRUE, TRUE)
+ )
> rownames(n.prop) <- c("A", "B", "C", "D", "E")
```

Now, we render the graphics.

```
> outputDir <- file.path(tempdir(), "RGraph2js_fixedNodes")
> g <- graph2js(a35, nodesProp=n.prop, outputDir=outputDir)
```

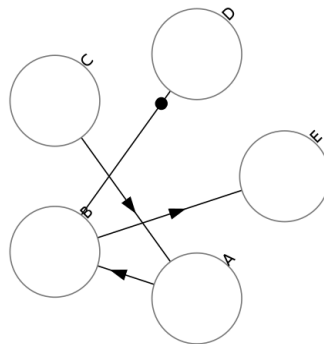


Figure 6: Fixed Node Coordinates Graph

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2.4 Time data or categories

RGraph2js implements a time-step functionality where, at each step, a different set of nodes can be highlighted with thicker borders and different colors. Each time-step is specified by an index starting at 1 and the highlighted nodes and their colors are specified by a dataframe as explained below.

Take for instance the following graph definition:

```
> v <- c(0, 0, 1,
+       1, 0, 0,
+       0, 0, 0,
+       0, -1, 0,
+       0, 1, 0)
> a35 <- matrix(v, 3, 5)
> colnames(a35) <- LETTERS[1:5]
> rownames(a35) <- LETTERS[1:3]
```

We specify 4 time-steps in the dataframe below using 2 prefixes:

- `leading.nodes.index` specifies the nodes to highlight with thicker border
- `highlight.index` specifies the colors for those leading nodes.

```
> numnodes <- 5
> nodesProp <- data.frame(leading.nodes.1=rbinom(numnodes, 1, 1/2),
+                         leading.nodes.2=rbinom(numnodes, 1, 1/2),
+                         leading.nodes.3=rbinom(numnodes, 1, 1/2),
+                         leading.nodes.4=rbinom(numnodes, 1, 1/2),
+                         highlight.1=rainbow(numnodes),
+                         highlight.2=rainbow(numnodes),
+                         highlight.3=rainbow(numnodes),
+                         highlight.4=rainbow(numnodes))
> rownames(nodesProp) <- LETTERS[1:5]
```

Renderering the Graph leads to:

```
> outputDir <- file.path(tempdir(), "RGraph2js_timeData")
> g <- graph2js(a35,
+              nodesProp=nodesProp,
+              outputDir=outputDir)
```

Clicking on the **LN** button will expand a new panel at the bottom containing a slider to navigate across the time steps.

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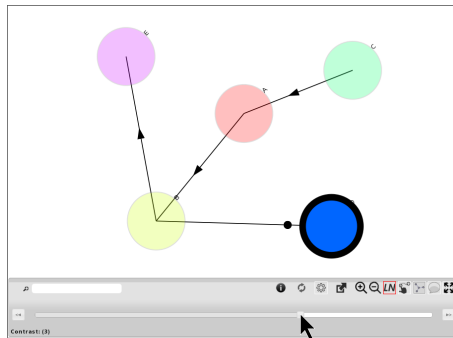


Figure 7: 4 Different states

2.5 Rendering barplots inside nodes

```
> v <- c(0, 0, 1,
+       1, 0, 0,
+       0, 0, 0,
+       0, -1, 0,
+       0, 1, 0)
> a35 <- matrix(v, 3, 5)
> colnames(a35) <- LETTERS[1:5]
> rownames(a35) <- LETTERS[1:3]
```

The `innerValues` parameter allows us to specify a barplot for each node and `innerColors` represent the bar colors. The order in both parameters is important for the barplots rendering.

```
> numnodes <- 5
> innerValues <- matrix(runif(numnodes * 8), numnodes, 8)
> rownames(innerValues) <- LETTERS[1:5]
> innerColors <- matrix(rainbow(numnodes * 8), numnodes, 8)
> rownames(innerColors) <- LETTERS[1:5]
```

```
> outputDir <- file.path(tempdir(), "RGraph2js_barplots")
> g <- graph2js(a35,
+             innerValues=innerValues,
+             innerColors=innerColors,
+             outputDir=outputDir)
```

After rendering, here is the result:

An alternate solution would be to display the barplot inside the node tooltips only, as shown below:

```
> opts <- getDefaultOptions()
> opts$displayBarPlotsInsideNodes <- FALSE
> opts$barplotInNodeTooltips <- TRUE
> g <- graph2js(a35,
+             opts=opts,
+             innerValues=innerValues,
+             innerColors=innerColors,
```

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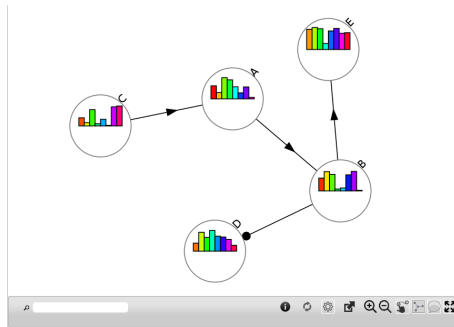


Figure 8: Barplots inside nodes

+ `outputDir=outputDir)`

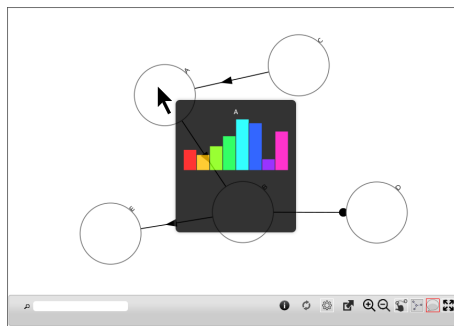


Figure 9: Barplots in tooltips only

2.6 Rendering starplots inside nodes

```
> v <- c(0, 0, 1,
+       1, 0, 0,
+       0, 0, 0,
+       0, -1, 0,
+       0, 1, 0)
> a35 <- matrix(v, 3, 5)
> colnames(a35) <- LETTERS[1:5]
> rownames(a35) <- LETTERS[1:3]
```

Define a starplot for each node. Each starplot has several parameters: the value (which determines its radius), the opacity, the color, the label, a URL and a tooltip. Additionally, a color and opacity can be specified for the starplot background.

```
> numnodes <- 5
> starplotValues <- matrix(runif(numnodes * 8), numnodes, 8)
> rownames(starplotValues) <- LETTERS[1:5]
> starplotColors <- matrix(rainbow(numnodes * 8), numnodes, 8)
> rownames(starplotColors) <- LETTERS[1:5]
> labels <- c("Sector1", "Sector2", "Sector3", "Sector4",
+           "Sector5", "Sector6", "Sector7", "Sector8")
> starplotLabels <- matrix(labels, numnodes, 8)
> rownames(starplotLabels) <- LETTERS[1:5]
> starplotTooltips <- matrix(labels, numnodes, 8)
> rownames(starplotTooltips) <- LETTERS[1:5]
> # add a url link for each sector
> urls <- c("http://d3js.org/", "http://jquery.com/",
+          "http://jqueryui.com/", "http://qtip2.com/",
+          "http://raphaeljs.com/", "http://www.bioconductor.org/",
+          "http://cran.r-project.org", "http://journal.r-project.org")
> starplotUrlLinks <- matrix(urls, numnodes, 8)
> rownames(starplotUrlLinks) <- LETTERS[1:5]
> starplotCircleFillColor <- matrix(rainbow(numnodes), numnodes, 1)
> rownames(starplotCircleFillColor) <- LETTERS[1:5]
> starplotCircleFillOpacity <- matrix(runif(numnodes,0,1), numnodes, 1)
> rownames(starplotCircleFillOpacity) <- LETTERS[1:5]
```

Render the Graph:

```
> outputDir <- file.path(tempdir(), "RGraph2js_starplots")
> output.filename <- "test.html"
> g <- graph2js(A=a35,
+             starplotColors=starplotColors,
+             starplotLabels=starplotLabels,
+             starplotValues=starplotValues,
+             starplotTooltips=starplotTooltips,
+             starplotUrlLinks=starplotUrlLinks,
+             starplotCircleFillColor=starplotCircleFillColor,
+             starplotCircleFillOpacity=starplotCircleFillOpacity,
+             outputDir=outputDir,
```

RGraph2js: Usage from an R session

```
+ filename=output.filename)
```

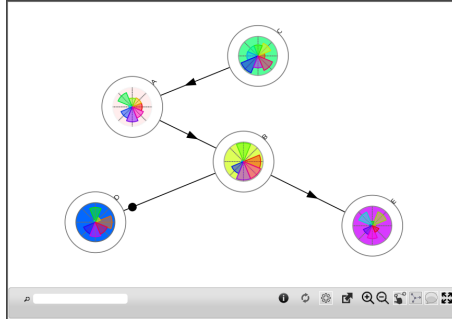


Figure 10: Starplots

Comment: Moving the mouse over the sectors will display a tooltip showing the sector name or label

Comment: Clicking on a sector will open the associated URL

2.7 Customizing the tooltip content

```
> v <- c(0, 0, 1,
+       1, 0, 0,
+       0, 0, 0,
+       0, -1, 0,
+       0, 1, 0)
> a35 <- matrix(v, 3, 5)
> colnames(a35) <- LETTERS[1:5]
> rownames(a35) <- LETTERS[1:3]
```

The content of the tooltip can be defined with the `nodesProperties` parameter which fully supports HTML content. *Comment: We can even add images like any other HTML content*

```
> numnodes <- 5
> someHtmlContent <- c(paste0("<table class=\|gridtable\|>",
+                           "<tr><th>Header 1</th><th>Header 2</th><th>",
+                           "Header 3</th></tr><tr><td>Text 1,1</td><td>",
+                           "Text 1,2</td><td>Text 1,3</td></tr><tr><td>",
+                           "Text 2,1</td><td>Text 2,2</td><td>Text 2,3",
+                           "</td></tr></table>"),
+                       "This is another <i>content</i>",
+                       "Yet another <font style=\|color:#00ff00;\|>one</font>",
+                       paste0("<table>",
+                               "<tr><th>Header 1</th><th>Header 2</th><th>",
+                               "Header 3</th></tr><tr><td>Text 1,1</td><td>",
+                               "Text 1,2</td><td>Text 1,3</td></tr><tr><td>",
+                               "Text 2,1</td><td>Text 2,2</td><td>Text 2,3",
+                               "</td></tr></table>"),
+                       "<h1>Header 1</h1><h2>Header 2</h2>")
> n.prop <- data.frame(tooltip=someHtmlContent)
> rownames(n.prop) <- LETTERS[1:5]
```

Since we specified a custom style `gridtable`, we can define it that way:

```
> userCssStyles <- "
+ <style type=\|text/css\|>
+ table.gridtable {
+   font-family: verdana,arial,sans-serif;
+   font-size:11px;
+   color:#333333;
+   border-width: 1px;
+   border-color: #666666;
+   border-collapse: collapse;
+ }
+ table.gridtable th {
+   border-width: 1px;
+   padding: 8px;
+   border-style: solid;
+   border-color: #666666;
+   background-color: #dedede;
```

RGraph2js: Usage from an R session

```
+ }
+ table.gridtable td {
+   border-width: 1px;
+   padding: 8px;
+   border-style: solid;
+   border-color: #666666;
+   background-color: #ffffff;
+ }
+ </style>
+ "
```

Render the Graph and provide custom styles with the `userCssStyles` parameter:

```
> outputDir <- file.path(tempdir(), "RGraph2js_tooltipContent")
> g <- graph2js(a35,
+   opts=opts,
+   nodesProp=n.prop,
+   userCssStyles=userCssStyles,
+   outputDir=outputDir)
```

The 5 tooltips will be rendered as follows:

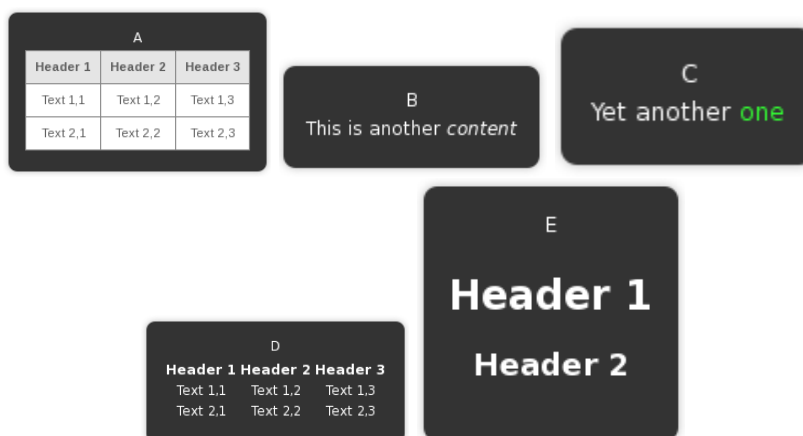


Figure 11: Custom Tooltips

RGraph2js: Usage from an R session

2.8 Use the DOT description language

This example requires the `sna`^[8] package which allows us to easily get an adjacency matrix from a DOT ²[9] file.

²<http://www.graphviz.org/doc/info/lang.html>

```
> library(sna)
> extdata.path <- file.path(path.package(package="RGraph2js"), "extdata")
> dot.file.path <- file.path(extdata.path, "nohosts.dot")
> adj.mat <- read.dot(dot.file.path)
```

Since the graph is rather large, we can save computing resources by displaying the graph every 100 iterations only, with the option `displayNetworkEveryNLayoutIterations`. Setting it at "zero" would mean to display the graph upon completion only.

```
> opts <- getDefaultOptions()
> opts$displayNetworkEveryNLayoutIterations <- 100
> opts$displayNodeLabels <- FALSE
> opts$layout_forceCharge <- -2400
> nodesGlobal <- list(color="#5544ff")
> outputDir <- file.path(tempdir(), "RGraph2js_dot")
> g <- graph2js(A=adj.mat,
+             nodesGlobal=nodesGlobal,
+             opts=opts,
+             outputDir=outputDir)
```

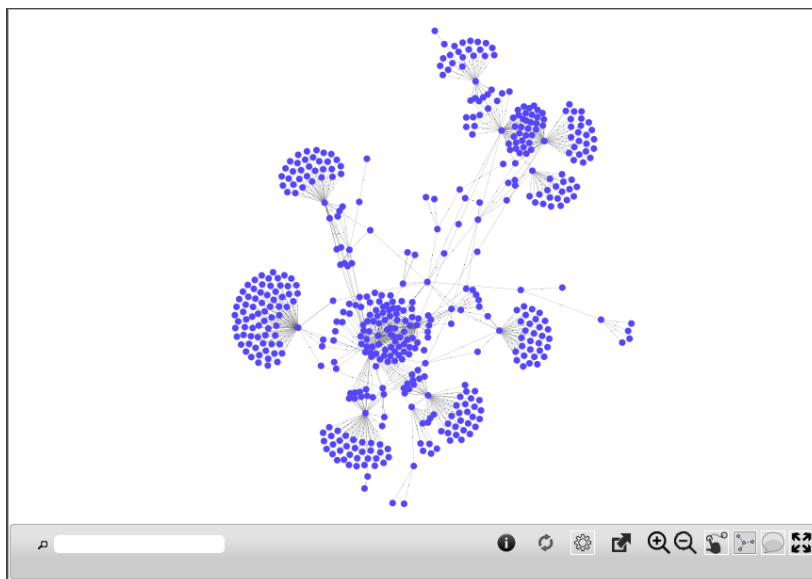


Figure 12: Generate a network from a DOT file

RGraph2js: Usage from an R session

2.9 Use a graph class

Instead of specifying an adjacency matrix, you can pass a *graph* class.

Here is an example with a `graphNEL` object `gnel`:

```
> library(graph)
> nodes <- c("A", "B", "C", "D", "E")
> edges <- list(
+   A=list(edges=c("A", "B"), weights=c(2, 2)),
+   B=list(edges=c("A", "E"), weights=c(0.25, 0.25)),
+   C=list(edges=c("A", "D"), weights=c(4, 4)),
+   D=list(edges=c("E"), weights=c(6)),
+   E=list(edges=c("A", "B"), weights=c(1, 1))
+ )
> gnel <- new("graphNEL", nodes=nodes, edgeL=edges, edgemode="directed")
```

The following shows how to graphically represent edges weights with the *Rgraphviz* package. As you can see, some extra steps are required.

```
> ew <- as.character(unlist(edgeWeights(gnel)))
> ew <- ew[setdiff(seq(along = ew), removedEdges(gnel))]
> names(ew) <- edgeNames(gnel)
> eAttrs <- list()
> eAttrs$label <- ew
> plot(gnel,
+   attrs=list(
+     edge=list(arrowsize=0.5)
+   ),
+   edgeAttrs=eAttrs)
```

Now, with *RGraph2js*, edges weights are translated into edges width by default. This default behaviour can be redefined by specifying edges properties.

```
> outputDir <- file.path(tempdir(), "RGraph2js_graphNELExample")
> g <- graph2js(A=gnel, outputDir=outputDir)
```

Please note the following limitations:

- links/edges representations are only translatable into "→" or "—"
- as mentioned earlier, loop connections are not rendered

RGraph2js: Usage from an R session

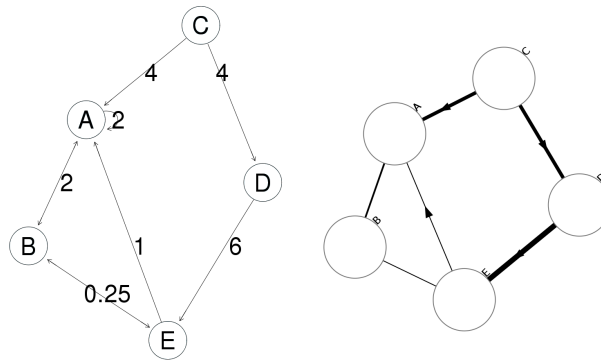


Figure 13: Comparison of the original `graphNEL` (left) and the `RGraph2js` output (right)

3 Interactions

3.1 Using the bottom panel buttons

All buttons are described in the next sections.



Figure 14: Buttons bottom panel

3.1.1 Search

The search field performs an incremental search on all node labels, highlighting matches with a tick red border. Clearing the search field cancels the search and resets the display.



Figure 15: Search field

3.1.2 About dialog

- Gives information about the software and its version

3.1.3 Reload

- Re-compute the layout

RGraph2js: Usage from an R session

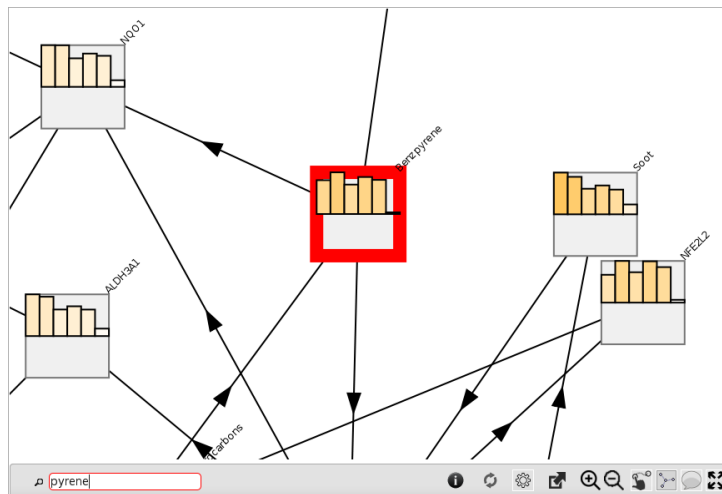


Figure 16: Search feature in action

3.1.4 Layout settings

⚙️ Toggle the sub-panel to customize the layout engine



Figure 17: Layout Settings

The parameters the user can control with sliders are:

- Charge
- Link distance

More details about the force layout can be found on the D3js wiki ³.

³<https://github.com/mbostock/d3/wiki/Force-Layout>

3.1.5 Export

📄 Lets you export the graph and save it as an SVG image

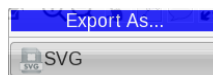


Figure 18: "Export As" popup menu

3.1.6 Zoom

🔍 Zoom in/out without using the mouse wheel

RGraph2js: Usage from an R session

3.1.7 Leading nodes


LN Expand a new panel at the bottom containing a slider to navigate across the time steps. Please note this button is present only when such data exist.



Figure 19: [Layout Settings](#)

RGraph2js: Usage from an R session

3.1.8 Dragging nodes

 Toggle the nodes dragging feature

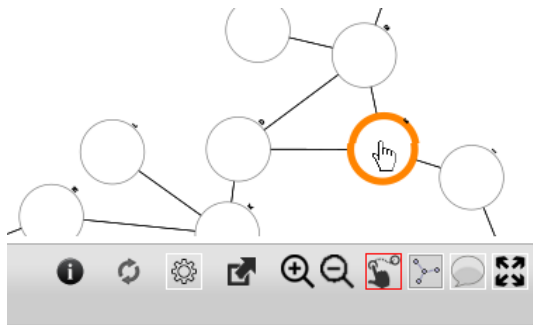



Figure 20: Dragging a node

3.1.9 Node neighbors

 Enable the highlight of the neighbors when hovering a node

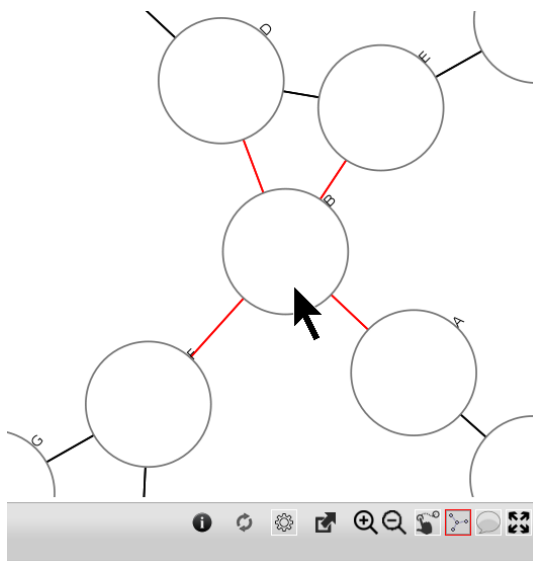


Figure 21: Highlighting of the neighbors

RGraph2js: Usage from an R session

3.1.10 Tooltips

🗨 Toggle the display of Tooltips when the mouse hovers a node or an edge

Below is an example of a node tooltip containing the node name with a barplot

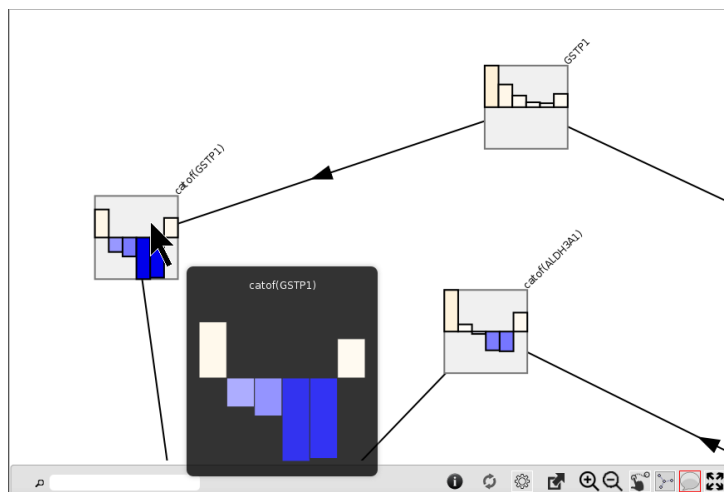


Figure 22: Tooltips

3.1.11 Magnify

🔍 Magnify the view area to fit to the browser current window size

3.2 Using the Mouse

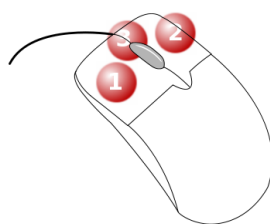


Figure 23: Mouse Buttons

Button (1) is used to drag the whole graph in the drawing area and to drag nodes when the corresponding mode 🖱 is activated. Double-clicking performs a zoom in.

Button (2) opens a popup menu when clicking a node.

Button (3), the mouse wheel allows to zoom in and out.

RGraph2js: Usage from an R session

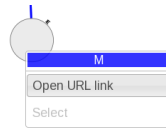


Figure 24: Node Popup Menu

References

- [1] Florian Martin, Alain Sewer, Marja Talikka, Yang Xiang, Julia Hoeng and Manuel C Peitsch, *Quantification of biological network perturbations for mechanistic insight and diagnostics using two-layer causal models*, BMC Bioinformatics 2014, 15:238. URL <http://www.biomedcentral.com/1471-2105/15/238>
- [2] Poussin Carine, Laurent Alexandra, Peitsch Manuel C., Hoeng Julia and De Leon Hector *Systems Biology Reveals Cigarette Smoke-Induced Concentration-Dependent Direct and Indirect Mechanisms That Promote Monocyte-Endothelial Cell Adhesion.*, Toxicological Sciences Journal, 2015.
Abstract:
<http://toxsci.oxfordjournals.org/content/early/2015/08/20/toxsci.kfv137.abstract>
Eprint:
<http://toxsci.oxfordjournals.org/content/early/2015/08/20/toxsci.kfv137.full.pdf+html>
- [3] Michael Bostock, Vadim Ogievetsky and Jeffrey Heer, *D3: Data-Driven Documents*, IEEE Trans. Visualization & Comp. Graphics (Proc. InfoVis), 2011. URL <http://vis.stanford.edu/papers/d3>
- [4] *jQuery*, URL <http://jquery.org/>
- [5] *Raphael*, URL <http://dmitrybaranovskiy.github.io/raphael/>
- [6] *jQueryUI*, URL <http://jqueryui.com/>
- [7] *qTip2*, URL <http://qtip2.com/>
- [8] Mark S. Handcock, David R. Hunter, Carter T. Butts, Steven M. Goodreau, and Martina Morris, *statnet: Software tools for the Statistical Modeling of Network Data*. URL <http://statnetproject.org>
- [9] Emden R. Gansner and Stephen C. North, *An open graph visualization system and its applications to software engineering.*, SOFTWARE - PRACTICE AND EXPERIENCE Journal, Volume 30 Number 11 pages 1203-1233, 2000. URL <http://www.graphviz.org>
- [10] R. Gentleman and Elizabeth Whalen and W. Huber and S. Falcon, *graph: A package to handle graph data structures*, URL <http://www.bioconductor.org/packages/2.10/bioc/html/graph/>