R Package diagram: visualising simple graphs, flowcharts, and webs

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Abstract

This document describes how to use the diagram package (Soetaert 2009a) for plotting small networks, flow charts, and (food) webs. Together with R-package shape (Soetaert 2009b) this package has been written to produce the figures of the book (Soetaert and Herman 2009b).
The electrical network symbols were added to produce a figure of the book (Soetaert, Cash, and Mazzia 2012)

Keywords: diagram, food web, flow chart, arrows, R.

1. Introduction

There are three ways in which package diagram can be used:

• function plotmat takes as input a matrix with transition coefficients or interaction strengths. It plots the corresponding network consisting of (labeled) boxes (the components) connected by arrows. Each arrow is labeled with the value of the coefficients.

• function plotweb takes as input a matrix with (flow) values, and plots a web. Here the components are connected by arrows whose thickness is determined by the value of the coefficients.

• Flowcharts can be made by adding separate objects (textboxes) to the figure and connecting these with arrows.

Three datasets have been included:

• Rigaweb, the planktonic food web of the Gulf of Riga (Donali, Olli, Heiskanen, and Andersen 1999).

• Takapotoweb, the Takapoto atoll planktonic food web (Niquil, Jackson, Legendre, and Delesalle 1998).

• Teasel, the transition matrix describing the population dynamics of Teasel, a European perennial weed ((Caswell 2001; Soetaert and Herman 2009b).
The food webs were generated using R packages LIM and limSolve (Soetaert, Van den Meersche, and van Oevelen 2009; Soetaert and van Oevelen 2009) which contain functions to read and solve food web problems respectively.

2. plotmat - plotting networks based on matrix input

This is the quickest method of plotting a network. The network is specified in a matrix, which gives the magnitudes of the links (from columns to rows).

The position of the elements (boxes) is specified by argument pos. Thus, setting pos=c(1,2,1) indicates that the 4 elements will be arranged in three equidistant rows; on the first row one element, on the second row two elements and on the third row one element.

2.1. Simple examples

Below are some simple examples of the use of plotmat. In the first graph - four simple boxes are put; no arrows drawn

The second graph contains round boxes with arrows, labeled "flow"

The third graph has diamond-shaped boxes including self-arrows.

The fourth graph has hexagonal-shaped boxes, with curved arrows. The arrows are enlarged and the arrowhead pointing from box 2 to 4 is colored red.

```r
> par(mar = c(1, 1, 1, 1), mfrow = c(2, 2))
> #
> names <- c("A", "B", "C", "D")
> M <- matrix(nrow = 4, ncol = 4, byrow = TRUE, data = 0)
> plotmat(M, pos = c(1, 2, 1), name = names, lwd = 1,
+ box.lwd = 2, cex.txt = 0.8, box.size = 0.1,
+ box.type = "square", box.prop = 0.5)
> #
> plotmat(M, pos = c(1, 2, 1), curve = 0, name = names, lwd = 1,
+ box.lwd = 2, cex.txt = 0.8, box.type = "circle", box.prop = 1.0)
> #
> #
> diag(M) <- "self"
> plotmat(M, pos = c(2, 2), curve = 0, name = names, lwd = 1, box.lwd = 2,
+ cex.txt = 0.8, self.cex = 0.5, self.shiftx = c(-0.1, 0.1, -0.1, 0.1),
+ box.type = "diamond", box.prop = 0.5)
> M <- matrix(nrow = 4, ncol = 4, data = 0)
> M[2, 1] <- 1 ; M[4, 2] <- 2 ; M[3, 4] <- 3 ; M[1, 3] <- 4
> Col <- M
> Col[,] <- "black"
> Col[4, 2] <- "darkred"
> pp <- plotmat(M, pos = c(1, 2, 1), curve = 0.2, name = names, lwd = 1,
+ box.lwd = 2, cex.txt = 0.8, arr.type = "triangle",
```
The contents of `pp` shows the position of the various items.

```r
> pp

$arr
row col Angle Value rad ArrowX ArrowY TextX TextY
1 2 1 53.1301 1 0.08333333 0.3085298 0.7169284 0.2843333 0.7346667
2 4 2 -53.1301 2 0.08333333 0.3077450 0.2841193 0.2843333 0.2653333
3 1 3 -53.1301 4 0.08333333 0.6918629 0.7164048 0.7156667 0.7346667
4 3 4 53.1301 3 0.08333333 0.6910769 0.2825485 0.7156667 0.2653333

$comp
x y
[1,] 0.50 0.8333333
[2,] 0.25 0.5000000
[3,] 0.75 0.5000000
[4,] 0.50 0.1666667

$radii
x y
[1,] 0.1 0.025
[2,] 0.1 0.025
[3,] 0.1 0.025
[4,] 0.1 0.025

$rect
xleft ybot xright ytop
[1,] 0.40 0.8083333 0.60 0.8583333
[2,] 0.15 0.4750000 0.35 0.5250000
[3,] 0.65 0.4750000 0.85 0.5250000
[4,] 0.40 0.1416667 0.60 0.1916667

2.2. a schematic representation of an ecosystem model

In the example below, first the main components and arrows are drawn (plotmat), and the output of this function written in list pp. This contains, a.o. the positions of the components (boxes), arrows, etc.. It is used to draw an arrow from the middle of the arrow connecting fish and zooplankton ("ZOO") to detritus. Function straightarrow (see below) is used to draw this arrow.
Figure 1: Four simple examples of `plotmat`

```r
> names <- c("PHYTO", "NH3", "ZOO", "DETRITUS", "BotDET", "FISH")
> M <- matrix(nrow = 6, ncol = 6, byrow = TRUE, data = c(
+ # p n z d b f
+ 0, 1, 0, 0, 0, 0, #p
+ 0, 0, 4, 10, 11, 0, #n
+ 2, 0, 0, 0, 0, 0, #z
+ 8, 0, 13, 0, 0, 12, #d
+ 9, 0, 0, 7, 0, 0, #b
+ 0, 0, 5, 0, 0, 0 #f
+ ))
> #
> pp <- plotmat(M, pos = c(1, 2, 1, 2), curve = 0, name = names,
+ lwd = 1, box.lwd = 2, cex.txt = 0.8,
+ box.type = "square", box.prop = 0.5, arr.type = "triangle",
+ arr.pos = 0.4, shadow.size = 0.01, prefix = "f",
+ main = "NPZZDD model")
> #
> phyto <- pp$comp[names=="PHYTO"]
> zoo <- pp$comp[names=="ZOO"]
> nh3 <- pp$comp[names=="NH3"]
> detritus <- pp$comp[names=="DETRITUS"]
> fish <- pp$comp[names=="FISH"]
> #
```
Figure 2: An NPZZDD model

```r
# flow5->detritus
> m2 <- 0.5*(zoo+fish)
> m1 <- detritus
> m1[1] <- m1[1] + pp$radii[4,1]
> mid <- straightarrow (to = m1, from = m2, arr.type = "triangle",
   arr.pos = 0.4, lwd = 1)
> text(mid[1], mid[2]+0.03, "f6", cex = 0.8)
# flow2->detritus
> m2 <- 0.5*(zoo+phyto)
> m1 <- detritus
> m1[1] <- m1[1] + pp$radii[3,1]*0.2
> mid <- straightarrow (to = m1, from = m2, arr.type = "triangle",
   arr.pos = 0.3, lwd = 1)
> text(mid[1]-0.01, mid[2]+0.03, "f3", cex = 0.8)
```
2.3. Plotting a transition matrix

The next example uses formulae to label the arrows. This is done by passing a data.frame rather than a matrix to function `plotmat`

```r
> # Create population matrix
> 
> #
> Numgenerations <- 6
> DiffMat <- matrix(data = 0, nrow = Numgenerations, ncol = Numgenerations)
> AA <- as.data.frame(DiffMat)
> AA[[1,4]] <- "f[3]"
> AA[[1,5]] <- "f[4]"
> AA[[1,6]] <- "f[5]"
> #
> AA[[2,1]] <- "s[list(0,1)]"
> AA[[3,2]] <- "s[list(1,2)]"
> AA[[4,3]] <- "s[list(2,3)]"
> AA[[5,4]] <- "s[list(3,4)]"
> AA[[6,5]] <- "s[list(4,5)]"
> #
> name <- c(expression(Age[0]), expression(Age[1]), expression(Age[2]),
> expression(Age[3]), expression(Age[4]), expression(Age[5]))
> #
> plotmat(A = AA, pos = 6, curve = 0.7, name = name, lwd = 2,
> + arr.len = 0.6, arr.width = 0.25, my = -0.2,
> + box.size = 0.05, arr.type = "triangle", dtext = 0.95,
> + main = "Age-structured population model 1")
```

2.4. Another transition matrix

The data set `Teasel` contains the transition matrix of the population dynamics model of teasel (Dipsacus sylvestris), a European perennial weed, (Caswell 2001; Soetaert and Herman 2009b)

```r
> Teasel

<table>
<thead>
<tr>
<th></th>
<th>DS 1yr</th>
<th>DS 2yr</th>
<th>R small</th>
<th>R medium</th>
<th>R large</th>
<th>F</th>
</tr>
</thead>
<tbody>
<tr>
<td>DS 1yr</td>
<td>0.000</td>
<td>0.000</td>
<td>0.000</td>
<td>0.000</td>
<td>0.000</td>
<td>322.380</td>
</tr>
<tr>
<td>DS 2yr</td>
<td>0.966</td>
<td>0.000</td>
<td>0.000</td>
<td>0.000</td>
<td>0.000</td>
<td>0.000</td>
</tr>
<tr>
<td>R small</td>
<td>0.013</td>
<td>0.01</td>
<td>0.125</td>
<td>0.000</td>
<td>0.000</td>
<td>3.448</td>
</tr>
<tr>
<td>R medium</td>
<td>0.007</td>
<td>0.00</td>
<td>0.125</td>
<td>0.238</td>
<td>0.000</td>
<td>30.170</td>
</tr>
<tr>
<td>R large</td>
<td>0.008</td>
<td>0.00</td>
<td>0.038</td>
<td>0.245</td>
<td>0.167</td>
<td>0.862</td>
</tr>
<tr>
<td>F</td>
<td>0.000</td>
<td>0.00</td>
<td>0.000</td>
<td>0.023</td>
<td>0.750</td>
<td>0.000</td>
</tr>
</tbody>
</table>
```

This dataset is plotted using curved arrows; we specify the curvature in a matrix called `curves`.

---

1This is now possible thanks to Yvonnick Noel, Univ. Rennes, France
3. plotweb - plotting webs based on matrix input

Given a matrix containing flow values (from rows to columns), function `plotweb` will plot a web. The elements are positioned on a circle, and connected by arrows; the magnitude of web flows determines the thickness of the arrow.

This function is less flexible than `plotmat`, although it does allow to color the arrows differently.

```r
> BB <- matrix(nrow = 20, ncol = 20, 1:20)
> diag(BB) <- 0
```
Teasel population model

Figure 4: The Teasel data set
3.1. Foodwebs

Dataset Rigaweb ((Donali et al. 1999) contains flow values for the food web of the Gulf of Riga planktonic system.

> Rigaweb

<table>
<thead>
<tr>
<th></th>
<th>P1</th>
<th>P2</th>
<th>B</th>
<th>N</th>
<th>Z</th>
<th>D</th>
<th>DOC</th>
<th>CO2</th>
</tr>
</thead>
<tbody>
<tr>
<td>P1</td>
<td>0.0000</td>
<td>0.0000</td>
<td>0.0000</td>
<td>4.12297</td>
<td>10.49431</td>
<td>0.000000</td>
<td>1.565910</td>
<td>17.22501</td>
</tr>
<tr>
<td>P2</td>
<td>0.0000</td>
<td>0.0000</td>
<td>0.0000</td>
<td>0.00000</td>
<td>16.79755</td>
<td>4.457164</td>
<td>2.723090</td>
<td>29.95399</td>
</tr>
</tbody>
</table>


4. functions to create flow charts

The various functions are given in table (1)\(^2\).

The code below generates a flow chart

```r
> plotweb(Rigaweb, main = "Gulf of Riga food web",
+         sub = "mgC/m3/d", val = TRUE)

\(^2\)textparallel was implemented by Michael Folkes, Canada
Figure 6: The Gulf of Riga data set
Table 1: Summary of flowchart functions

<table>
<thead>
<tr>
<th>Function</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>openplotmat</td>
<td>creates an empty plot</td>
</tr>
<tr>
<td>coordinates</td>
<td>calculates coordinates of elements, neatly arranged in rows/columns</td>
</tr>
<tr>
<td>bentarrow</td>
<td>adds 2-segmented arrow between two points</td>
</tr>
<tr>
<td>curvedarrow</td>
<td>adds curved arrow between two points</td>
</tr>
<tr>
<td>segmentarrow</td>
<td>adds 3-segmented arrow between two points</td>
</tr>
<tr>
<td>selfarrow</td>
<td>adds a circular self-pointing arrow</td>
</tr>
<tr>
<td>splitarrow</td>
<td>adds a branched arrow between several points</td>
</tr>
<tr>
<td>straightarrow</td>
<td>adds straight arrow between two points</td>
</tr>
<tr>
<td>treearrow</td>
<td>adds dendrogram-like branched arrow between several points</td>
</tr>
<tr>
<td>shadowbox</td>
<td>adds a box with a shadow to a plot</td>
</tr>
<tr>
<td>textdiamond</td>
<td>adds lines of text in a diamond-shaped box to a plot</td>
</tr>
<tr>
<td>textellipse</td>
<td>adds lines of text in an ellipse-shaped box to a plot</td>
</tr>
<tr>
<td>textempty</td>
<td>adds lines of text on a colored background to a plot</td>
</tr>
<tr>
<td>texthexa</td>
<td>adds lines of text in a hexagonal box to a plot</td>
</tr>
<tr>
<td>textmulti</td>
<td>adds lines of text in a multigonal box to a plot</td>
</tr>
<tr>
<td>textparallel</td>
<td>adds lines of text in a parallelogram to a plot</td>
</tr>
<tr>
<td>textplain</td>
<td>adds lines of text to a plot</td>
</tr>
<tr>
<td>textround</td>
<td>adds lines of text in a rectangular-shaped box to a plot</td>
</tr>
</tbody>
</table>

The different types of text boxes are generated with the following code:

```r
> openplotmat(main = "textbox shapes")
> rx <- 0.1
> ry <- 0.05
> pos <- coordinates(c(1, 1, 1, 1, 1, 1, 1, 1), mx = -0.2)
> textdiamond(mid = pos[1,], radx = rx, rady = ry, lab = LETTERS[1],
+    cex = 2, shadow.col = "lightblue")
```

```r
> openplotmat(main = "textbox shapes")
> rx <- 0.1
> ry <- 0.05
> pos <- coordinates(c(1, 1, 1, 1, 1, 1, 1, 1), mx = -0.2)
> textdiamond(mid = pos[1,], radx = rx, rady = ry, lab = LETTERS[1],
+    cex = 2, shadow.col = "lightblue")
```
Figure 7: A flow chart
5. functions to draw electrical networks

Since version 1.6, it is possible to use `diagram` to draw electrical networks. Below I give an example of a small transistor circuit.

```r
> layoutmat <- matrix(data = c(rep(1, 12), 2, 3, 4, 5),
+ nrow = 4, ncol = 4, byrow = TRUE)
```
Figure 8: The text boxes
Figure 9: The arrow types
> nf <- layout(layoutmat, respect = FALSE)
> par(lwd = 1.5)
> par(mar = c(0, 0, 2, 0))
> emptyplot(main = "transistor Amplifier", asp = FALSE, +    ylim = c(-0.1, 1), xlim = c(-0.1, 1.1))
> x1 <- 0; x2 <- 0.2; x3 <- 0.4; x4 <- 0.6; x5 <- 0.8; x6 <- 1
> y1 <- 0.05; y2 <- 0.4; y3 <- 0.5; y4 <- 0.6; y5 <- 0.95
> x23 <- (x2 + x3)/2
> x56 <- (x5 + x6)/2
> lines(c(x2, x6, x6, x2, x1, x1, x23, x3, x3), 
>    c(y1, y1, y5, y5, y1, y1, y3, y3, y4, y5))
> lines(c(x23, x3, x3), c(y3, y2, y1))
> lines(c(x3, x4, x4), c(y2, y2, y1))
> lines(c(x3, x5, x5), c(y4, y4, y1))
> en.Amplifier(c(x23, y3), r = 0.035)
> en.Signal(c(x1, 0.2), lab = expression("U"["in"]))
> en.Signal(c(x6, y2), lab = expression("U"["b"]))
> straightarrow(c(x1 - 0.05, 0.23), c(x1 - 0.05, 0.17),
>    arr.pos = 1, arr.type = "triangle", lwd = 1)
> straightarrow(c(x6 + 0.05, y2 + 0.03), c(x6 + 0.05, y2 - 0.03),
>    arr.pos = 1, arr.type = "triangle", lwd = 1)
> en.Node(c(x1, y3), lab = "u1")
> en.Node(c(x2, y3), lab = "u2")
> en.Node(c(x3, y2), lab = "u3", pos = 1.5)
> en.Node(c(x3, y4), lab = "u4", pos = 2.5)
> en.Node(c(x5, y4), lab = "u5")
> en.Capacitator(c(0.5*(x1 + x2), y3), lab = "C1", vert = FALSE)
> en.Capacitator(c(x4, y4), lab = "C3", vert = FALSE)
> en.Capacitator(c(x4, 0.5*(y1+y2)), lab = "C2", vert = TRUE)
> en.Resistor(c(x1, y2), lab = "R0")
> en.Resistor(c(x2, 0.5*(y1+y2)), lab = "R1")
> en.Resistor(c(x2, 0.5*(y4+y5)), lab = "R2")
> en.Resistor(c(x3, 0.5*(y4+y5)), lab = "R4")
> en.Resistor(c(x3, 0.5*(y1+y2)), lab = "R3")
> en.Resistor(c(x5, 0.5*(y1+y2)), lab = "R5")
> en.Ground(c(1.0, 0.05))
> par(mar=c(2, 2, 2, 2))
> emptyplot(main = "transistor")
> lines(c(0.1, 0.5, 0.9), c(0.5, 0.5, 0.9))
> lines(c(0.5, 0.9), c(0.5, 0.1))
> lines(c(0.5, 0.5), c(0.4, 0.6))
> text(0.2, 0.4, "Gate", font = 3)
> text(0.8, 0.9, "Drain", font = 3, adj = 1)
> text(0.8, 0.1, "Source", font = 3, adj = 1)
> en.Amplifier(c(0.5, 0.5), r = 0.15)
> box(col = "grey")
> emptyplot(main = "capacitator")
This vignette was created using Sweave (Leisch 2002).
The package is on CRAN, the R-archive website ((R Development Core Team 2008))
More examples can be found in the demo’s of package ecolMod (Soetaert and Herman 2009a)

References


Figure 10: Drawing an electrical network with package diagram


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