

Overview of the spnet package

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Introduction

Social networks analysis has received special attention over the past decade, and a lot of tools for manipulating and rendering social networks have emerged. In several situations a social network is associated with a spatial dimension, and behaviors observed within the network cannot be interpreted without taking into account the location of each of its nodes regarding to the other nodes. This is the case, for example, when studying inflows/outflows between cities or companies, or when studying people debating in a room. Based on the `sp` package, which provides efficient classes for storing spatial data and methods for handling and rendering them, the `spnet` package aims at facilitating the rendering of (social) networks on maps. Furthermore, fixing network node positions allows to more easily monitor time-varying networks and observe how connections and flows evolve over time.

The `spnet` package offers methods for dealing with spacial social networks. It allows to plot networks for which actors have a specific location on a map (participants in a political debate, cities, etc.). `SpatialPolygons` objects from the `sp` package are supported.

```
global.par.mar <- c(0,0,0,0)
```

Gallery

The `spnet.example.basic` function provides a working example involving basic fonctionnalities of the `spnet` package. Don't hesitate to look at its code and take what you need.

```
net1 <- spnet.example.basic()
plot(net1)
```

Tips

- By default, arrows start and end in the centre of gravity of areas. If you use labels, this behaviour can make labels unreadable. To fix this you can (1) shorten arrows with `graph.network.arrow.shorten` or/and (2) shift on *x* and *y* axis with respectively `graph.network.arrow.shift.x` and `graph.network.arrow.shift.y`.

Usage

Creating a `spnet` object

Creating a new `spnet` object (formal class `SpatialNetwork`) is done by the `spnet.create` function. You will have to supply two required parameters: the node IDs, and the corresponding position IDs on the map (which will be set in a second step). Note that in a `spnet` object data are stored in a `data.frame`. In this `data.frame` the two required parameters will be supplied as variables 'NODE' and 'POSITION'. Here is an example:

```
node <- c("France", "United States", "Brazil", "Australia")

# We load the world map
data(world.map.simplified)

# We retrieve position IDs for our nodes
# As countries are numbered from 0 to 245 on the map, we subtract 1
```

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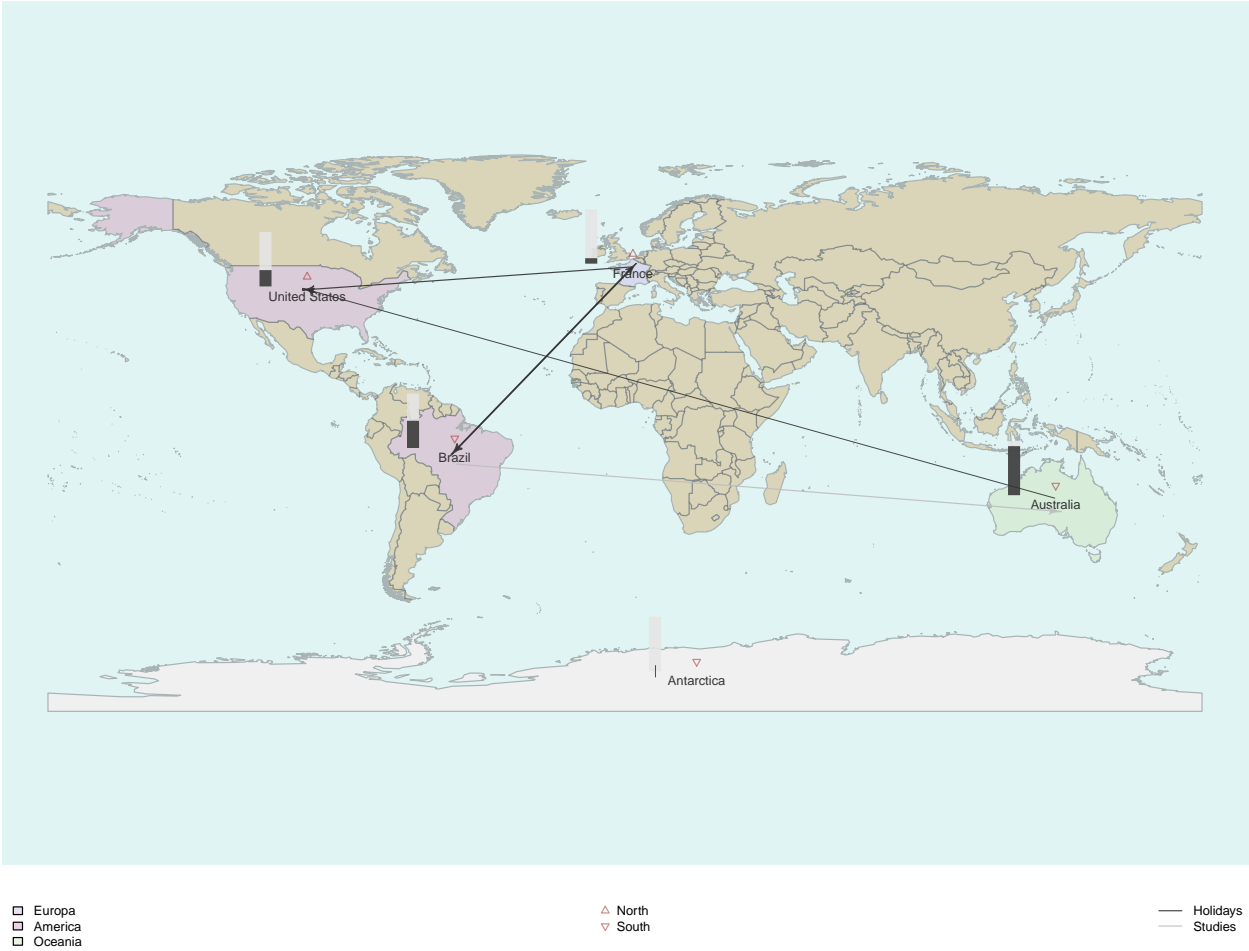


Figure 1:

```
position <- match(node, world.map.simplified@data[, 'NAME']) - 1

net1 <- spnet.create(
  data.frame(
    'NODE' = node,
    'POSITION' = position
  )
)

class(net1)
```

```
## [1] "SpatialNetwork"
## attr(,"package")
## [1] "spnet"
```

Setting a map

The next step is to define the map to use for plotting. This is done with the `graph.map` function. Currently only `SpatialPolygons` objects are supported. If your map comes on another format (e. g. a shape file .shp) you can use instead the facilities of the `maptools` package to convert your map data into a `SpatialPolygons` object.

Setting labels

The IDs for nodes have to be valid variable names. You should avoid special character or spaces. Currently network data have to be provide as a `matrix`, with node IDs in row names an column names. Thus, the `spnet` object will automatically match your network data with the node positions on the map. To be valid row and column names, node IDs must not contain special characters or spaces.

In case you need to have more suitable label for your nodes when plotting the `spnet` object (e.g. containing spaces or accented characters), here is the protocol to follow:

First, we start with our basic `spnet` object containing a simple room map:

```
net1 <- spnet.example.basic.map()
plot(net1)
```

Then we introduce the data. In this case we use the following frame/matrix as an example:

```
data.frame(net1)
```

```
##           NODE POSITION
## 1      France      64
## 2 United States  208
## 3      Brazil    20
## 4   Australia    8
## 5   Antarctica  144
```

```
net1$label <- c("France", "United States", "Brazil", "Australia", "Antartica")
graph.label.variable(net1) <- 'label'
```

Now you can change the size and the color of labels:

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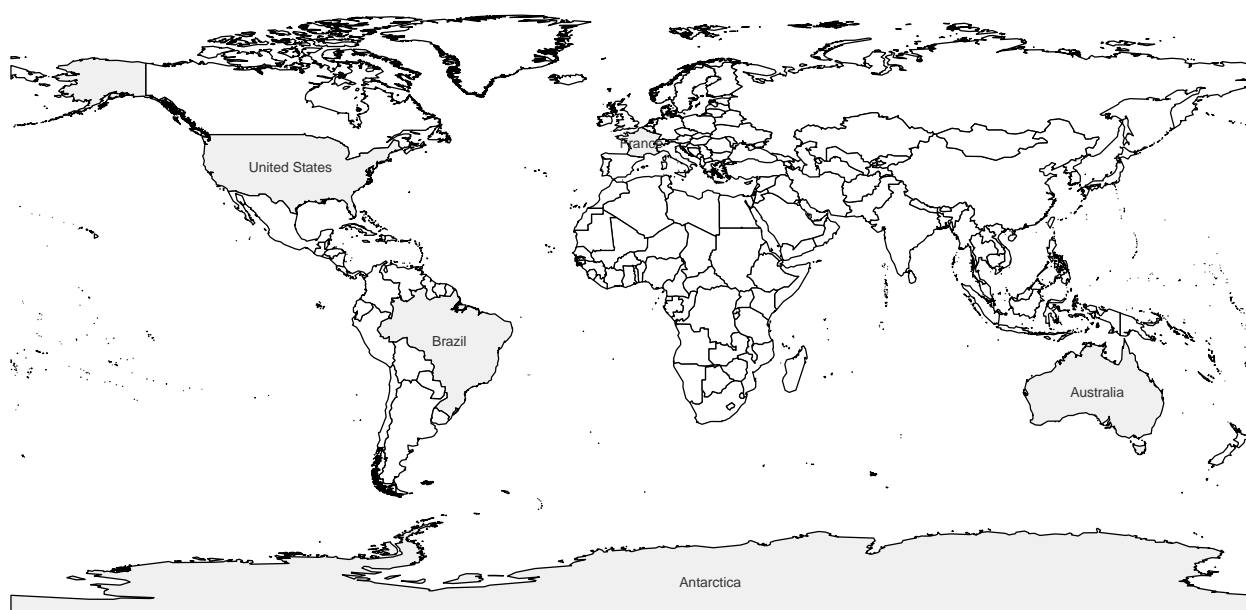


Figure 2:

```
graph.label.cex(net1) <- 0.7
graph.label.color(net1) <- '#FF0000'

plot(net1)
```

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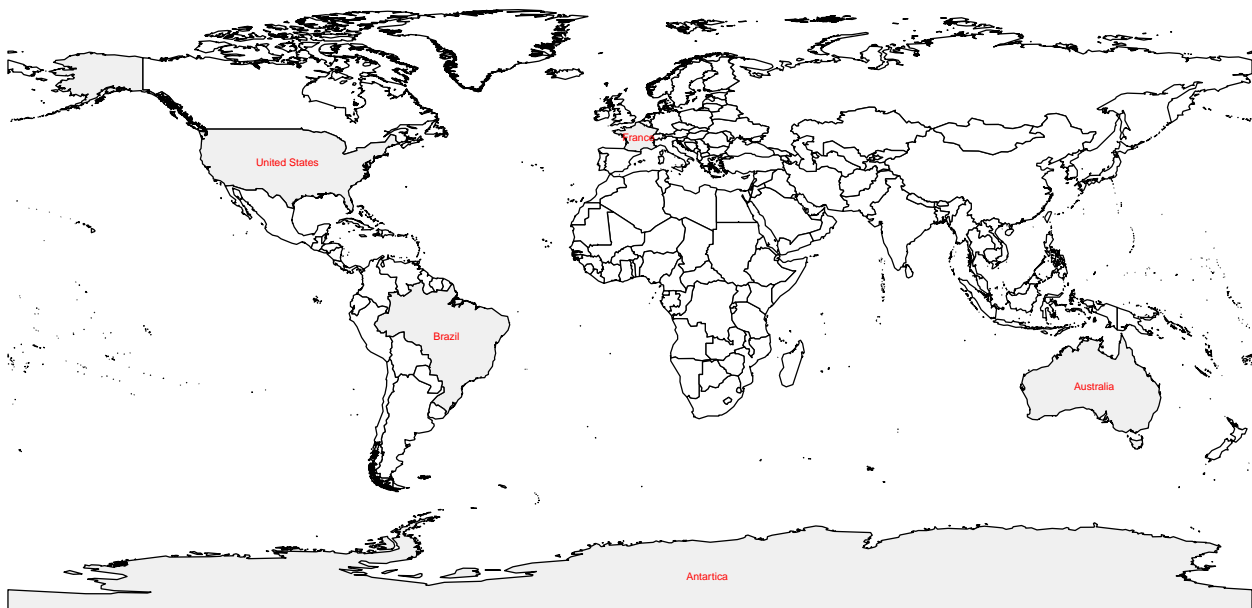


Figure 3:

If you want to disable label printing, you can add an empty variable and use it as the labelling variable:

```
net1$label.empty <- rep("", nrow(net1))
graph.label.variable(net1) = 'label.empty'

plot(net1)
```

Setting colors

To set colors you need:

- A categorical variable affecting each node to a class
- a legend of color specifying the color to use for each class

Here is a practical example. First, we start with our basic `spnet` object containing a map.

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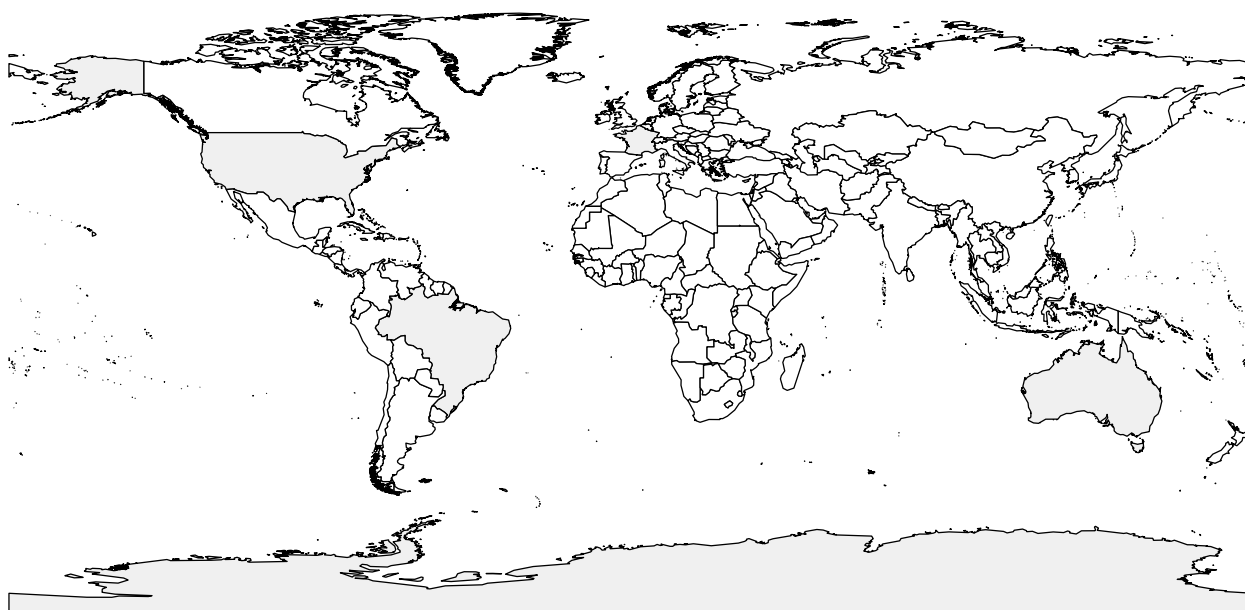


Figure 4:

```
net1 <- spnet.example.basic.map()
plot(net1)
```

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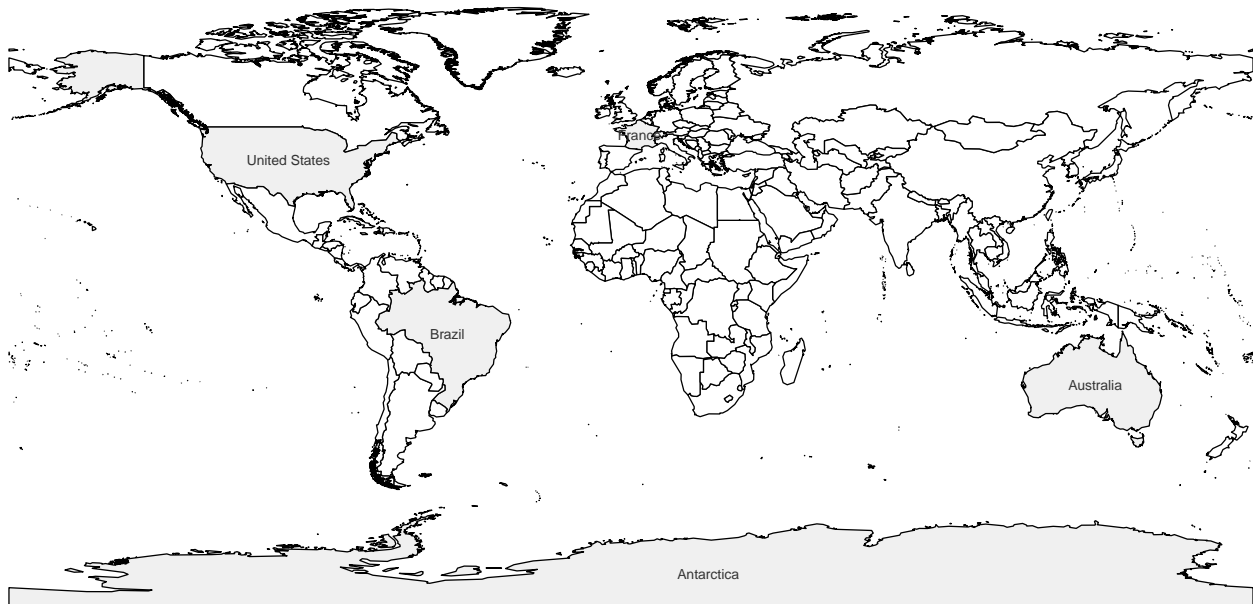


Figure 5:

To make the graphic nicer, we can set the background color, border color, and a default color for coloring the regions:

```
graph.color.background(net1) <- "#B3E4E6" # light blue
graph.color.border(net1) <- "#555555" # grey
graph.color.region(net1) <- "#D2A6F6" # light orange
plot(net1)
```

Now, we add a categorical variable affecting each node to a class:

```
net1$continent <- c("Europa", "America", "America", "Oceania", "Antartica")
```

Therefore, the data is changed to:

```
data.frame(net1)
```

```
##          NODE POSITION continent
## 1         France         64  Europa
```


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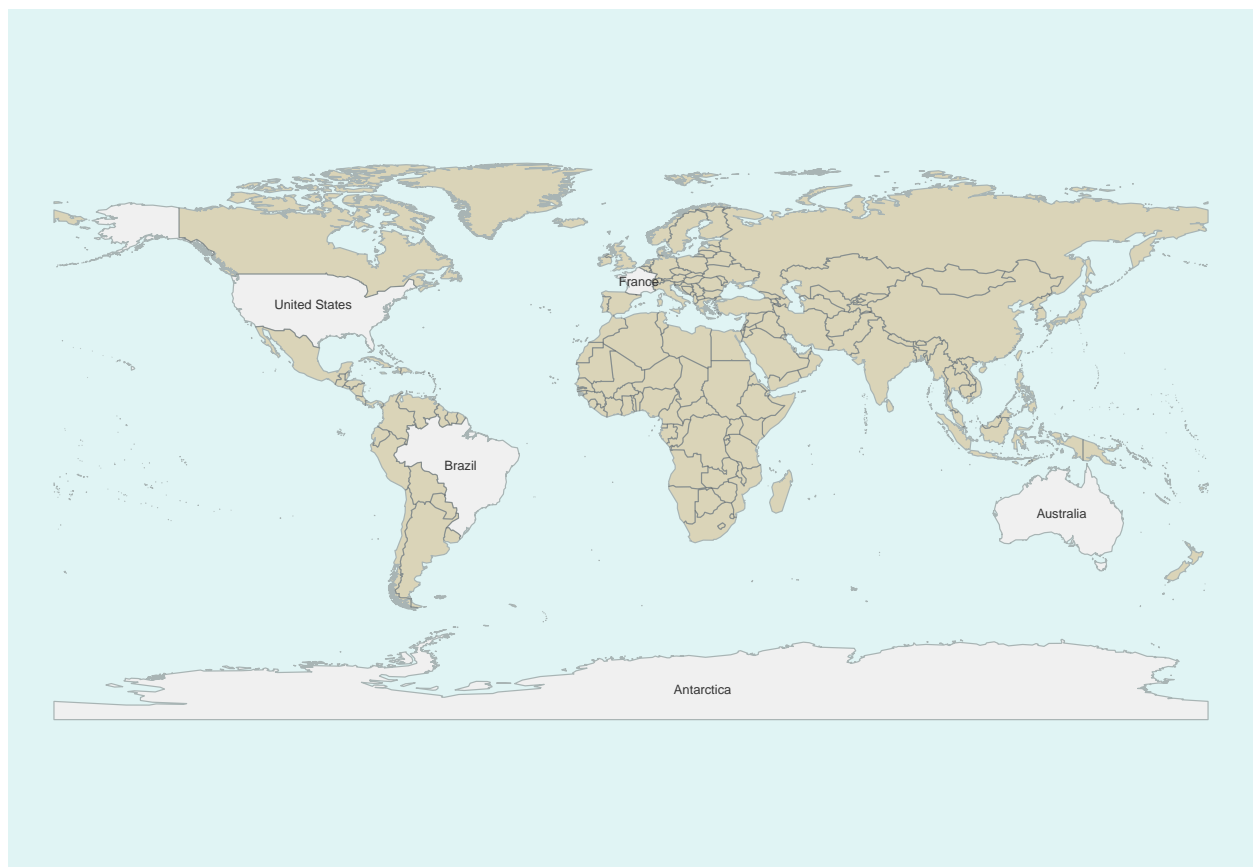


Figure 6:

```
## 2 United States      208   America
## 3      Brazil        20   America
## 4      Australia      8   Oceania
## 5      Antarctica   144 Antarctica
```

Then we specify that we want to use the variable `parti` in order to colorize the map:

```
graph.color.variable(net1) <- "continent"
```

Finally we specify the colors to use:

```
graph.color.legend(net1) <- c('Europa' = "#CBB3E466", 'America' = "#D490B366", 'Oceania' = "#CBE4B366")
```

Now the `plot` function is able to colorize the graphics just by running:

```
plot(net1)
```

Dealing with a quantitative covariate: rendering individual barplots

We may need to render a quantitative attribute related to each node of the network. To that purpose we provide a simple barplot tool. This section details how to use it. We start with a fresh `spnet` object and equip it with a map.

```
ex.bp <- net1
```

A fresh `spnet` object contains the following default barplot settings:

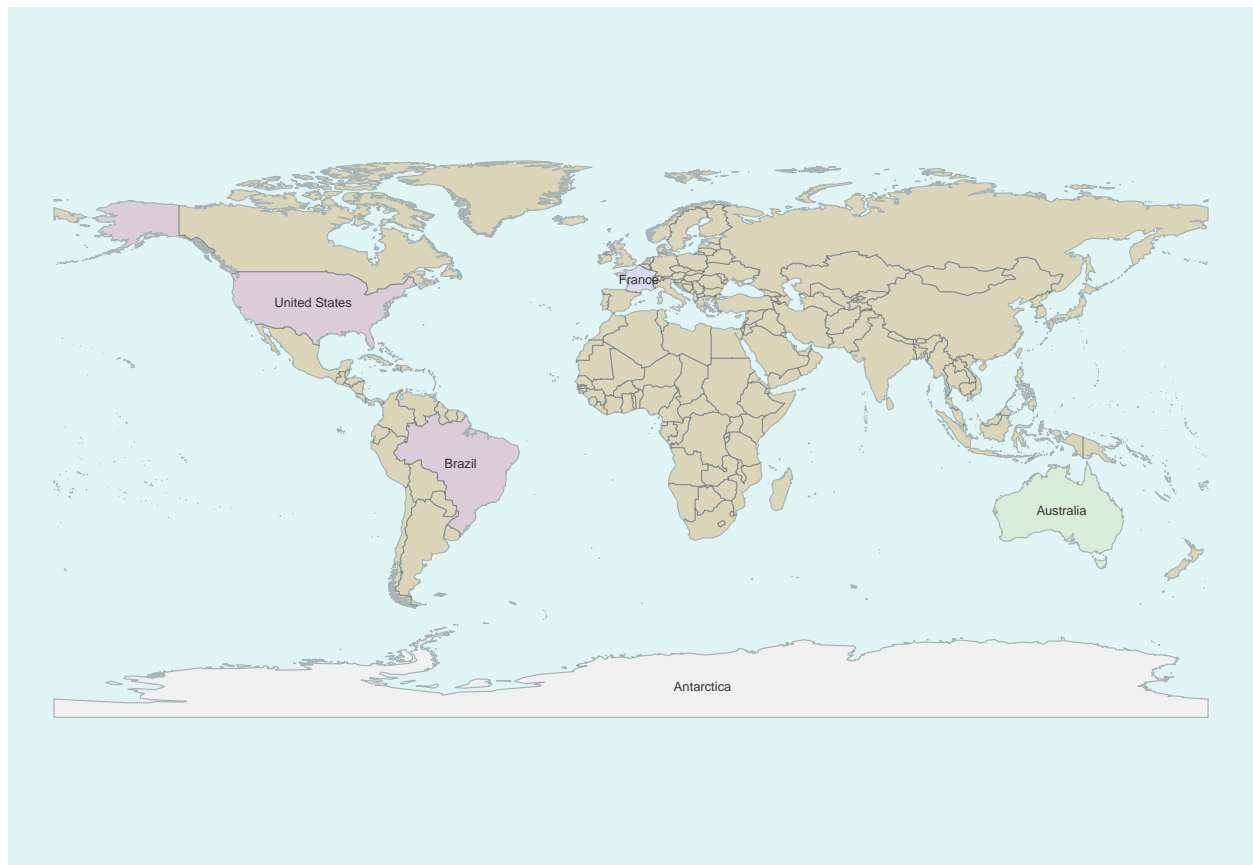
```
graph.barplot.list(ex.bp)
```

```
## $variable
## [1] ""
##
## $bound.lower
## [1] -0.5 -0.5
##
## $bound.upper
## [1] 0.5 -0.5
##
## $fgcolor
## [1] "#666666"
##
## $bgcolor
## [1] "#eeeeee"
##
## $width
## [1] 8
```

The first point is to define a variable to fill in the individual barplot. This variable can either be defined manually in a vector or imported. The value has to be normalized between 0 and 1.

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- Europa
- America
- Oceania

Figure 7:

```
ex.bp$content <- c(0.1,0.3,0.5,0.9,0.0)
ex.bp
```

```
## This is a valid 'SpatialNetwork' object.
##
## - Data: (first rows)
##
##           NODE POSITION continent content
## 1      France         64   Europa    0.1
## 2 United States      208  America    0.3
## 3      Brazil        20   America    0.5
## 4   Australia         8  Oceania    0.9
## 5   Antarctica      144 Antarctica  0.0
##
## - Map:
##   Length: 246
##
## - Plotting options:
##   Variable used to colorize: 'continent'
```

Now the 'content' variable is added into the database.

The second step is to define the barplot variable. This operation is confirmed on the first line of the function `graph.barplot.list()`.

```
graph.barplot.variable(ex.bp) <- "content"
graph.barplot.list(ex.bp)
```

```
## $variable
## [1] "content"
##
## $bound.lower
## [1] -0.5 -0.5
##
## $bound.upper
## [1] 0.5 -0.5
##
## $fgcolor
## [1] "#666666"
##
## $bgcolor
## [1] "#eeeeee"
##
## $width
## [1] 8
```

```
ex.bp
```

```
## This is a valid 'SpatialNetwork' object.
##
## - Data: (first rows)
##
```

```
##          NODE POSITION continent content
## 1      France         64   Europa    0.1
## 2 United States      208  America    0.3
## 3      Brazil        20   America    0.5
## 4    Australia        8   Oceania    0.9
## 5    Antarctica     144 Antarctica  0.0
##
## - Map:
##   Length: 246
##
## - Plotting options:
##   Variable used to colorize: 'continent'
##   Variable used to draw barplots: 'content'
```

```
plot(ex.bp)
```

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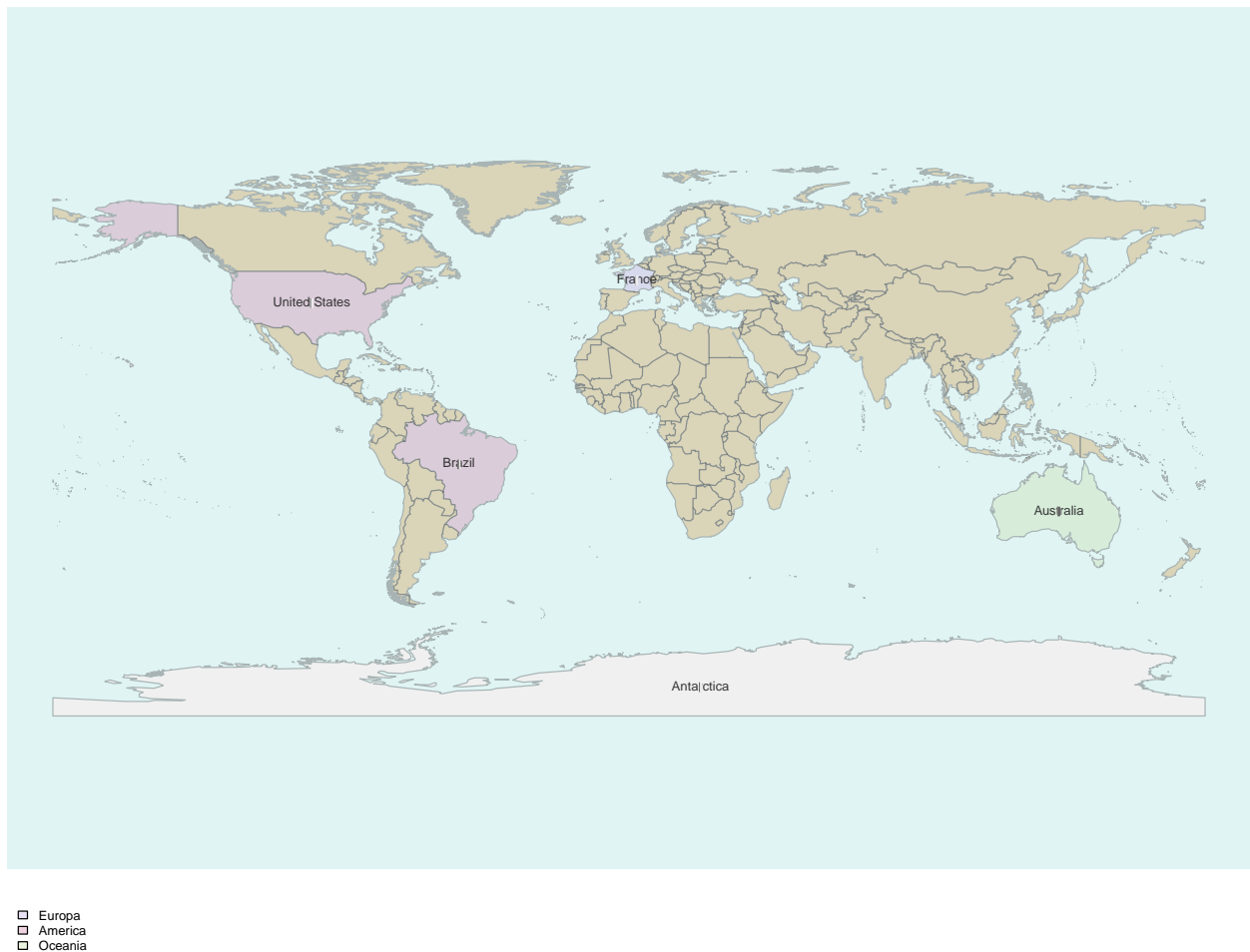


Figure 8:

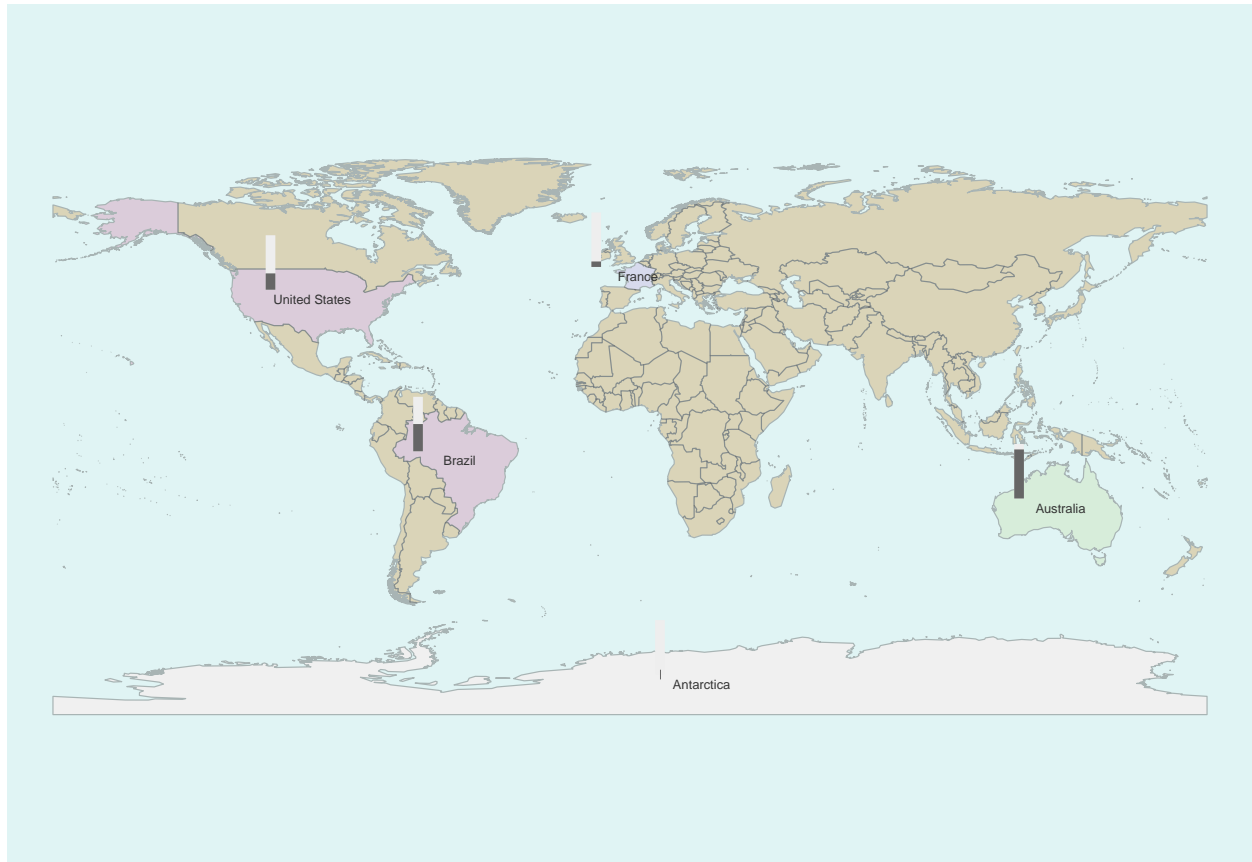
The plot is by default in black and white. On the graphs, individual barplots appear at the bottom of the boxes. To pull them up a little, use the functions `graph.barplot.bound.lower` and `graph.barplot.bound.upper`

to adjust. The coordinate system is computed from the center of the grid (as for the other functions to adjust positions).

```
graph.barplot.bound.upper(ex.bp) <- c(-13,20)
graph.barplot.bound.lower(ex.bp) <- c(-13,3)
plot(ex.bp)
```

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□ Europe
□ America
□ Oceania

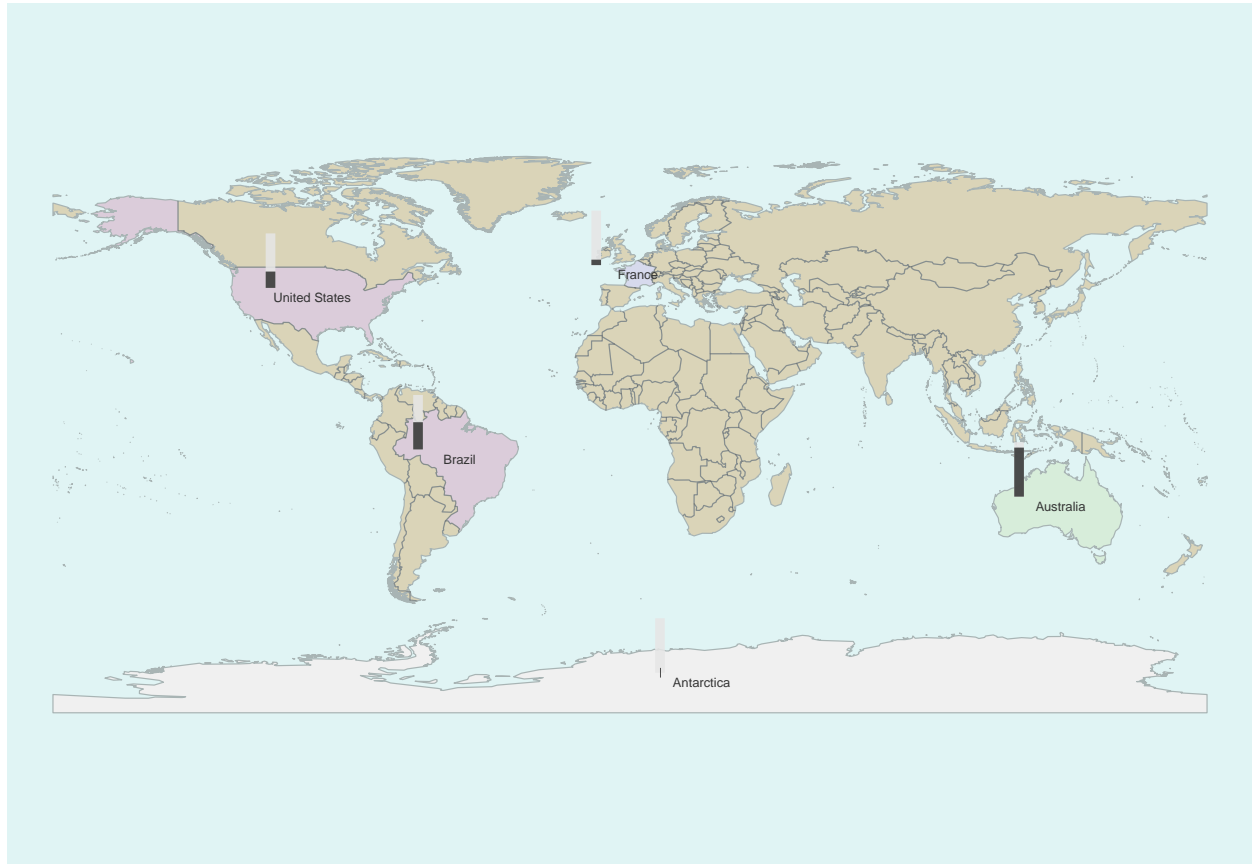
Figure 9:

To change the colors of the barplot, you can define a fill color `graph.barplot.fgcolor` and a background color `graph.barplot.bgcolor`.

```
graph.barplot.fgcolor(ex.bp) <- "#333333DD"
graph.barplot.bgcolor(ex.bp) <- "#E6E6E6DD"
plot(ex.bp)
```

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- Europa
- America
- Oceania

To find colors, you can go to a web repository such as [ColorPicker.com](https://colorpicker.com). To determine opacity, specify a number from 1 (transparent) to 100 (full opacity) at the end of the string.

To change size of the barplot is possible with the `graph.barplot.width` function:

```
graph.barplot.width(ex.bp) <- 10
plot(ex.bp)
```

Put symbols on the map

You can add a symbol for any additional covariate:

```
ex.bp$hemisphere <- c('North', 'North', 'South', 'South', 'South')
graph.symbol.variable(ex.bp) <- 'hemisphere'
graph.symbol.legend(ex.bp) <- c('North' = 'triangle.up', 'South' = 'triangle.down')
plot(ex.bp)
```

To fix size of the symbol:

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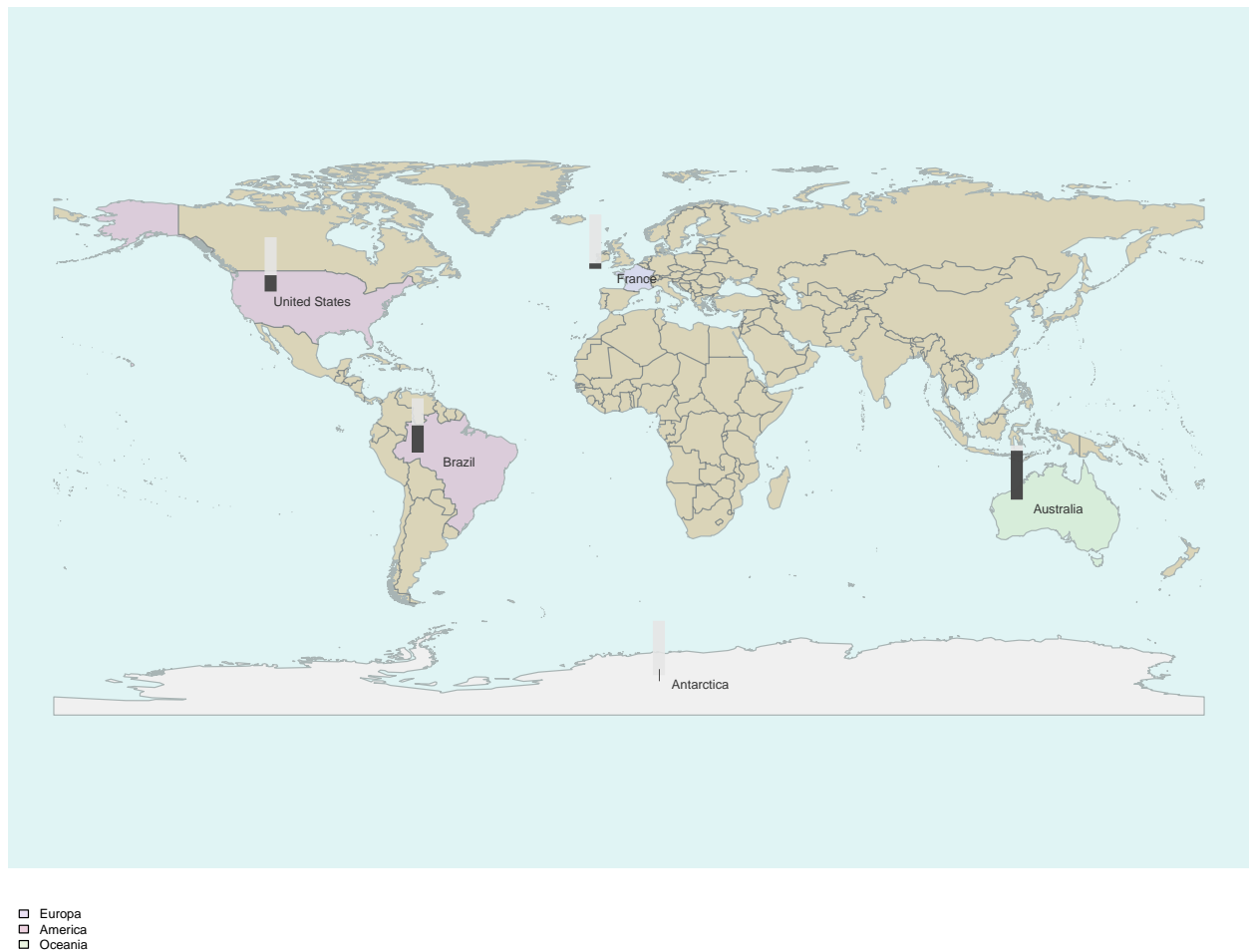


Figure 10:

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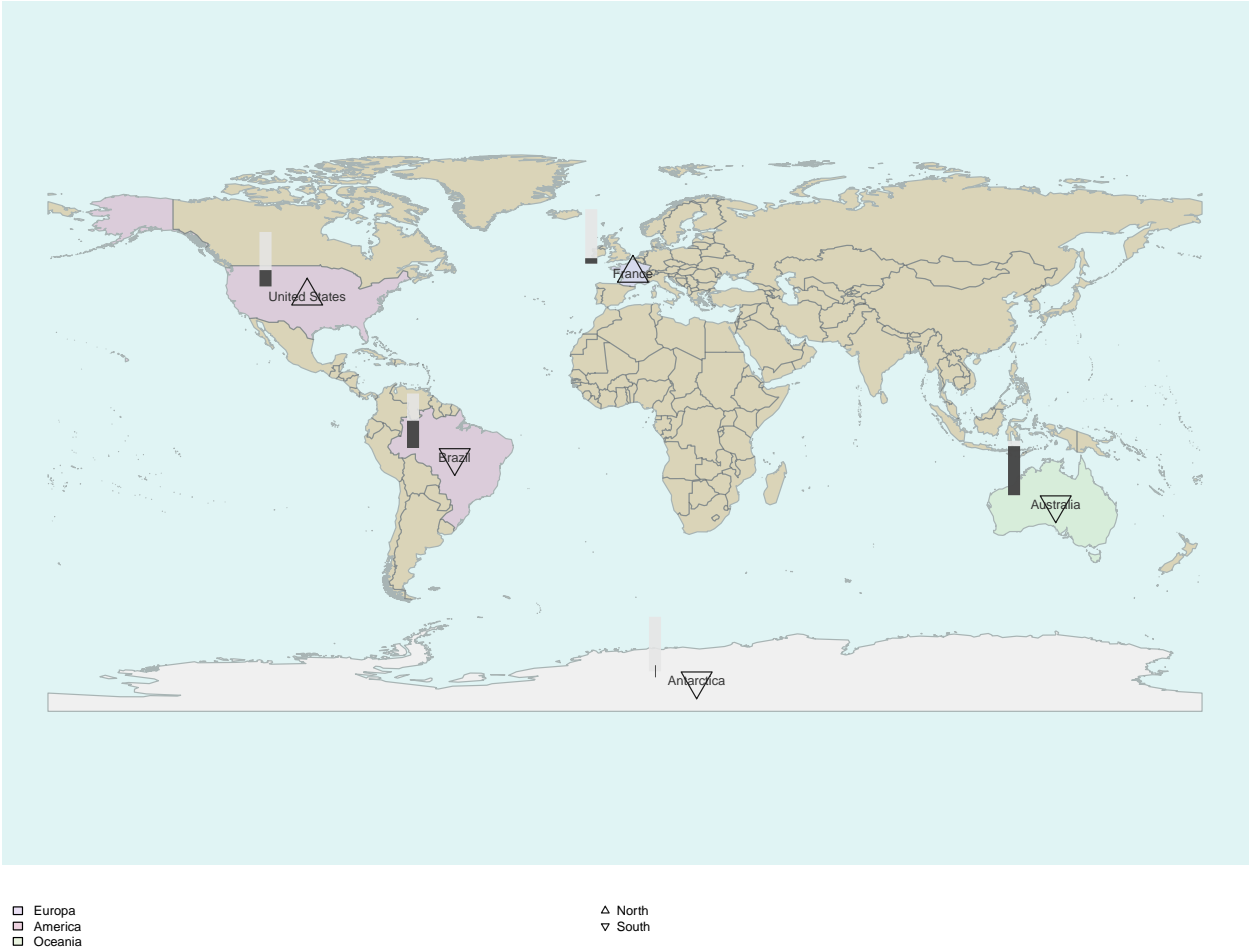


Figure 11:

```
graph.symbol.cex(ex.bp) <- 1
plot(ex.bp)
```

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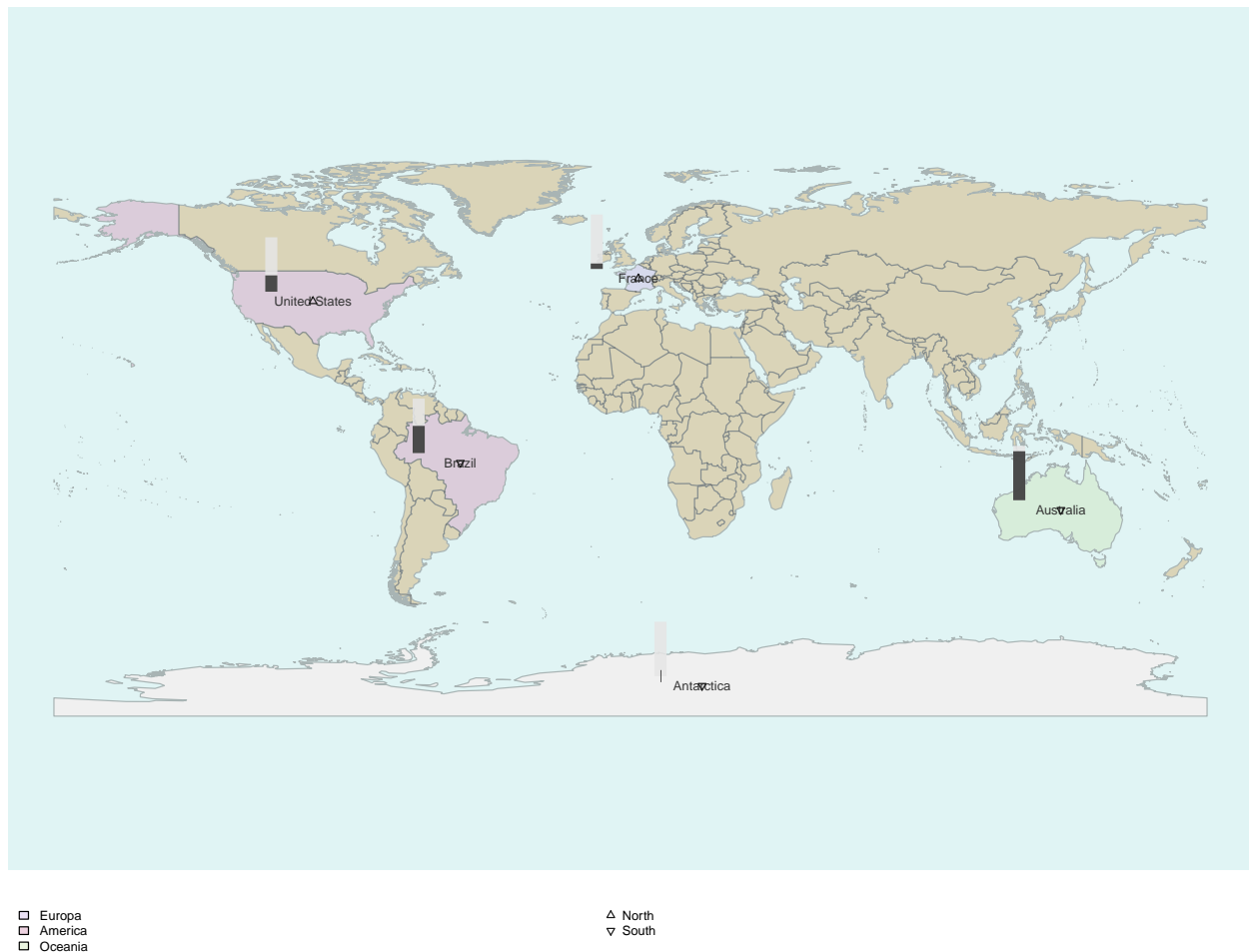


Figure 12:

To fix position of the symbol:

```
graph.symbol.shift.y(ex.bp) <- 4
plot(ex.bp)
```

To fix color of the symbol:

```
graph.symbol.color(ex.bp) <- '#A52A2A88'
plot(ex.bp)
```

Define and draft networks on the graph

To add a network on the map:

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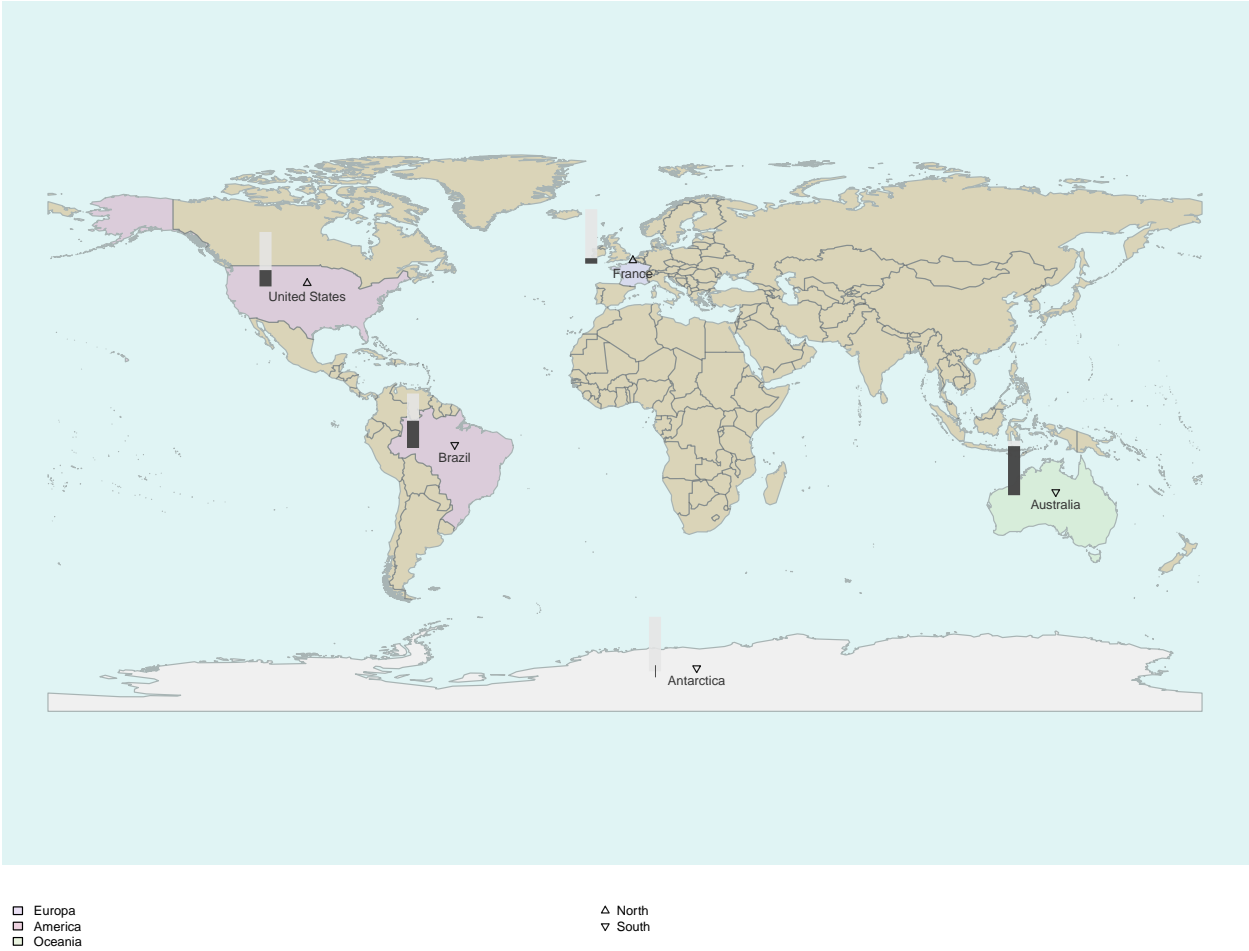


Figure 13:

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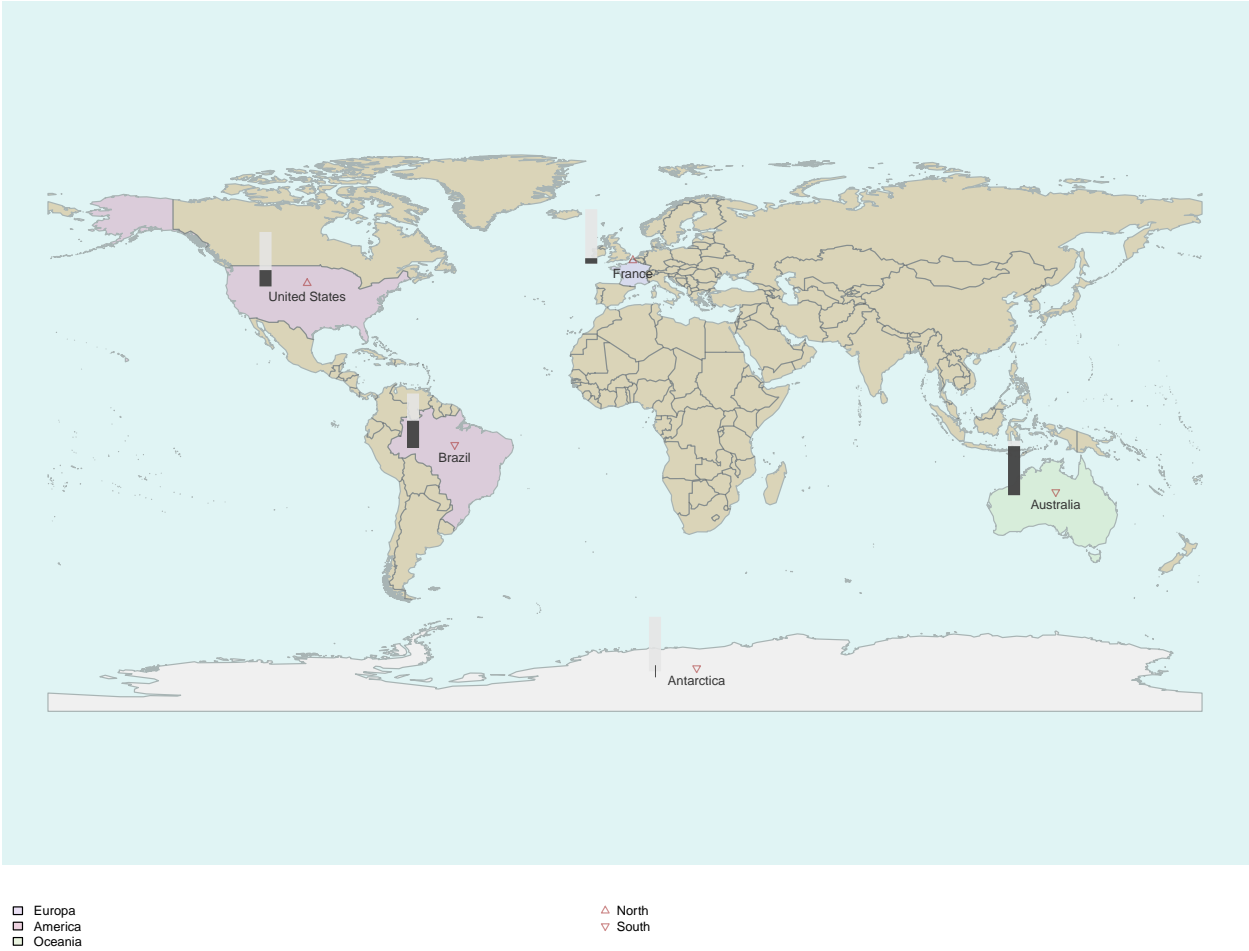


Figure 14:

```
graph.networks.add(ex.bp) <- 'n1'
graph.network.label(ex.bp, 'n1') <- 'Holidays'
ex.bp
```

This is a valid 'SpatialNetwork' object.

##

- Data: (first rows)

##

	NODE	POSITION	continent	content	hemisphere
## 1	France	64	Europa	0.1	North
## 2	United States	208	America	0.3	North
## 3	Brazil	20	America	0.5	South
## 4	Australia	8	Oceania	0.9	South
## 5	Antarctica	144	Antartica	0.0	South

##

- Map:

Length: 246

##

- Network data:

Number of network(s): 1

##

- Plotting options:

Variable used to colorize: 'continent'

Variable used to draw symbols: 'hemisphere'

Variable used to draw barplots: 'content'

In the “Network data” sectionDetail of ex.bp object mentions “Number of network(s): 1”.

Now, to add a second network:

```
graph.networks.add(ex.bp) <- 'n2'
graph.network.label(ex.bp, 'n2') <- 'Studies'
```

We define a matrix manually; for this example we will use:

```
matrix_int <- matrix(
  rep(0, length(node)^2),
  nrow = length(node),
  dimnames = list(node, node)
)
```

And now we attach it to our spnet object:

```
graph.network.data(ex.bp, 'n1') <- matrix_int
graph.network.data(ex.bp, 'n2') <- matrix_int
```

A matrix can also be imported from a data file.

We fill our network manually directly into the spnet object:

```

#Network 1
graph.network.data(ex.bp, 'n1')['France', 'United States'] <- 2
graph.network.data(ex.bp, 'n1')['Australia', 'United States'] <- 1
graph.network.data(ex.bp, 'n1')['France', 'Brazil'] <- 3
graph.network.data(ex.bp, 'n1')['Brazil', 'France'] <- 2

#Network 2
graph.network.data(ex.bp, 'n2')['Brazil', 'Australia'] <- 2
plot(ex.bp)

```

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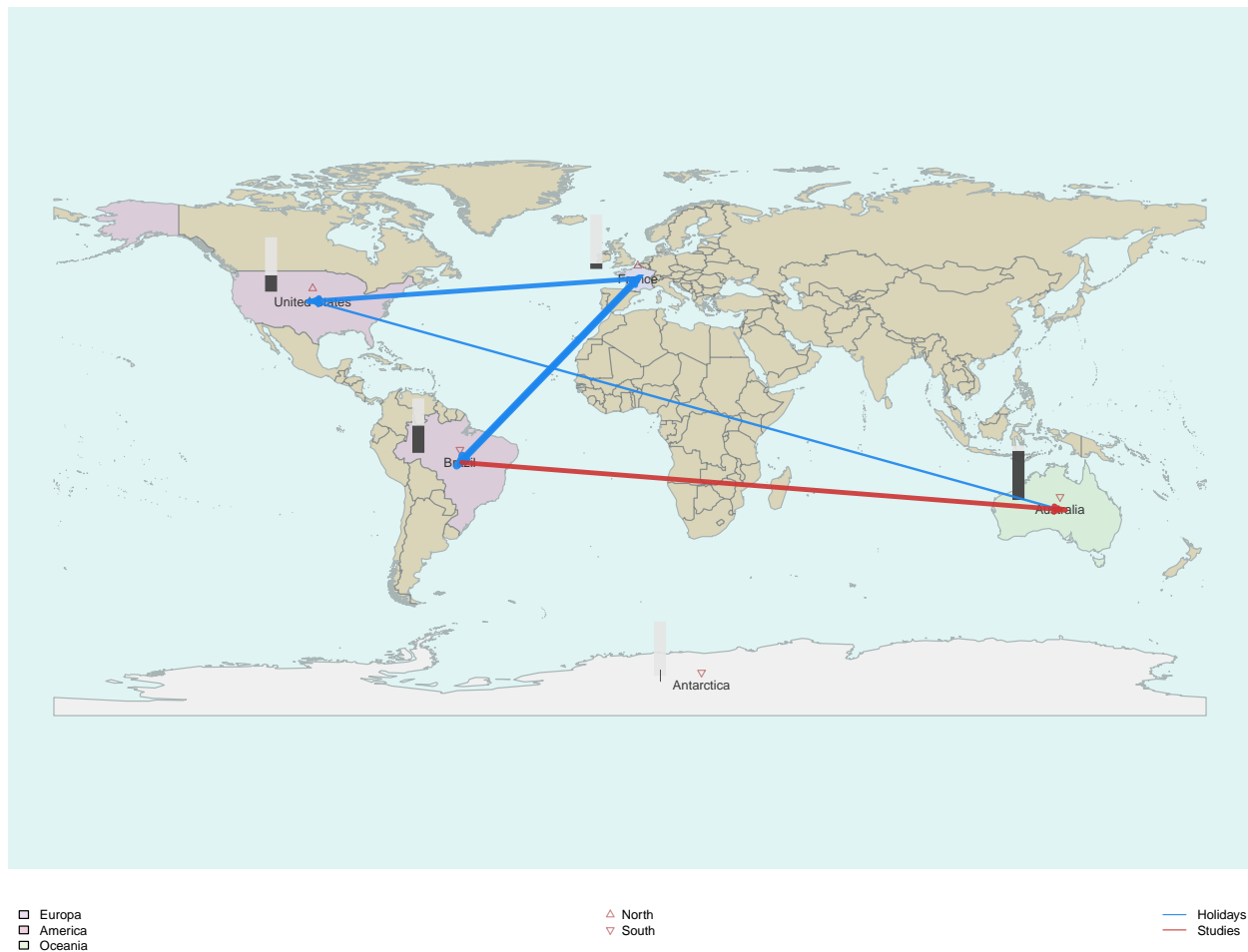


Figure 15:

There are 'by default' colors, but we can change them by using:

```

graph.network.arrow.color(ex.bp, 'n1') <- '#33333366'
graph.network.arrow.color(ex.bp, 'n2') <- 'grey'
plot(ex.bp)

```

We can reduce as well the thickness and length of arrows:

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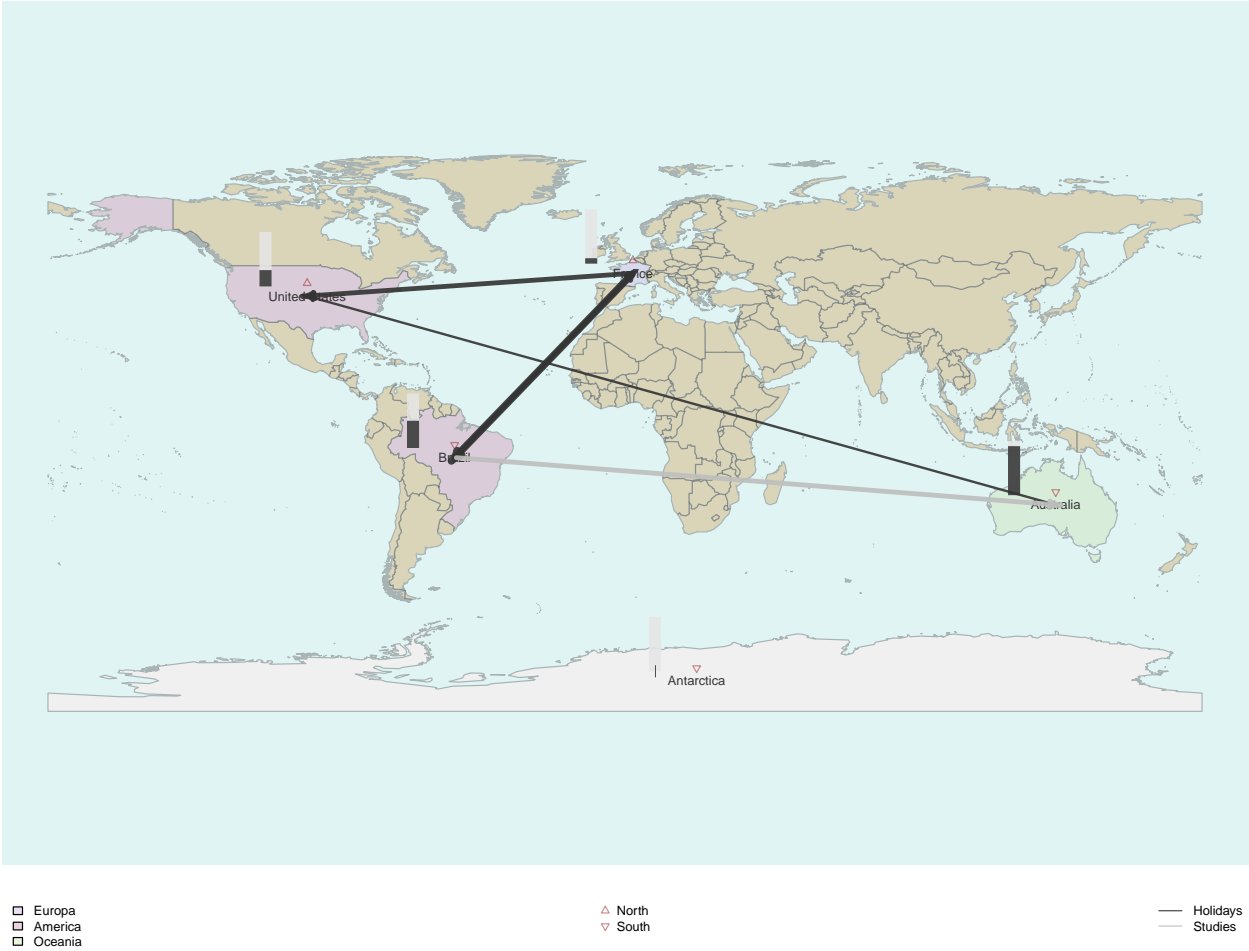


Figure 16:

```
graph.network.arrow.thickness(ex.bp, 'n1') <- 0.5
graph.network.arrow.thickness(ex.bp, 'n2') <- 0.5

graph.network.arrow.shorten(ex.bp, 'n1') <- 0.9
graph.network.arrow.shorten(ex.bp, 'n2') <- 0.9
plot(ex.bp)
```

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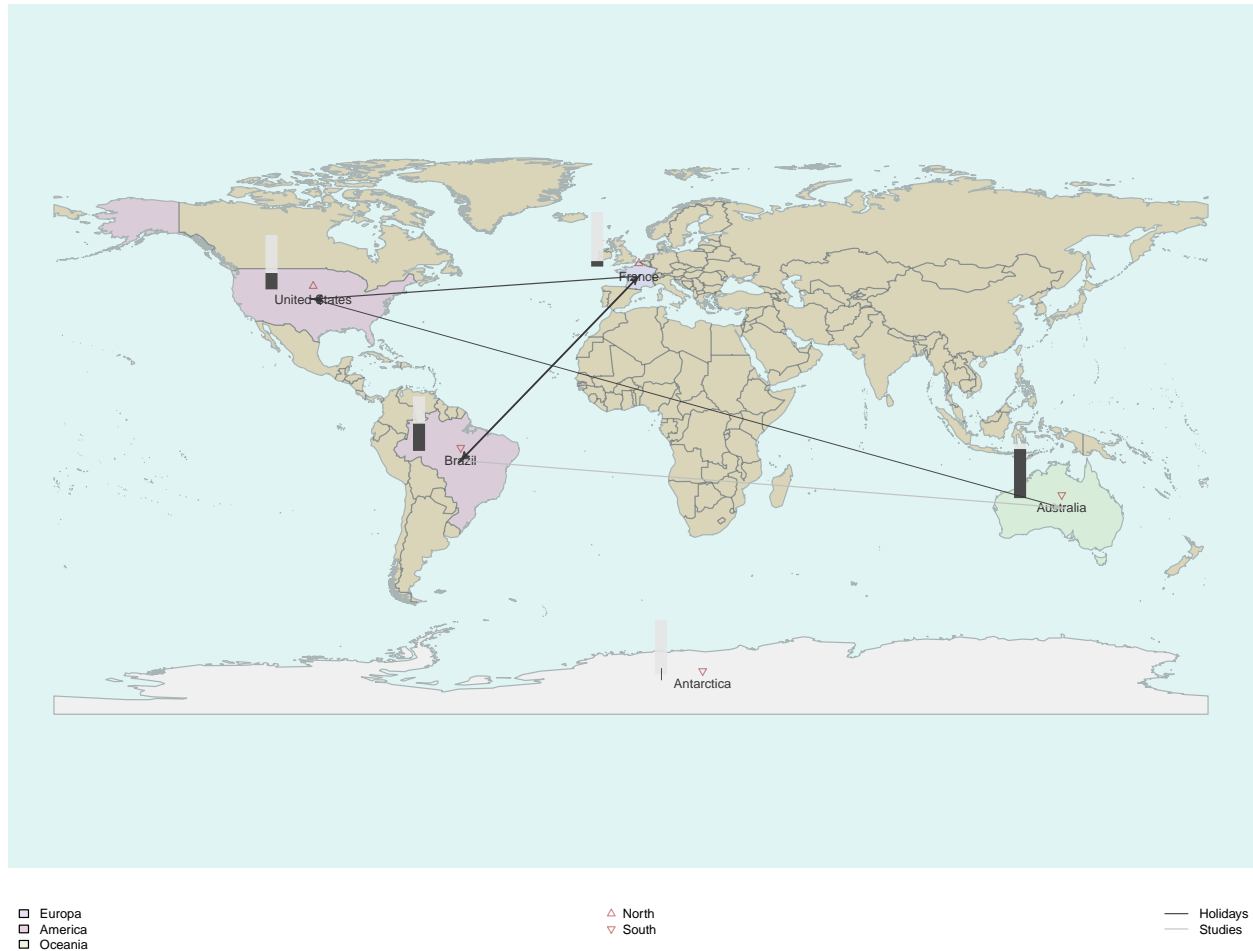


Figure 17:

Some examples to grasp how to edit the graph:

We shift 'Holidays' a little bit upper:

```
graph.network.arrow.shift.y(ex.bp, 'n1') <- 2
plot(ex.bp)
```

And we shift 'Studies' a little bit lower and give it some transparency:

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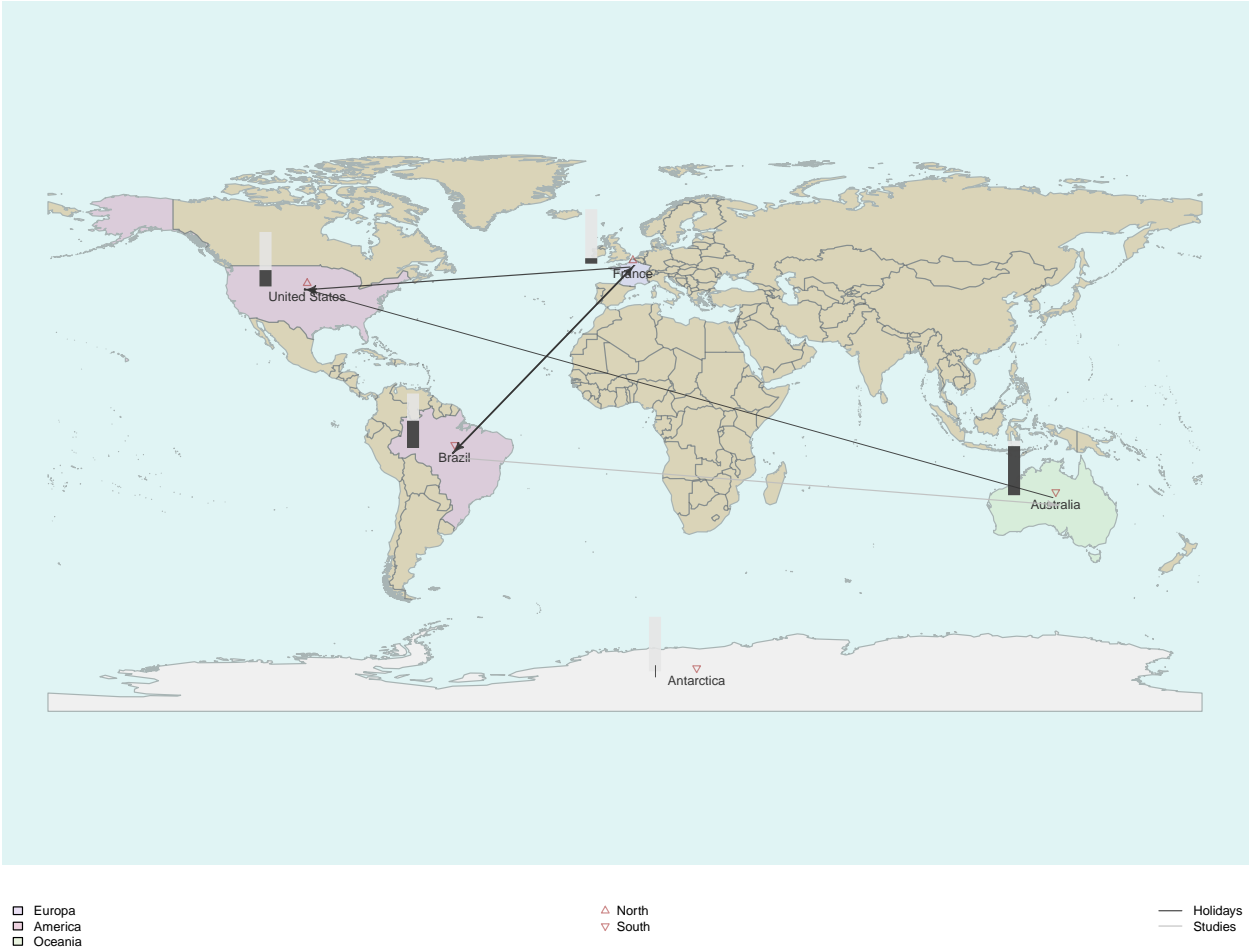


Figure 18:

```
graph.network.arrow.shift(ex.bp, 'n2') <- -2
graph.network.arrow.opacity(ex.bp, 'n2') <- 0.9
plot(ex.bp)
```

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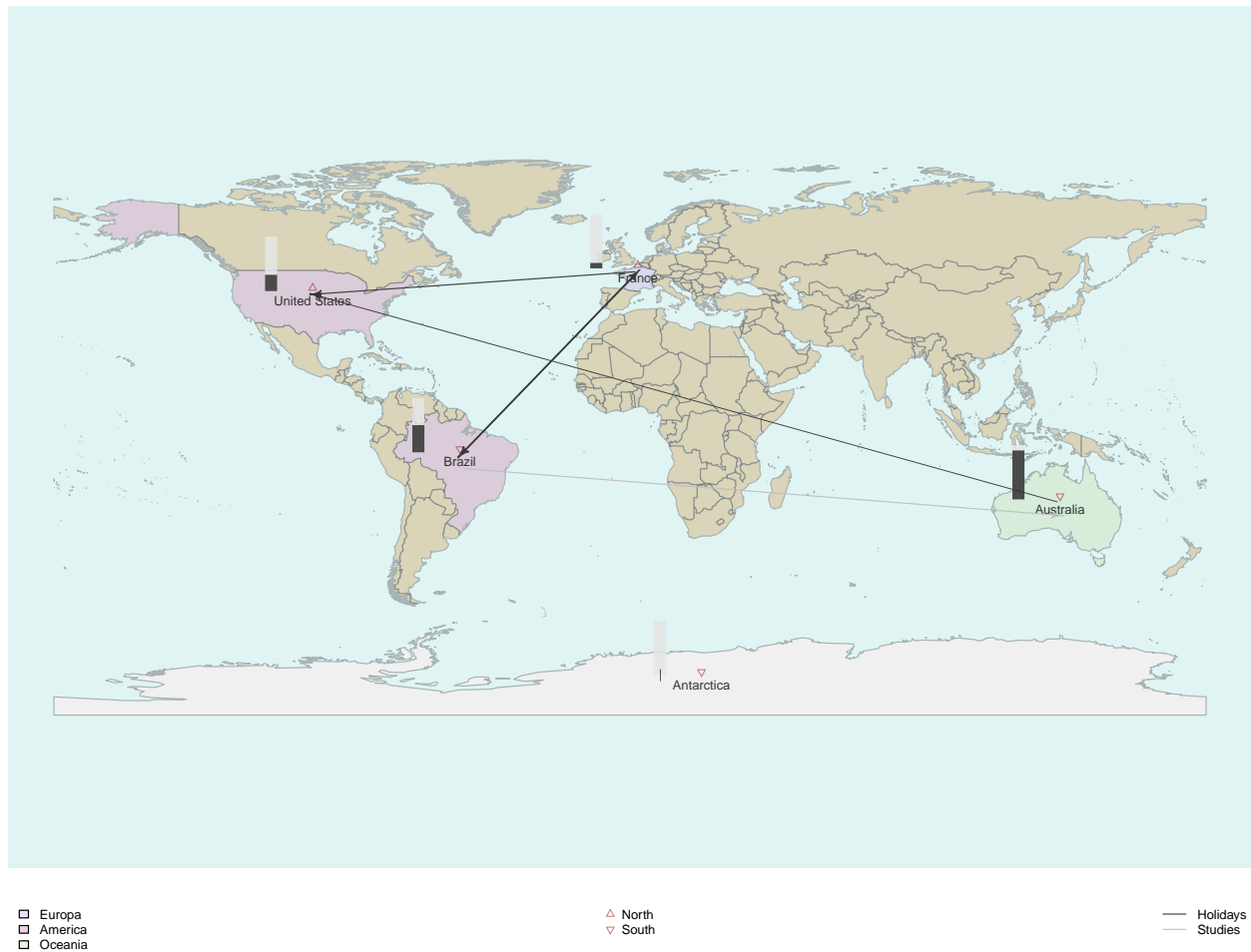


Figure 19:

Title and legend

To give a title to the graph we use:

```
graph.title.main(ex.bp) <- "Places visited by John, Elsa, Brian and Kate"
graph.title.sub(ex.bp) <- "For holidays and studies"
plot(ex.bp)
```

In order to make the legend a little bit bigger, network lines more visible and colored variable legend on 3 columns, for instance, we use:

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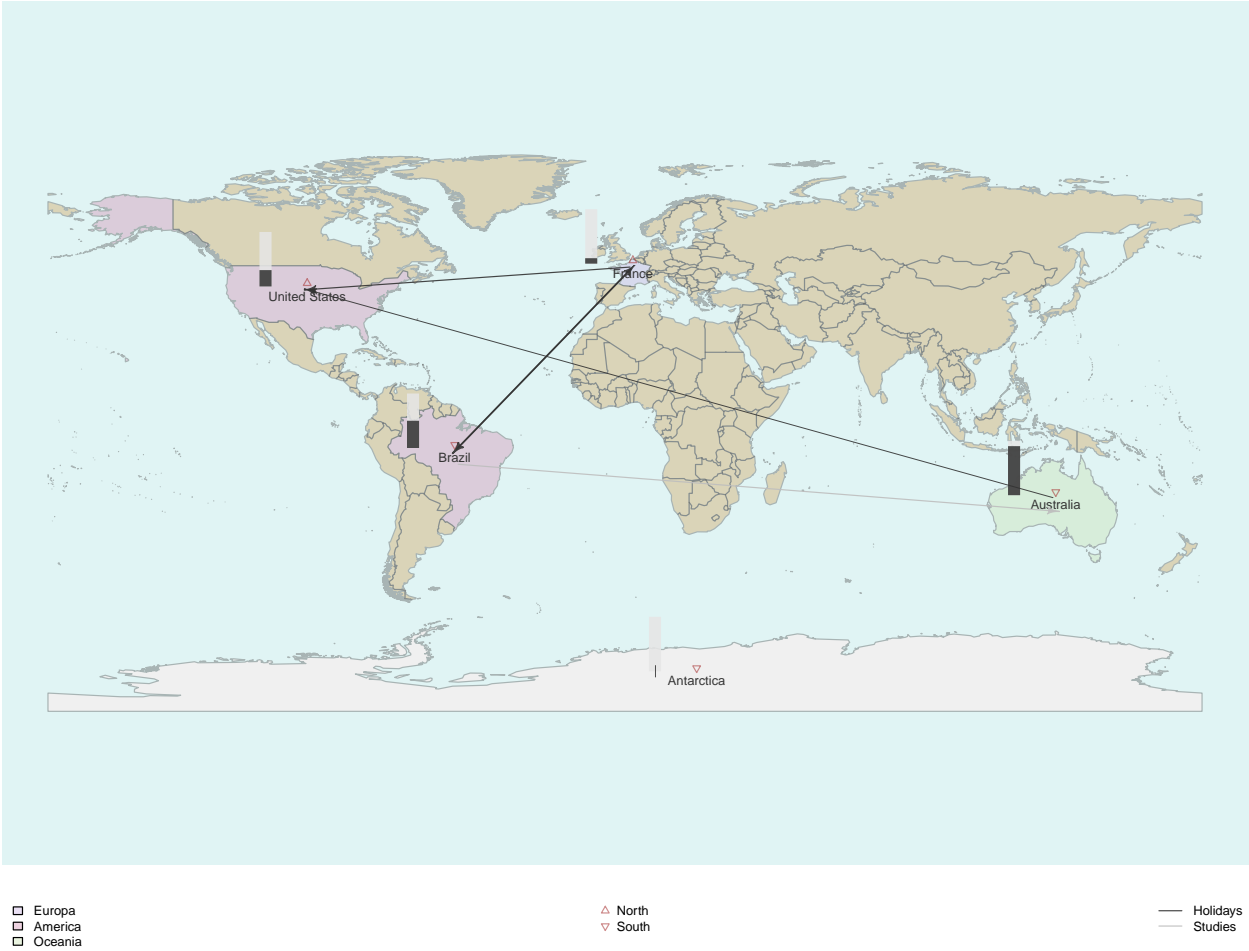


Figure 20:

```
graph.legend.cex(ex.bp) <- 1.2
graph.legend.line.width(ex.bp) <- 2
graph.legend.ncol(ex.bp) <- 3
plot(ex.bp)
```

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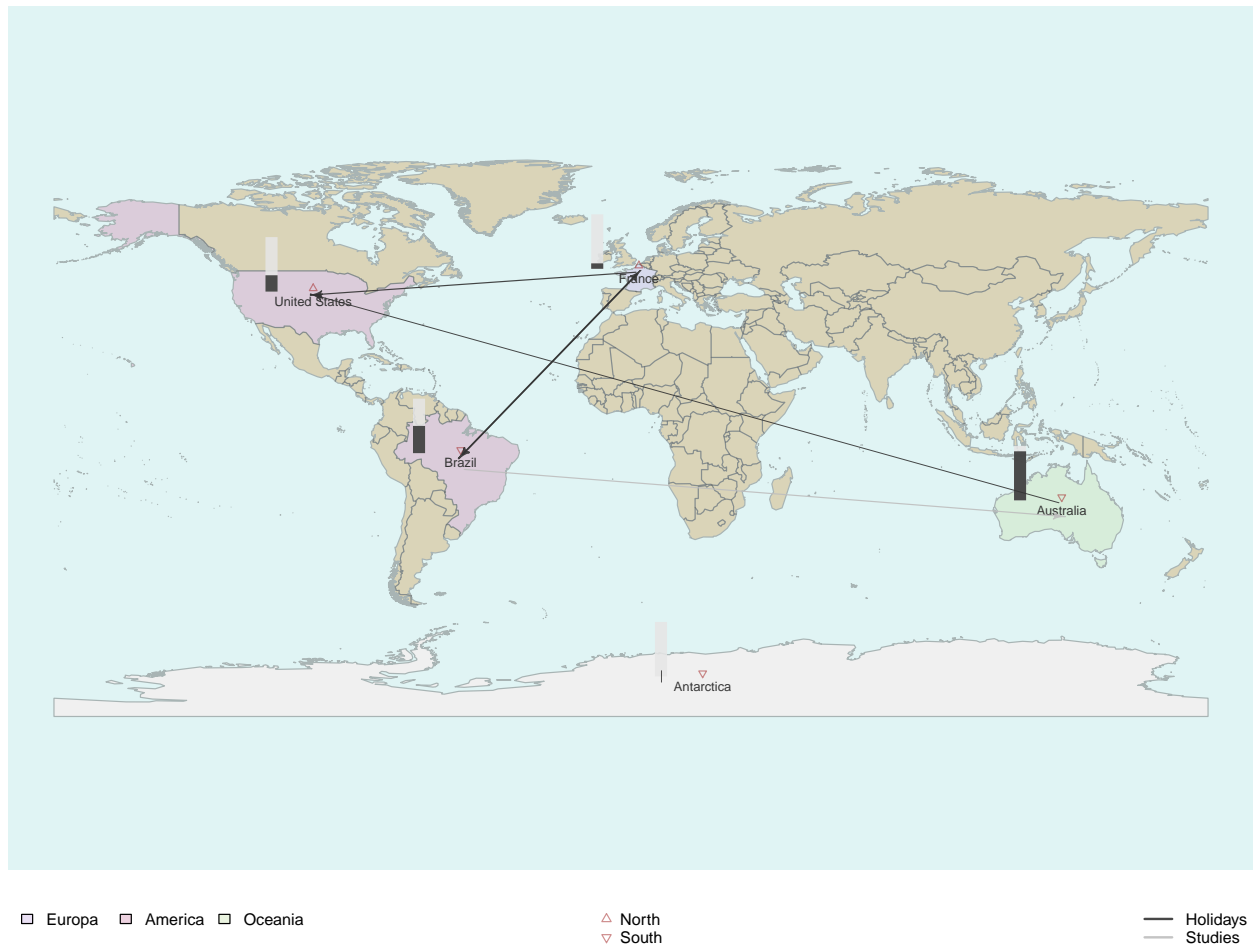


Figure 21:

To hide the legend, we should run:

```
graph.legend.print(ex.bp) <- FALSE
plot(ex.bp)
```

Black & white mode

The package provides a B&W mode that allows you (1) to easily preview how you graph will render when printing in B&W and (2) optimize readability. To enable the B&W mode run:

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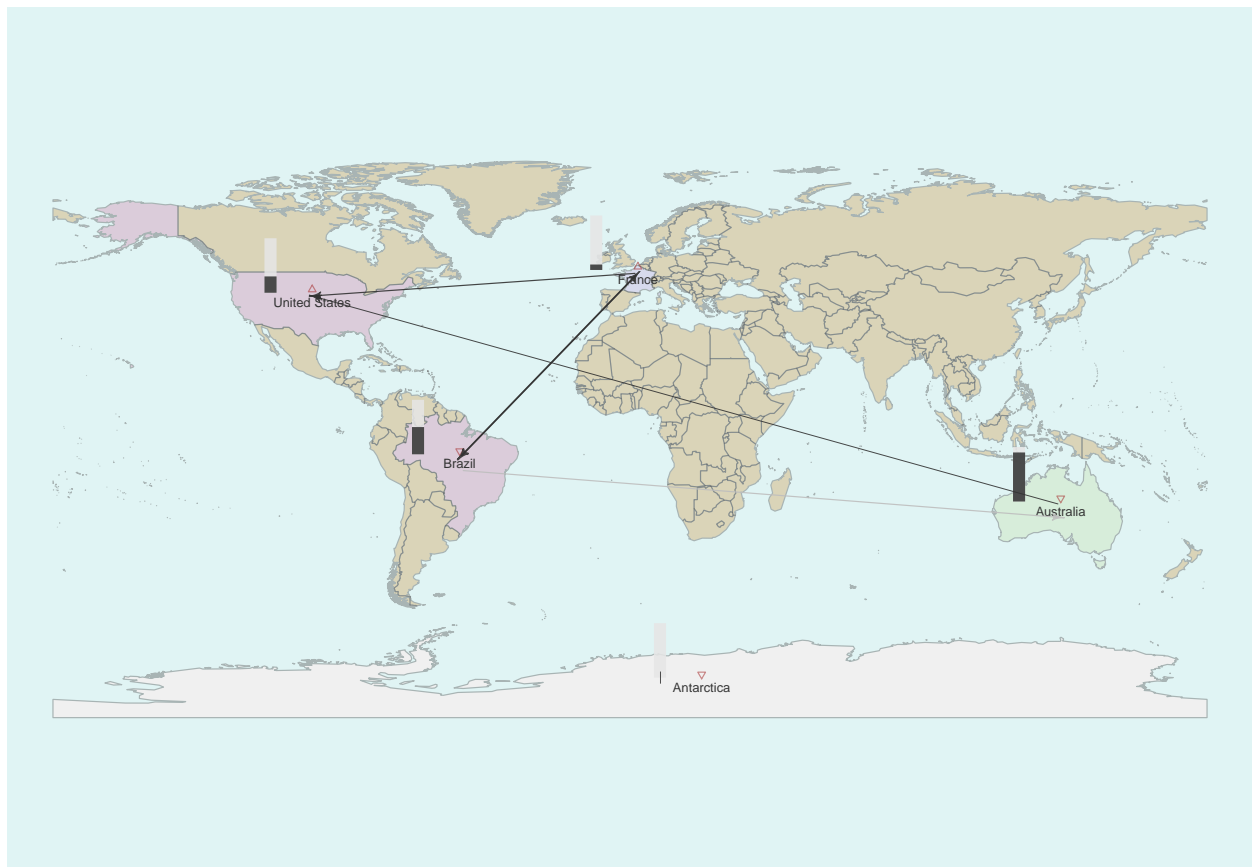


Figure 22:

```
graph.blackwhite.enable(ex.bp) <- TRUE
plot(ex.bp)
```

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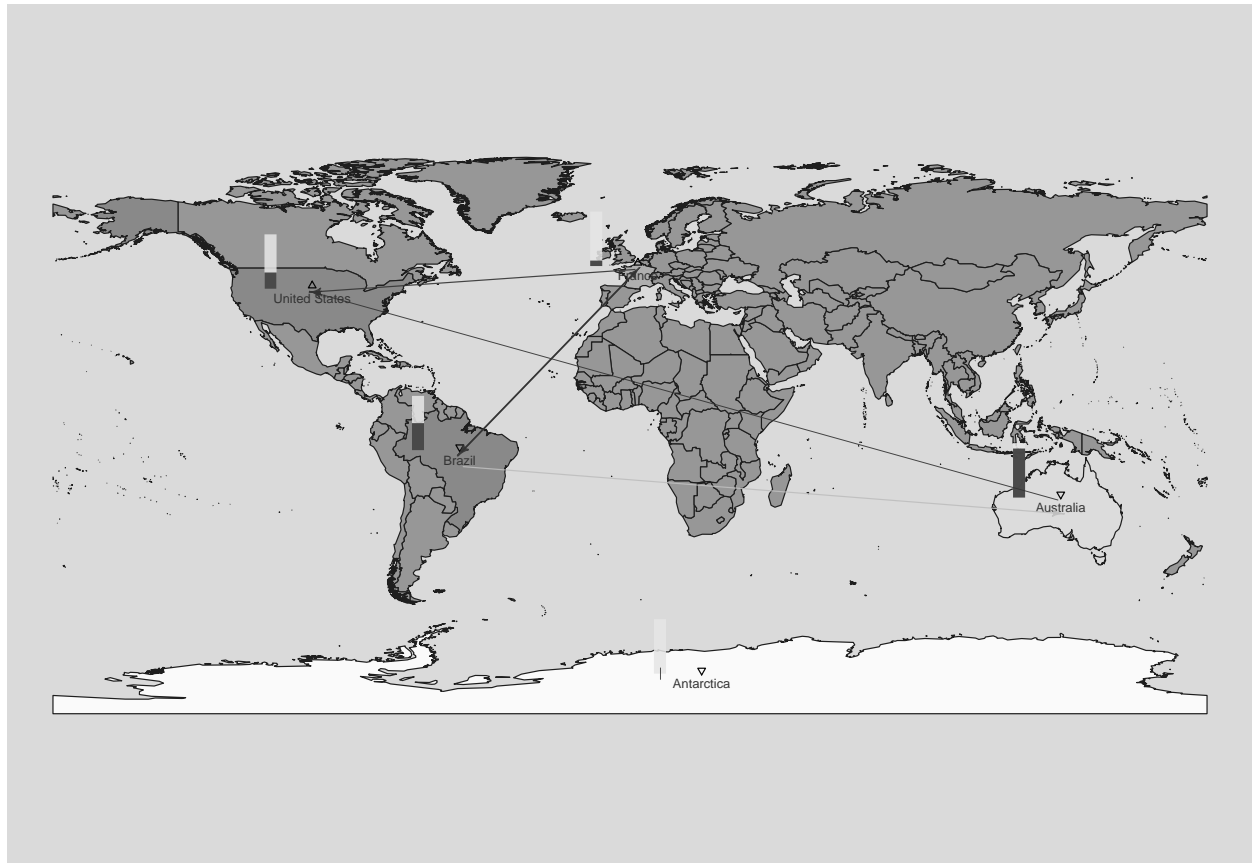


Figure 23:

The conversion from color to gray levels is performed using the luminosity method ($0.21 \text{ red} + 0.72 \text{ green} + 0.07 \text{ blue}$). Then, to enhance readability, graph contrast is increased by linearly scaling grey levels. You can customize the minimal and maximal value of grey with the `graph.blackwhite.min` and `graph.blackwhite.max` function and specifying a value in $[0;1]$ (0 for pure white and 1 for pure black). For instance:

```
graph.blackwhite.min(ex.bp) <- 0.0
graph.blackwhite.max(ex.bp) <- 0.6
plot(ex.bp)
```

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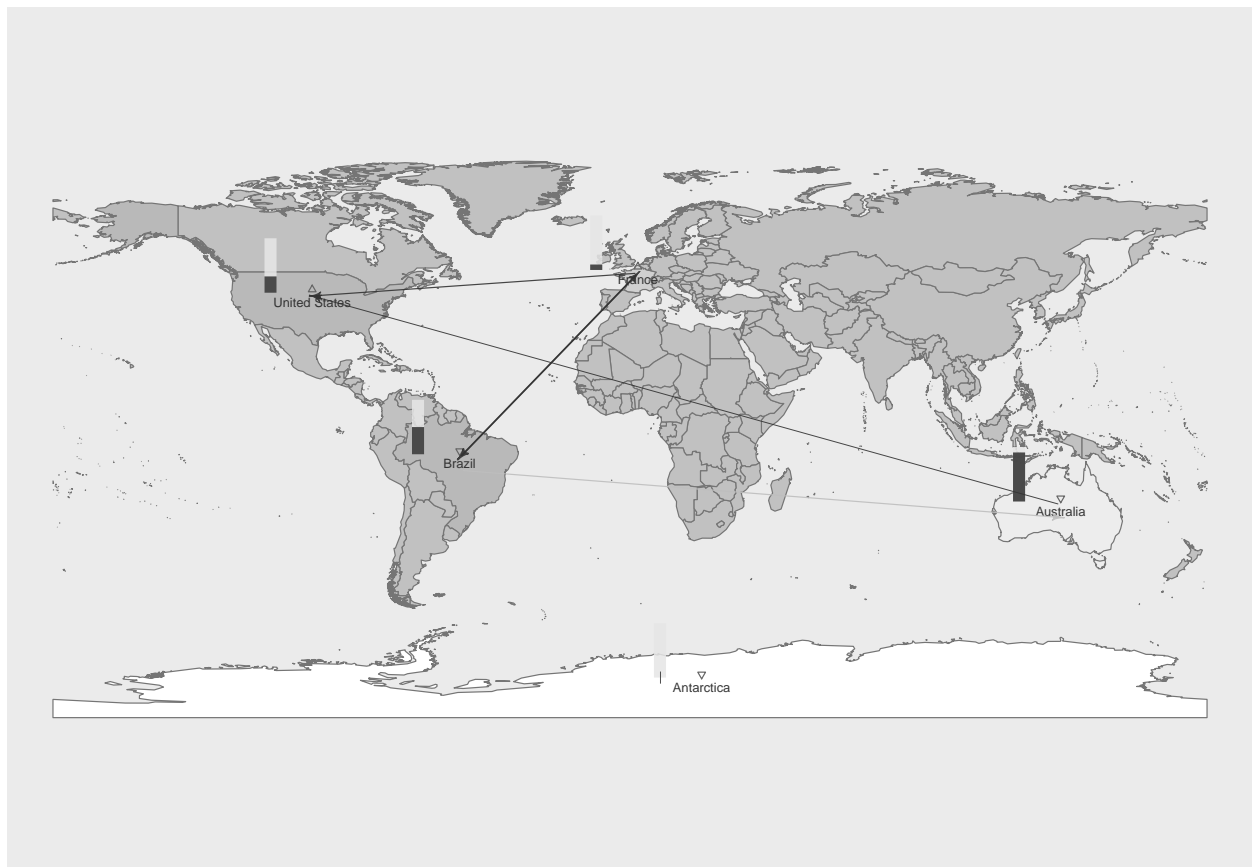


Figure 24: