

rworldmap vignette

Andy South*

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rworldmap is a package for visualising global data, concentrating on data referenced by country codes or gridded at half degree resolution.



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*Centre for Environment, Fisheries and Aquaculture Science (Cefas), Lowestoft, NR33 OHT, UK. southandy at gmail.com

1 Introduction

Package `rworldmap` is loaded by

```
> library(rworldmap)
```

This vignette shows a few examples of the main `rworldmap` functions to get you started. To access the full help system, type `?rworldmap` in the R console. The functions are designed to operate with few specified parameters in which case default values are used, but can also accept user input to allow flexibility e.g. in size, data categorisation, and colour schemes.

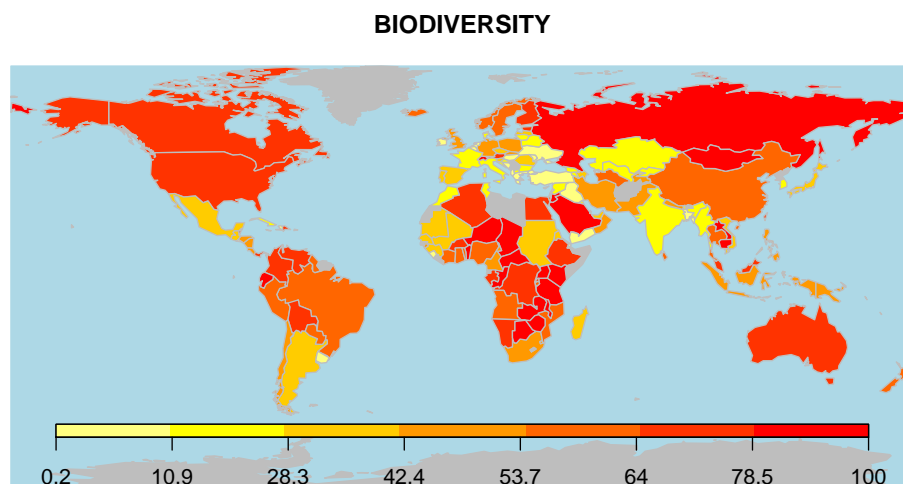


Figure 1: An example from `mapCountryData`

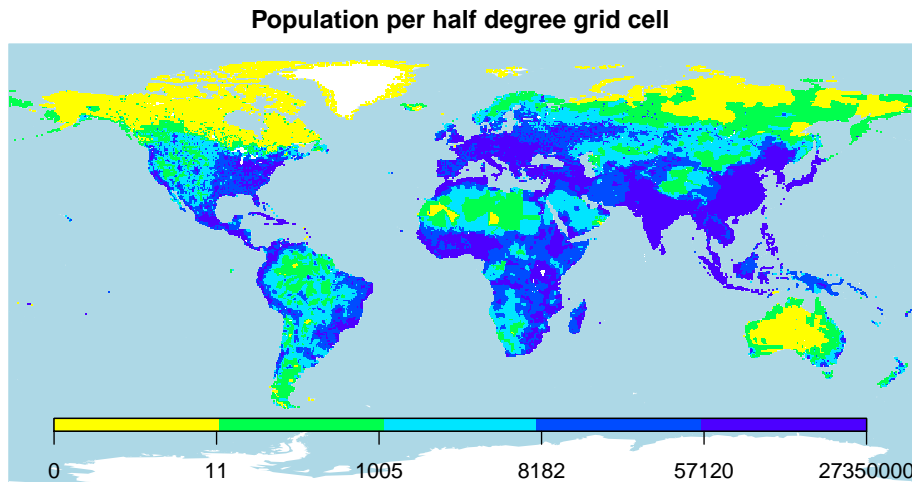


Figure 2: An example from `mapGriddedData`

2 Mapping your own country level data

To map your own data you will need it in columns with one row per country, one column containing country identifiers, and other columns containing your data.

The mapping process then involves 3 steps (or 2 if your data are already in an R dataframe).

1. read data into R
2. join data to a map (using `joinCountryData2Map()`)
3. display the map (using `mapCountryData()`)

There is an example dataset within the package that can be accessed using the `data` command, and the command below shows how to display a subset of the rows and columns.

```
> data(countryExData)
> countryExData[5:10,1:5]
```

	ISO3V10	Country	EPI_regions
5	ARM	Armenia	Middle East and North Africa
6	AUS	Australia	East Asia and the Pacific
7	AUT	Austria	Europe
8	AZE	Azerbaijan	Central and Eastern Europ
9	BDI	Burundi	Sub-Saharan Africa
10	BEL	Belgium	Europe

	GEO_subregion	Population2005
5	Eastern Europe	3016.3
6	Australia + New Zealand	20155.1
7	Western Europe	8189.4
8	Eastern Europe	8410.8

9	Eastern Africa	7547.5
10	Western Europe	10419.1

2.1 Reading data into R

To read in your own data from a space or comma delimited text file you will need to use : `read.csv(filename.csv)` or `read.txt(filename.txt)`, type `?read.table` from the R console to get help on this.

2.2 Joining data to a country map

To join the data to a map use `joinCountryData2Map`, and you will need to specify the name of column containing your country identifiers (`nameJoinColumn`) and the type of code used (`joinCode`) e.g. "ISO3" for ISO 3 letter codes or "UN" for numeric country codes. If you only have country names rather than codes use `joinCode="NAME"`, you can expect more mismatches because there is greater variation in what a single country may be named.

```
> data(countryExData)
> sPDF <- joinCountryData2Map( countryExData,
+                               joinCode = "ISO3",
+                               nameJoinColumn = "ISO3V10")
```

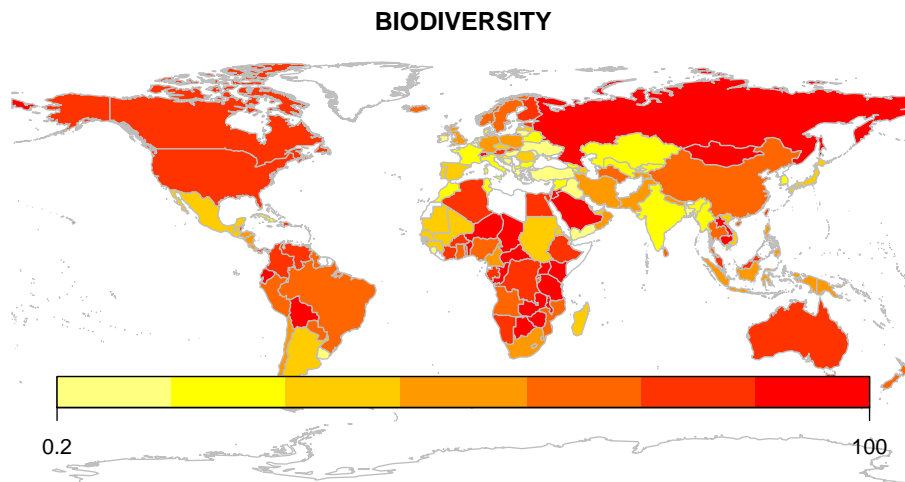
```
149 codes from your data successfully matched countries in the map
0 codes from your data failed to match with a country code in the map
97 codes from the map weren't represented in your data
```

You can see that a summary of how many countries are successfully joined is output to the console. You can specify `verbose=TRUE` to get a full list of countries. The object returned (named `sPDF` in this case) is of type `SpatialPolygonsDataFrame` from the package `sp`. This object is required for the next step, displaying the map.

2.3 Displaying a countries map

`mapCountryData` requires as a minimum a `SpatialPolygonsDataFrame` object and a specification of the name of the column containing the data to plot. The first line starting `par ...` below and in subsequent plots simply ensures the plot fills the available space on the page.

```
> par(mai=c(0,0,0.2,0),xaxs="i",yaxs="i")
> mapCountryData( sPDF, nameColumnToPlot="BIODIVERSITY" )
```



In this small map the default legend is rather large. This could be fixed by calling the `addMapLegend` function as in the code below.

```
> mapParams <- mapCountryData( sPDF, nameColumnToPlot="BIODIVERSITY", addLegend=F
> do.call( addMapLegend, c(mapParams, legendWidth=0.5, legendMar = 2))
```

Using `do.call` allows the output from `mapCountryData` to be used in `addMapLegend` to ensure the legend matches the map while also allowing easy modification of extra parameters such as `legendWidth`.

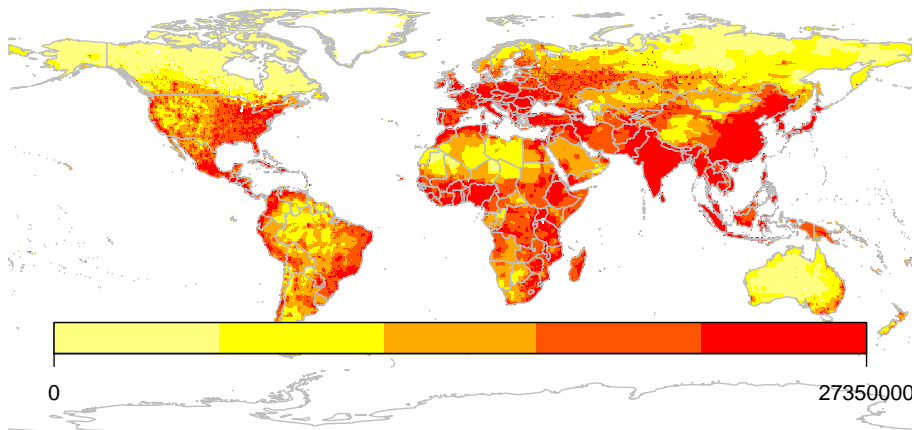
3 Mapping your own half degree gridded data

The `mapGriddedData` function can accept either

1. an object of type `SpatialGridDataFrame`, as defined in the package `sp`
2. the name of an ESRI `gridAscii` file as a character string
3. a 2D R matrix or array (rows by columns)

`rworldmap` contains an example `SpatialGridDataFrame` that can be accessed and printed as shown in the code below.

```
> par(mai=c(0,0,0.2,0),xaxs="i",yaxs="i")
> data(gridExData)
> mapGriddedData(gridExData)
```

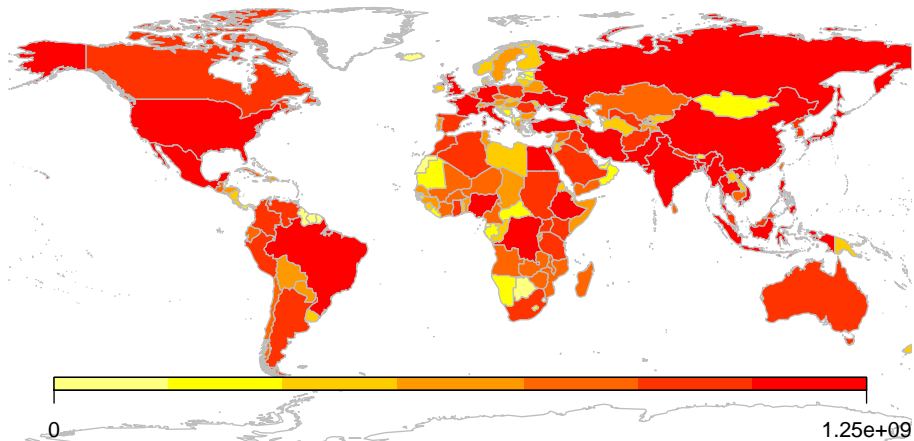


4 Aggregating half degree gridded data to a country level

`mapHalfDegreeGridToCountries()` takes a gridded input file, and aggregates, to a country level and plots the map, it accepts most of the same arguments as `mapCountryData()`. In the example below the trick from above of modifying the legend using `addMapLegend()` is repeated.

```
> par(mai=c(0,0,0.2,0),xaxs="i",yaxs="i")
> mapParams <- mapHalfDegreeGridToCountries(gridExData, addLegend=FALSE)
> do.call( addMapLegend, c(mapParams, legendWidth=0.5, legendMar = 2))
```

sum_pa2000.asc



5 Aggregating country level data to global regions

Country level data can be aggregated to global regions specified by `regionType` in `country2Region` which outputs as text, and `mapByRegion` which produces a map plot. The regional classifications available include SRES, GEO3, Stern and GBD.

```
> #Using country2Region to calculate mean Environmental Health index in Stern reg
> sternEnvHealth <- country2Region(inFile=countryExData,
```

```

+           nameDataColumn="ENVHEALTH",
+           joinCode="ISO3",
+           nameJoinColumn="ISO3V10",
+           regionType="Stern",
+           FUN="mean"
+         )
> print(sternEnvHealth)

```

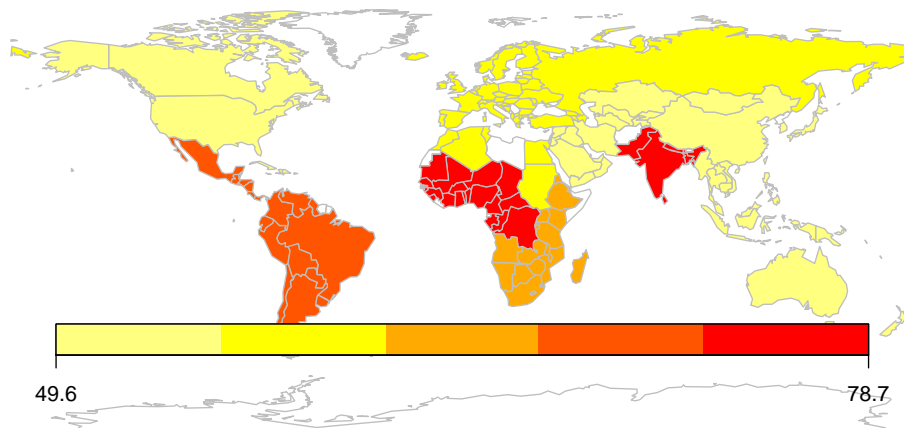
	meanENVHEALTHbyStern
Australasia	78.86000
Caribbean	82.18000
Central America	82.78750
Central Asia	77.24000
East Asia	75.52308
Europe	95.19762
North Africa	77.38000
North America	98.70000
South America	83.62727
South Asia	61.96000
South+E Africa	49.06316
West Africa	36.99474
West Asia	82.78000

```

> par(mai=c(0,0,0.2,0),xaxs="i",yaxs="i")
> mapByRegion(countryExData,
+           nameDataColumn="CLIMATE",
+           joinCode="ISO3",
+           nameJoinColumn="ISO3V10",
+           regionType="Stern",
+           FUN="mean"
+         )

```

mean CLIMATE by Stern regions



6 Map display options common across the plotting methods

The following arguments can be specified to alter the appearance of your plots.

- **catMethod** method for categorisation of data "pretty", "fixedWidth", "diverging", "logfixedWidth", "quantiles", "categorical", or a numeric vector defining breaks.
- **numCats** number of categories to classify the data into, may be modified if that exact number is not possible for the chosen catMethod.
- **colourPalette** a string describing the colour palette to use, choice of :
 1. "palette" for the current palette
 2. a vector of valid colours, e.g. c("red", "white", "blue") or output from RColourBrewer
 3. one of "heat", "diverging", "white2Black", "black2White", "topo", "rainbow", "terrain", "negpos8", "negpos9"
- **addLegend** set to TRUE for a default legend, if set to FALSE the function addMapLegend() or addMapLegendBoxes() can be used to create a more flexible legend.
- **mapRegion** a region to zoom in on, can be set to a country name from getMap()\$NAME or one of "eurasia", "africa", "latin america", "uk", "oceania", "asia"

7 Example maps with settings modified

```
> par(mai=c(0,0,0.2,0),xaxs="i",yaxs="i")
> #joining the data to a map
> sPDF <- joinCountryData2Map(countryExData,
+                             joinCode = "ISO3",
+                             nameJoinColumn = "ISO3V10",
+                             projection = "none",
+                             )
> #creating a user defined colour palette
> op <- palette(c('green','yellow','orange','red'))
> #find quartile breaks
> cutVector <- quantile(sPDF@data[["BIODIVERSITY"]],na.rm=TRUE)
> #classify the data to a factor
> sPDF@data[["BIOcategories"]] <- cut(sPDF@data[["BIODIVERSITY"]], cutVector, include.lowest=TRUE)
> #rename the categories
> levels(sPDF@data[["BIOcategories"]]) <- c('low', 'med', 'high', 'vhigh')
> #mapping
> mapCountryData(sPDF, nameColumnToPlot='BIOcategories', catMethod='categorical',
+ mapTitle='Biodiversity categories',colourPalette='palette', oceanCol='lightblue',
+ missingCountryCol='white')
```

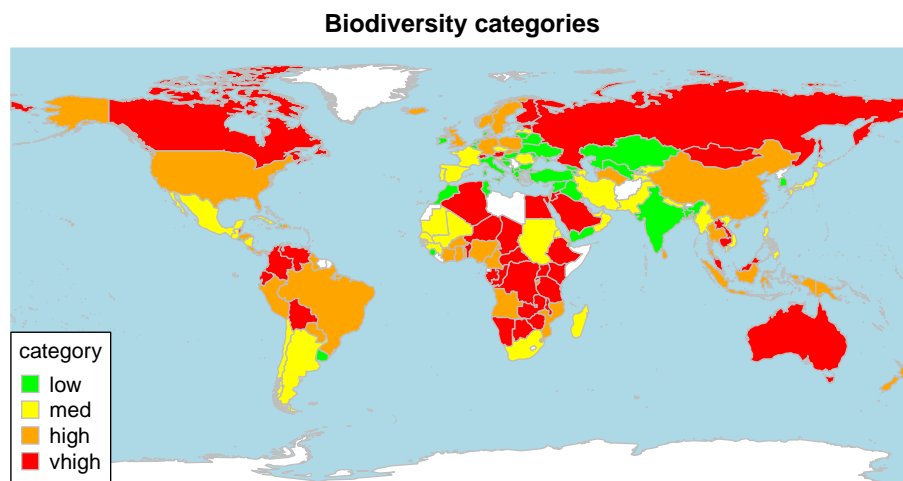
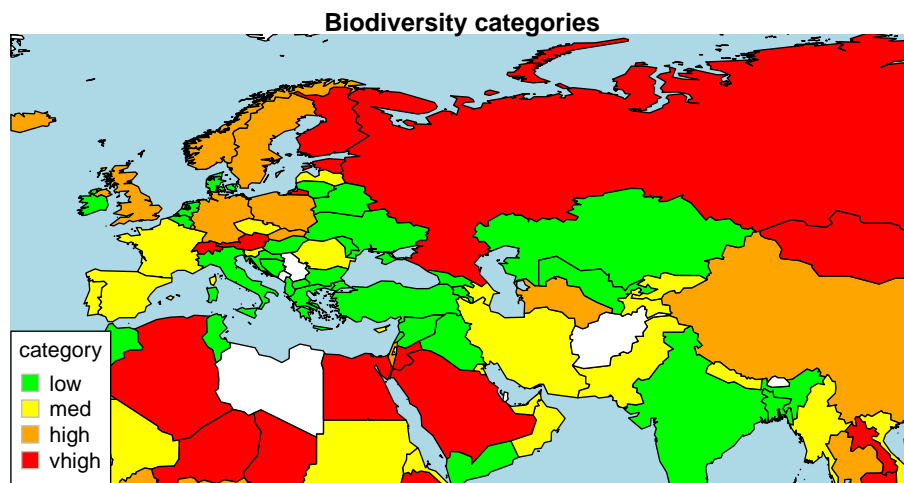


Figure 3: An example of a categorical map produced from `mapCountryData`

This demonstrates how continuous data can be put into categories outside of the `rworldmap` functions and how a user defined colour palette can be used. Because the `catMethod="categorical"` was used a legend with separate boxes rather than a colour bar is added.

You can zoom in on a map by specifying `mapRegion="Eurasia"` (or by specifying `xlim` and `ylim`) and the country outlines can be changed by `borderCol="black"`.

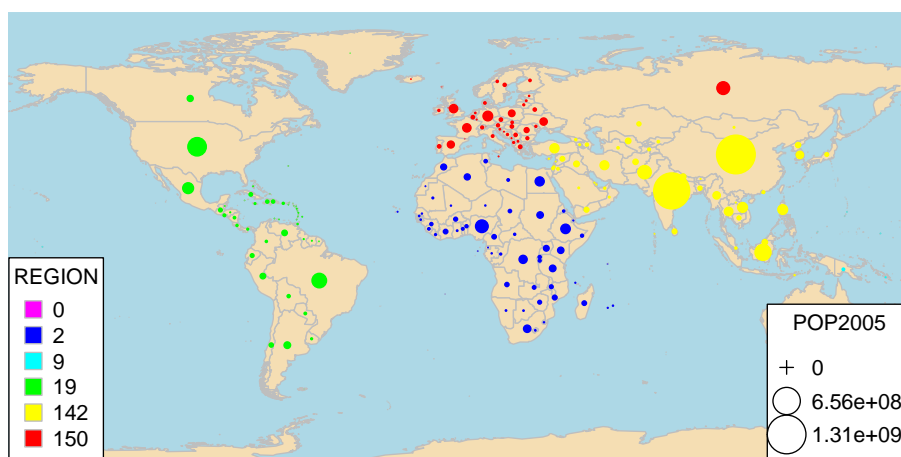
```
> par(mai=c(0,0,0.2,0),xaxs="i",yaxs="i")
> mapCountryData(sPDF, nameColumnToPlot='BIOcategories', catMethod='categorical',
+ mapTitle='Biodiversity categories',colourPalette='palette', oceanCol='lightblue',
+ missingCountryCol='white', mapRegion='Eurasia', borderCol='black')
> ## At end of plotting, reset palette to previous settings:
> palette(op)
>
```



8 Bubble plots

The `mapBubbles` function allows flexible creation of bubble plots on global maps. You can specify data columns that will determine the sizing and colouring of the bubbles (using `nameZsize` and `nameZColour`). The function also accepts other `spatialDataFrame` objects or data frames as long as they contain columns specifying the x and y coordinates. The interactive function `identifyCountries` allows the user to click on bubbles and the country name and optionally an attribute variable will be printed on the map.

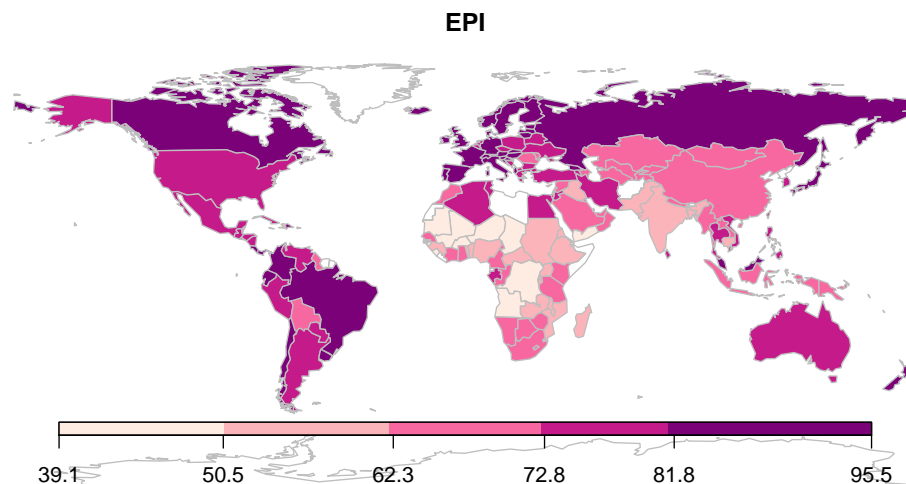
```
> par(mai=c(0,0,0.2,0),xaxs="i",yaxs="i")
> mapBubbles(dF=getMap(), nameZSize="POP2005", nameZColour="REGION",colourPalette
>
>
```



9 Combining rworldmap with other packages classInt and RColorBrewer

Whilst rworldmap sets many defaults internally there is also an option to use other packages to have greater flexibility. In this example the package classInt is used to create the classification and RColorBrewer to specify the colours. The following page demonstrates how multiple maps can be generated in the same figure and shows a selection of different RColorBrewer palettes.

```
> par(mai=c(0,0,0.2,0),xaxs="i",yaxs="i")
> library(classInt)
> library(RColorBrewer)
> #getting example data and joining to a map
> data("countryExData",envir=environment(),package="rworldmap")
> sPDF <- joinCountryData2Map(countryExData,joinCode = "ISO3",nameJoinColumn = "1")
> #getting class intervals using a 'jenks' classification in classInt package
> classInt <- classIntervals( sPDF[["EPI"]], n=5, style="jenks")
> catMethod = classInt[["brks"]]
> #getting a colour scheme from the RColorBrewer package
> colourPalette <- brewer.pal(5,'RdPu')
> #calling mapCountryData with the parameters from classInt and RColorBrewer
> mapParams <- mapCountryData(sPDF,nameColumnToPlot="EPI", addLegend=FALSE, catMe
> do.call(addMapLegend, c(mapParams,legendLabels="all",legendWidth=0.5,legendInte
```



```

> #uses sPDF from the previous chunk
>
> op <- par(fin=c(7,9),mfcol=c(5,2),mai=c(0,0,0.2,0),xaxs="i",yaxs="i")
> brewerList <- c("Greens","Greys","Oranges","OrRd","PuBuGn","Purples","YlGn","YlGnBu","YlOrBr","YlOrRd")
> for(i in 1:10)
+   {
+     #getting a colour scheme from the RColorBrewer package
+     colourPalette <- brewer.pal(7,brewerList[i])
+
+     #calling mapCountryData with the parameters from RColorBrewer
+     mapParams <- mapCountryData(sPDF,nameColumnToPlot="CLIMATE", addLegend=FALSE, colourPalette=colourPalette)
+     do.call(addMapLegend, c(mapParams,horizontal=FALSE,legendLabels="none",legendWidth=0.7))
+   }
> par(op)

```

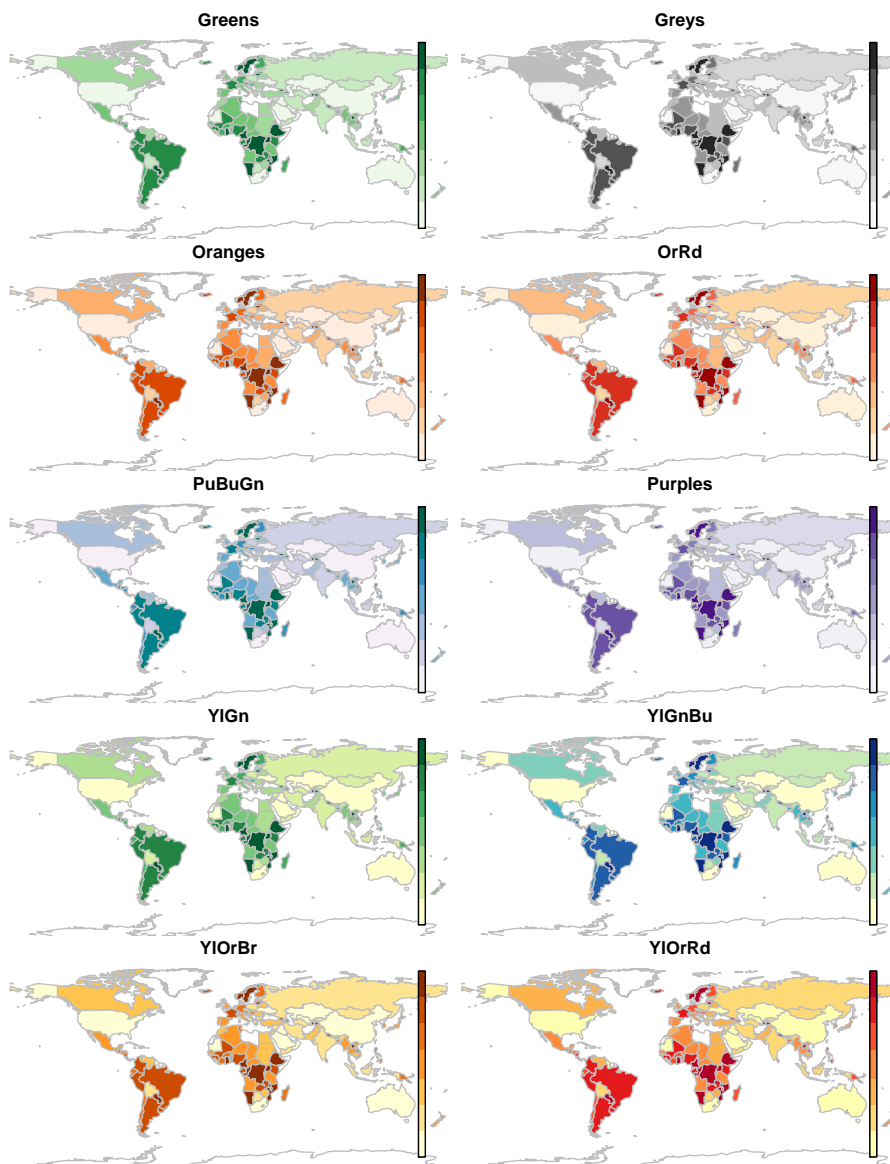


Figure 4: Different RColorBrewer palettes applied to the same data