rworldmap vignette

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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.







Funded by:

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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.

BIODIVERSITY

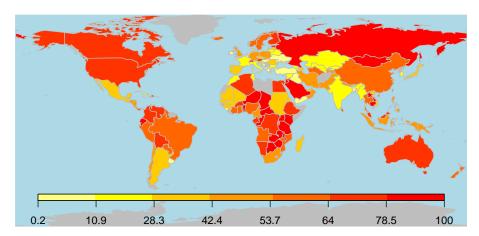


Figure 1: An example from mapCountryData

Population per half degree grid cell

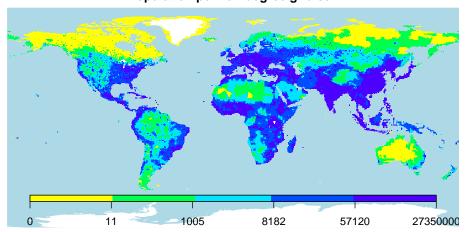


Figure 2: An example from mapGriddedData

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	${\tt Country}$			EPI_:	regions
5	ARM	Armenia	Middle	East aı	nd North	Africa
6	AUS	Australia	East	Asia a	and the	Pacific
7	AUT	Austria				Europe
8	AZE	Azerbaijan	Cent	ral and	d Easter	n Europ
9	BDI	Burundi		Sub-	-Saharan	Africa
10	BEL	Belgium				Europe
		GEO_subre	gion Po	pulatio	on2005	
5		Eastern Eu	rope	3	3016.3	
6	Australi	ia + New Zea	land	20	0155.1	
7		Western Eu	rope	8	3189.4	
8		Eastern Eu	rope	8	3410.8	

9	Eastern Af	rica	7547.5
10	Western Eu	rope	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 <code>joinCountryData2Map</code>, 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.

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

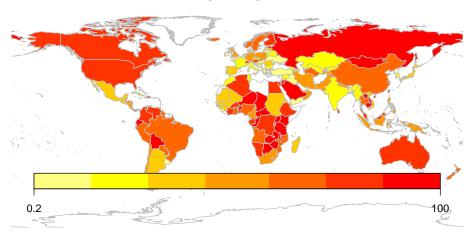
You can see that a summary of how many countries are successfully joined is output to the console. You can specify verbose=TRUE to get a full list of countries. The object returned (named sPDF in this case) is of type SpatialPolygonsDataFrame from the package sp. This object is required for the next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

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=FALSE )
> 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

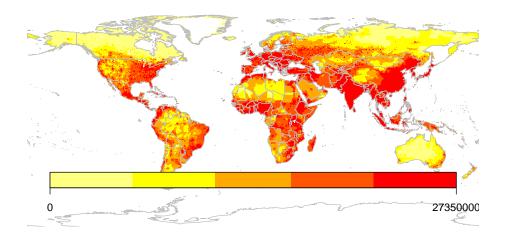
The mapGriddedData function can accept either

- 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")
```

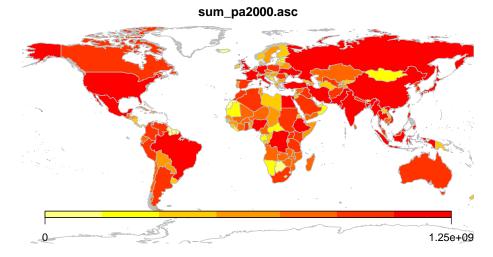
- > data(gridExData)
- > mapGriddedData(gridExData)



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 map-CountryData(). 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))



5 Aggregating country level data to global regions

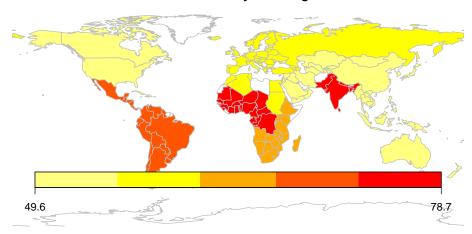
Country level data can be aggregated to global regions specified by regionType in country2Region which outputs as text, and mapByRegion which produces a map plot. The regional classifications available include SRES, GEO3, Stern and GBD.

- > #Using country2Region to calculate mean ENVHEALTH in Stern regions.
- > sternEnvHealth <- country2Region(inFile=countryExData

meanENVHEALTHbyStern

J ~ ~ ~
78.86000
82.18000
82.78750
77.24000
75.52308
95.19762
77.38000
98.70000
83.62727
61.96000
49.06316
36.99474
82.78000

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 RColour-Brewer
 - 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

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.

```
> 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'))</pre>
> #find quartile breaks
> cutVector <- quantile(sPDF@data[["BIODIVERSITY"]],na.rm=TRUE)</pre>
> #classify the data to a factor
> sPDF@data[["BIOcategories"]] <- cut(sPDF@data[["BIODIVERSITY"]]</pre>
                                       , cutVector
                                       , include.lowest=TRUE )
> #rename the categories
> levels(sPDF@data[["BIOcategories"]]) <- c('low', 'med', 'high', 'vhigh')</pre>
> #mapping
> mapCountryData( sPDF
                 , nameColumnToPlot='BIOcategories'
                 , catMethod='categorical'
                 , mapTitle='Biodiversity categories'
                 , colourPalette='palette'
                 , oceanCol='lightblue'
                  missingCountryCol='white'
```

Biodiversity categories

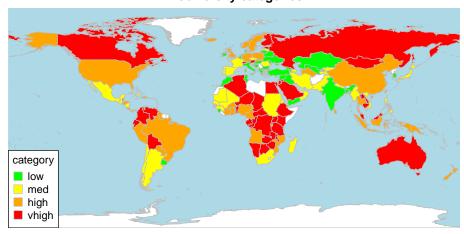
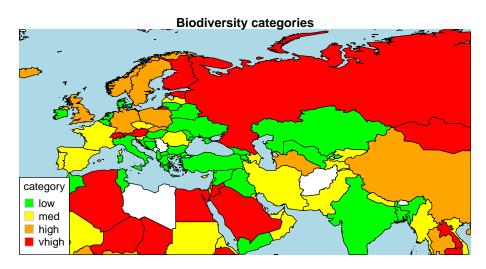


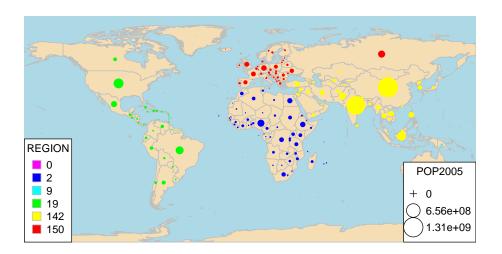
Figure 3: An example of a categorical map produced from mapCountryData

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".



8 Bubble plots

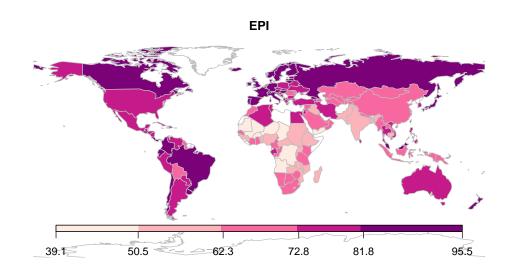
The mapBubbles function allows flexible creation of bubble plots on global maps. You can specify data columns that will determine the sizing and colouring of the bubbles (using nameZsize and nameZColour). The function also accepts other spatialDataFrame objects or data frames as long as they contain 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.



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</pre>
+
                              , joinCode = "ISO3"
                              , nameJoinColumn = "ISO3V10"
                              , mapResolution='coarse'
+
> #getting class intervals using a 'jenks' classification in classInt package
> classInt <- classIntervals( sPDF[["EPI"]], n=5, style="jenks")</pre>
> catMethod = classInt[["brks"]]
> #getting a colour scheme from the RColorBrewer package
> colourPalette <- brewer.pal(5,'RdPu')</pre>
> #calling mapCountryData with the parameters from classInt and RColorBrewer
> mapParams <- mapCountryData( sPDF</pre>
+
                              , nameColumnToPlot="EPI"
+
                              , addLegend=FALSE
+
                              , catMethod = catMethod
                              , colourPalette = colourPalette )
> do.call( addMapLegend
         , c( mapParams
+
            , legendLabels="all"
            , legendWidth=0.5
            , legendIntervals="data"
            , legendMar = 2 ) )
```



```
> #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"</pre>
                   ,"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])</pre>
      #calling mapCountryData with the parameters from RColorBrewer
      mapParams <- mapCountryData( sPDF</pre>
                                     , nameColumnToPlot="CLIMATE"
                                     , addLegend=FALSE
                                     , colourPalette=colourPalette
                                     , mapTitle=brewerList[i] )
      do.call( addMapLegend
              , c(\texttt{mapParams}, \texttt{horizontal=FALSE}, \texttt{legendLabels="none"}, \texttt{legendWidth=0.7}))
> par(op)
```

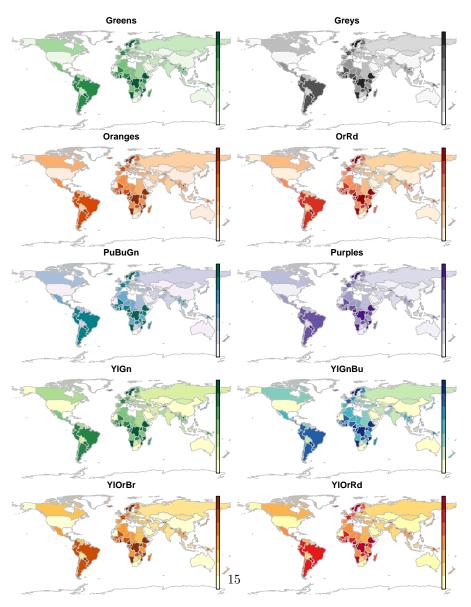


Figure 4: Different RColorBrewer palettes applied to the same data