Package ‘treemap’

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Description

A treemap is a space-filling visualization of hierarchical structures. This package offers great flexibility to draw treemaps.

Details

The main function is `treemap`. See also `itreemap` for a graphical user interface to create treemaps. By default Tree Colors are used, which are colors from the HCL color space. Use `treecolors` to experiment with the parameter settings.

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Description

Fictitious (aggregated) business statistics data. The index variables (NACE1 to NACE4) are derived from the Statistical Classification of Economic Activities in the European Community (NACE). The variables `turnover(.prev)` and `employees(.prev)` have values for NACE codes in the business economy domain only.

References

Statistical Classification of Economic Activities in the European Community (NACE) Eurostat - Structural business statistics (SBS)
Description

Gross national income (per capita) in dollars and population totals per country in 2014.

Details

The GNI numbers from the World Bank are based on the Atlas. The population data are taken from Natural Earth Data.

References

The World Bank - GNI per capita ranking Natural Earth Data

itreemap

Interactive user interface for treemap

Description

This function is an interactive user interface for creating treemaps. Interaction is provided for the four main input arguments of (treemap) besides the data.frame itself, namely index, vSize, vColor and type. Zooming in and out is possible. Command line outputs are generated in the console.

Usage

itreemap(
  dtf = NULL,
  index = NULL,
  vSize = NULL,
  vColor = NULL,
  type = NULL,
  height = 700,
  command.line.output = TRUE
)

Arguments

dtf a data.frame (treemap) If not provided, then the first data.frame in the global workspace is loaded.
index index variables (up to four). See treemap.
vSize name of the variable that determine the rectangle sizes.
vColor name of the variable that determine the rectangle colors. See treemap.
random.hierarchical.data

Create random hierarchical data

Description

This function generates random hierarchical data. Experimental.

Usage

```r
random.hierarchical.data(
  n = NULL,
  method = "random",
  number.children = 3,
  children.root = 4,
  depth = 3,
  nodes.per.layer = NULL,
  labels = c("LETTERS", "numbers", "letters"),
  labels.prefix = NULL,
  sep = ".",
  colnames = c(paste("index", 1:depth, sep = ""), "x"),
  value.generator = rlnorm,
  value.generator.args = NULL
)
```
Arguments

**n**  
number of leaf nodes. This is a shortcut argument. If specified, the method is set to "random.arcs" with a nodes.per.layer such that the average number of children per layer is as constant as possible.

**method**  
one of
- "random": Random tree where for each node, the number of children, is determined by a random poisson generator with \( \lambda = \text{number.children} \), until the maximum depth specified by depth is reached. The number of children of the root node is set to children.root.
- "random.arcs": Random tree where the exact number of nodes in each layer must be specified by nodes.per.layer. The arcs between the layers are random, with the restriction that each node is connected.
- "full.tree": Each node has exactly number.children children.

**number.children**  
the number of children. For method="random" this is the average number of children and for method="full.tree", it is the exact number of children. In the latter case, it can also be a vector that specifies the number of children for each layer.

**children.root**  
number of children of the root node. For method="random" only.

**depth**  
deepth of the tree. Note that for method="random", this depth may not be reached.

**nodes.per.layer**  
extact number of nodes per layer, that is needed for method="random.arcs"

**labels**  
one of "letters", "LETTERS", "numbers", "numbers1", "numbers0", "hex", "bits". The label set for "numbers1" is 1:9, and for "numbers0" it is 0:9. "numbers" is equal to "numbers0", except that starts from 1.

**labels.prefix**  
vector of label prefixes, one for each layer

**sep**  
separator character

**colnames**  
names of the columns. The first depth columns are the index columns (from highest to lowest hierarchical layer), and the last column is stored with random values

**value.generator**  
function that determine the random values for the leaf nodes

**value.generator.args**  
list of arguments passed to value.generator

Examples

```r
d <- random.hierarchical.data(200)
treemap(d, index=names(d)[1:(ncol(d)-1)], vSize="x")

d <- random.hierarchical.data(number.children=5)
treemap(d, index=names(d)[1:(ncol(d)-1)], vSize="x")

d <- random.hierarchical.data(method="full.tree", number.children=3, value.generator=runif)
treemap(d, index=names(d)[1:(ncol(d)-1)], vSize="x")
```
tmPlot

Create a treemap (deprecated)

Description
This function is migrated to `treemap`.

Usage
```
tmPlot(...)  
```

Arguments
```
... passed on to treemap
```

treecolors

Interactive tool to experiment with Tree Colors

Description
Tree Colors are color palettes for tree data structures. They are used in `treemap` by default (`type="index"`). With this tool, users can experiment with the parameters (in `treemap` stored in `palette.HCL.options`). Tree Colors can directly be obtained by `treepalette` with method="HCL".

Usage
```
treecolors(height = 700)
```

Arguments
```
height height of the plotted treemap in pixels. Tip: decrease this number if the treemap doesn’t fit conveniently.
```

Examples
```
## Not run:  
treecolors()

## End(Not run)
```
Create a tree graph

Description

This function draws a tree graph. By default, a radial layout is used.

Usage

treegraph(
  dtf,
  index = names(dtf),
  directed = FALSE,
  palette.HCL.options,
  show.labels = FALSE,
  rootlabel = "",
  vertex.layout = "reingold.tilford",
  vertex.layout.params,
  truncate.labels = NULL,
  vertex.size = 3,
  vertex.label.dist = 0.3,
  vertex.label.cex = 0.8,
  vertex.label.family = "sans",
  vertex.label.color = "black",
  mai = c(0, 0, 0, 0),
  ...
)

Arguments

dtf a data.frame or data.table. Required.
index the index variables of dtf (see treemap)
directed logical that determines whether the graph is directed (TRUE) or undirected (FALSE)
palette.HCL.options list of advanced options to obtain Tree Colors from the HCL space (when palette="HCL"). This list contains:
  hue_start: number between 0 and 360 that determines the starting hue value (default: 30)
  hue_end: number between hue_start and hue_start + 360 that determines the ending hue value (default: 390)
  hue_perm: boolean that determines whether the colors are permuted such that adjacent levels get more distinguishable colors. If FALSE, then the colors are equally distributed from hue_start to hue_end (default: TRUE)
  hue_rev: boolean that determines whether the colors of even-numbered branched are reversed (to increase discrimination among branches)
hue_fraction: number between 0 and 1 that determines the fraction of the hue circle that is used for recursive color picking: if 1 then the full hue circle is used, which means that the hue of the colors of lower-level nodes are spread maximally. If 0, then the hue of the colors of lower-level nodes are identical of the hue of their parents. (default: .5)

chroma: chroma value of colors of the first-level nodes, that are determined by the first index variable (default: 60)

luminance: luminance value of colors of the first-level nodes, i.e. determined by the first index variable (default: 70)

chroma_slope: slope value for chroma of the non-first-level nodes. The chroma values for the second-level nodes are chroma+chroma_slope, for the third-level nodes chroma+2*chroma_slope, etc. (default: 5)

luminance_slope: slope value for luminance of the non-first-level nodes (default: -10)

For "depth" and "categorical" types, only the first two items are used. Use treecolors to experiment with these parameters.

show.labels show the labels

rootlabel name of the root label

vertex.layout layout algorithm name. See layout for options. The name corresponds to the layout function name after the period symbol, e.g. "auto", "random", etc. The default is "reingold.tilford" with a circular layout.

vertex.layout.params list of arguments passed to vertex.layout

truncate.labels number of characters at which the levels are truncated. Either a single value for all index variables, or a vector of values for each index variable

vertex.size vertex.size (see igraph.plotting)

vertex.label.dist vertex.label.dist (see igraph.plotting)

vertex.label.cex vertex.label.cex (see igraph.plotting)

vertex.label.family vertex.label.family (see igraph.plotting)

vertex.label.color vertex.label.color (see igraph.plotting)

mai margins see par

... arguments passed to plot.igraph

Value

(invisible) igraph object
Examples

```r
data(business)
treegraph(business, index=c("NACE1", "NACE2", "NACE3", "NACE4"), show.labels=FALSE)
treegraph(business[business$NACE1=="F - Construction",],
  index=c("NACE2", "NACE3", "NACE4"), show.labels=TRUE, truncate.labels=c(2,4,6))
treegraph(business[business$NACE1=="F - Construction",],
  index=c("NACE2", "NACE3", "NACE4"), show.labels=TRUE, truncate.labels=c(2,4,6),
  vertex.layout="fruchterman.reingold")
```

Description

A treemap is a space-filling visualization of hierarchical structures. This function offers great flexibility to draw treemaps. Required is a data.frame (`dtf`) that contains one or more hierarchical index columns given by `index`, a column that determines the rectangle area sizes (`vSize`), and optionally a column that determines the rectangle colors (`vColor`). The way how rectangles are colored is determined by the argument `type`.

Usage

```r
treemap(
  dtf,
  index,
  vSize,
  vColor = NULL,
  stdErr = NULL,
  type = "index",
  fun.aggregate = "sum",
  title = NA,
  title.legend = NA,
  algorithm = "pivotSize",
  sortID = "-size",
  mirror.x = FALSE,
  mirror.y = FALSE,
  palette = NA,
  palette.HCL.options = NULL,
  range = NA,
  mapping = NA,
  n = 7,
  na.rm = TRUE,
  na.color = "#DDDDDD",
  na.text = "Missing",
  fontsize.title = 14,
  fontsize.labels = 11,
  fontsize.legend = 12,
)```
fontcolor.labels = NULL,
fontName.labels = c("bold", rep("plain", length(index) - 1)),
fontfamily.title = "sans",
fontfamily.labels = "sans",
fontfamily.legend = "sans",
border.col = "black",
border.lwds = c(length(index) + 1, (length(index) - 1):1),
lowerbound.cex.labels = 0.4,
inflate.labels = FALSE,
bg.labels = NULL,
force.print.labels = FALSE,
overlap.labels = 0.5,
align.labels = c("center", "center"),
xmod.labels = 0,
ymod.labels = 0,
eval.labels = FALSE,
position.legend = NULL,
reverse.legend = FALSE,
format.legend = NULL,
drop.unused.levels = TRUE,
aspRatio = NA,
vp = NULL,
draw = TRUE,
...
)

Arguments

dtf
a data.frame. Required.

index
vector of column names in dtf that specify the aggregation indices. It could
contain only one column name, which results in a treemap without hierarchy.
If multiple column names are provided, the first name is the highest aggrega-
tion level, the second name the second-highest aggregation level, and so on.
Required.

vSize
name of the column in dtf that specifies the sizes of the rectangles. Required.

vColor
name of the column that, in combination with type, determines the colors of the
rectangles. The variable can be scaled by the addition of "*<scale factor>" or
"/<scale factor>". Note: when omitted for "value" treemaps, a constant value of
1 is taken.

stdErr
name of the column that contains standard errors. These are not used for the
treemaps, but only aggregated accordingly and returned as item of the output
list.

type
type of the treemap, which determines how the rectangles are colored:

"index": colors are determined by the index variables. Different branches
in the hierarchical tree get different colors. For this type, vColor
is not needed.
"value": the numeric vColor-column is directly mapped to a color palette. This palette is diverging, so that values of 0 are assigned to the mid color (white or yellow), and negative and positive values are assigned to color based on two different hues colors (by default reds for negative and greens for positive values). For more freedom, see "manual".

"comp": colors indicate change of the vSize-column with respect to the numeric vColor-column in percentages. Note: the negative scale may be different from the positive scale in order to compensate for the ratio distribution.

"dens": colors indicate density. This is analogous to a population density map where vSize-values are area sizes, vColor-values are populations per area, and colors are computed as densities (i.e. population per squared km).

"depth": each aggregation level (defined by index) has a distinct color. For this type, vColor is not needed.

"categorical": vColor is a factor column that determines the color.

"color": vColor is a vector of colors in the hexadecimal (#RRGGBB) format

"manual": The numeric vColor-column is directly mapped to a color palette. Both palette and range should be provided. The palette is mapped linearly to the range.

fun.aggregate aggregation function, only used in "value" treemaps. This function determines how values of the lowest aggregation level are aggregated. By default, it takes the sum. Other sensible functions are mean and weighted.mean. In the latter case, the weights are determined by the vSize variable. Other arguments can be passed on. For weighted.mean, it is possible to assign a variable name for its w argument.

title title of the treemap.
title.legend title of the legend.

algorithm name of the used algorithm: "squarified" or "pivotSize". The squarified treemap algorithm (Bruls et al., 2000) produces good aspect ratios, but ignores the sorting order of the rectangles (sortID). The ordered treemap, pivot-by-size, algorithm (Bederson et al., 2002) takes the sorting order (sortID) into account while aspect ratios are still acceptable.

sortID name of the variable that determines the order in which the rectangles are placed from top left to bottom right. Only applicable when algorithm="pivotSize". Also the values "size" and "color" can be used, which refer to vSize and vColor respectively. To inverse the sorting order, use "-" in the prefix. By default, large rectangles are placed top left.

mirror.x logical that determines whether the rectangles are mirrored horizontally

mirror.y logical that determines whether the rectangles are mirrored vertically

palette one of the following:

**a color palette**: i.e., a vector of hexadecimal colors (#RRGGBB)

**a name of a Brewer palette**: See RColorBrewer::display.brewer.all() for the options. The palette can be reversed by prefixing with a ".". For treemap types "value" and "comp", a diverging palette should be chosen (default="RdYIgN"), for type "dens" a sequential (default="OrRd"). The default value for "depth" is "Set2".
"HCL": Tree Colors are color schemes derived from the Hue-Chroma-Luminance color space model. This is only applicable for qualitative palettes, which are applied to the treemap types "index", "depth", and "categorical". For "index" and "categorical" this is the default value.

palette.HCL.options

list of advanced options to obtain Tree Colors from the HCL space (when palette="HCL"). This list contains:

hue_start: number between 0 and 360 that determines the starting hue value (default: 30)

hue_end: number between hue_start and hue_start + 360 that determines the ending hue value (default: 390)

hue_perm: boolean that determines whether the colors are permuted such that adjacent levels get more distinguishable colors. If FALSE, then the colors are equally distributed from hue_start to hue_end (default: TRUE)

hue_rev: boolean that determines whether the colors of even-numbered branched are reversed (to increase discrimination among branches)

hue_fraction: number between 0 and 1 that determines the fraction of the hue circle that is used for recursive color picking: if 1 then the full hue circle is used, which means that the hue of the colors of lower-level nodes are spread maximally. If 0, then the hue of the colors of lower-level nodes are identical of the hue of their parents. (default: .5)

chroma: chroma value of colors of the first-level nodes, that are determined by the first index variable (default: 60)

luminance: luminance value of colors of the first-level nodes, i.e. determined by the first index variable (default: 70)

chroma_slope: slope value for chroma of the non-first-level nodes. The chroma values for the second-level nodes are chroma+chroma_slope, for the third-level nodes chroma+2*chroma_slope, etc. (default: 5)

luminance_slope: slope value for luminance of the non-first-level nodes (default: -10)

For "depth" and "categorical" types, only the first two items are used. Use treecolors to experiment with these parameters.

range

range of values (so vector of two) that correspond to the color legend. By default, the range of actual values, determined by vColor, is used. Only applicable for numeric types, i.e. "value", "comp", "dens", and "manual". Note that the range doesn’t affect the colors in the treemap itself for "value" and "manual" types; this is controlled by mapping.

mapping

vector of three values that specifies the mapping of the actual values, determined by vColor, to palette. The three values are respectively the minimum value, the mid value, and the maximum value. The mid value is particularly useful for diverging color palettes, where it defined the middle, neutral, color which is typically white or yellow. The mapping should cover the range. By default, for "value" treemaps, it is c(-max(abs(values)), 0, max(abs(values))), where values are the actual values defined by vColor. For "manual" treemaps, the default setting is c(min(values), mean(range(values)), max(values)). A vector of two can also be specified. In that case, the mid value will be the average of those. Only applicable for "value" and "manual" type treemaps.
preferred number of categories by which numeric variables are discretized.

ignore missing values for the vSize variable (by default TRUE)

color for missing values for the vColor variable

legend label for missing values for the vColor variable

font size of the title

font size(s) of the data labels, which is either a single number that specifies the font size for all aggregation levels, or a vector that specifies the font size for each aggregation level. Use value 0 to omit the labels for the corresponding aggregation level.

font size for the legend

Specifies the label colors. Either a single color value, or a vector of color values one for each aggregation level. By default, white and black colors are used, depending on the background (bg.labels).

either a single value, or a vector of values one for each aggregation level. Values can be integers If an integer, following the R base graphics standard: 1 = plain, 2 = bold, 3 = italic, 4 = bold italic, or characters: "plain", "bold", "italic", "oblique", and "bold.italic".

font family of the title. Standard values are "serif", "sans", "mono", "symbol". Mapping is device dependent.

font family of the labels in each rectangle. Standard values are "serif", "sans", "mono", "symbol". Mapping is device dependent.

font family of the legend. Standard values are "serif", "sans", "mono", "symbol". Mapping is device dependent.

color of borders drawn around each rectangle. Either one color for all rectangles or a vector of colors, or one for each aggregation level

thicknesses of border lines. Either one number specifies the line thicknesses (widths) for all rectangles or a vector of line thicknesses for each aggregation level.

multiplier between 0 and 1 that sets the lowerbound for the data label font sizes: 0 means draw all data labels, and 1 means only draw data labels if they fit (given fontsize.labels).

logical that determines whether data labels are inflated inside the rectangles. If TRUE, fontsize.labels does not determine the fontsize anymore, but it still determines the minimum fontsize in combination with lowerbound.cex.labels.

background color of high aggregation labels. Either a color, or a number between 0 and 255 that determines the transparency of the labels. In the latter case, the color itself is determined by the color of the underlying rectangle.
For "value" and "categorical" treemaps, the default is (slightly) transparent grey ("#CCCCCCDC"), and for the other types slightly transparent: 220.

force.print.labels

logical that determines whether data labels are being forced to be printed if they don’t fit.

overlap.labels

number between 0 and 1 that determines the tolerance of the overlap between labels. 0 means that labels of lower levels are not printed if higher level labels overlap, 1 means that labels are always printed. In-between values, for instance the default value .5, means that lower level labels are printed if other labels do not overlap with more than .5 times their area size.

align.labels

object that specifies the alignment of the labels. Either a character vector of two values specifying the horizontal alignment ("left", "center", or "right") and the vertical alignment ("top", "center", or "bottom"), or a list of such character vectors, one for each aggregation level.

xmod.labels

the horizontal position modification of the labels in inches. Options: a single value, a vector or a list that specifies the modification for each aggregation level. If a list is provided, each list item consists of a single value or a named vector that specify the modification per label.

ymod.labels

the vertical position modification of the labels in inches. Options: a single value, a vector or a list that specifies the modification for each aggregation level. If a list is provided, each list item consists of a single value or a named vector that specify the modification per label.

eval.labels

should the text labels, i.e. the factor labels of the index variables, be evaluated as expressions? Useful for printing mathematical symbols or equations.

position.legend

position of the legend: "bottom", "right", or "none". For "categorical" and "index" treemaps, "right" is the default value, for "index" treemap, "none", and for the other types, "bottom".

reverse.legend

should the legend be reversed?

format.legend

a list of additional arguments for the formatting of numbers in the legend to pass to format(); only applies if type is "value", "dens" or "manual".

drop.unused.levels

logical that determines whether unused levels (if any) are shown in the legend. Applicable for "categorical" treemap type.

aspRatio

preferred aspect ratio of the main rectangle, defined by width/height. When set to NA, the available window size is used.

vp

viewport to draw in. By default it is not specified, which means that a new plot is created. Useful when drawing small multiples, or when placing a treemap in a custom grid based plot.

draw

logical that determines whether to draw the treemap.

... 

arguments to be passed to other functions. Currently, only fun.aggregate takes optional arguments.
Value

A list is silently returned:

`tm` a `data.frame` containing information about the rectangles: indices, sizes, original color values, derived color values, depth level, position (x0, y0, w, h), and color.

type argument type
vSize argument vSize
cColor argument vColor
stdErr standard errors
algorithm argument algorithm
vpCoorX x-coordinates of the treemap within the whole plot
vpCoorY y-coordinates of the treemap within the whole plot
aspRatio aspect ratio of the treemap
range range of the color values scale

References


Examples

#########################################
### quick example with Gross National Income data
#########################################
data(GNI2014)
treemap(GNI2014, 
  index=c("continent", "iso3"),
  vSize="population",
  cColor="GNI",
  type="value",
  format.legend = list(scientific = FALSE, big.mark = " "))

#########################################
### extended examples with fictive business statistics data
#########################################
data(business)

#########################################
### treemap types
#########################################

# index treemap: colors are determined by the index argument
## Not run:
# large example which takes some time...
```r
treemap(business,
    index=c("NACE1", "NACE2", "NACE3"),
    vSize="turnover",
    type="index")
```
## End(Not run)
```r
treemap(business[business$NACE1=="C - Manufacturing",],
    index=c("NACE2", "NACE3"),
    vSize=c("employees"),
    type="index")
```

# value treemap: colors are derived from a numeric variable given by vColor
# (when omitted, all values are set to 1 as in the following example)
```r
treemap(business,
    index=c("NACE1", "NACE2"),
    vSize="employees",
    title.legend="number of NACE4 categories",
    type="value")
```

# comparison treemaps: colors indicate change of vSize with respect to vColor
```r
treemap(business,
    index=c("NACE1", "NACE2"),
    vSize="employees",
    vColor="employees.prev",
    type="comp")
```

# density treemaps: colors indicate density (like a population density map)
```r
treemap(business,
    index=c("NACE1", "NACE2"),
    vSize="turnover",
    vColor="employees/1000",
    type="dens")
```
## Not run:
# depth treemap: show depth
```r
treemap(business,
    index=c("NACE1", "NACE2", "NACE3"),
    vSize="turnover",
    type="depth")
```
## End(Not run)

# categorical treemap: colors are determined by a categorical variable
```r
business <- transform(business, data.available = factor(!is.na(turnover)), x = 1)
treemap(business,
    index=c("NACE1", "NACE2"),
    vSize="x",
    vColor="data.available",
    type="categorical")
```
## Not run:
# color treemap

business$color <- rainbow(nlevels(business$NACE2))[business$NACE2]
treemap(business,
  index=c("NACE1", "NACE2"),
  vSize="x",
  vColor="color",
  type="color")

# manual

business$color <- rainbow(nlevels(business$NACE2))[business$NACE2]
treemap(business,
  index=c("NACE1", "NACE2"),
  vSize="turnover",
  vColor="employees",
  type="manual",
  palette=terrain.colors(10))

## End(Not run)

#################################################################
### graphical options: control fontsizes
#################################################################

## Not run:
# draw labels of first index at fontsize 12 at the center,
# and labels of second index at fontsize 8 top left

treemap(business,
  index=c("NACE1", "NACE2"),
  vSize="employees",
  fontsize.labels=c(12, 8),
  align.labels=list(c("center", "center"), c("left", "top")),
  lowerbound.cex.labels=1)

# draw all labels at fontsize 12 (only if they fit)

treemap(business,
  index=c("NACE1", "NACE2"),
  vSize="employees",
  fontsize.labels=12,
  lowerbound.cex.labels=1)

# draw all labels at fontsize 12, and if they don't fit, reduce to a minimum of .6*12

treemap(business,
  index=c("NACE1", "NACE2"),
  vSize="employees",
  fontsize.labels=12,
  lowerbound.cex.labels=.6)

# draw all labels at maximal fontsize

treemap(business,
  index=c("NACE1", "NACE2"),
  vSize="employees",
  lowerbound.cex.labels=0,
treemap(business, index=c("NACE1", "NACE2"),
vSize="employees",
fontsize.labels=10,
lowerbound.cex.labels=1,
force.print.labels=TRUE)

# draw all labels at fixed fontsize, even if they don’t fit
	treemap(business, index=c("NACE1", "NACE2"),
vSize="employees",
fontsize.labels=10,
lowerbound.cex.labels=1,
force.print.labels=TRUE)

############################################
### graphical options: color palettes
############################################

## for comp and value typed treemaps all diverging brewer palettes can be chosen

treemap(business, index=c("NACE1", "NACE2"),
vSize="employees",
vColor="employees.prev",
type="comp",
palette="RdBu")

## draw warm-colored index treemap

draw warm-colored index treemap

palette.HCL.options <- list(hue_start=270, hue_end=360+150)

treemap(business, index=c("NACE1", "NACE2"),
vSize="employees",
type="index",
palette.HCL.options=palette.HCL.options)

# terrain colors

business$employees.growth <- business$employees - business$employees.prev
treemap(business, index=c("NACE1", "NACE2"),
vSize="employees",
vColor="employees.growth",
type="value",
palette=terrain.colors(10))

# Brewer’s Red-White-Grey palette reversed with predefined legend range

treemap(business, index=c("NACE1", "NACE2"),
vSize="employees",
vColor="employees.growth",
type="value",
palette="-RdGy",
range=c(-20000,30000))

# More control over the color palette can be achieved with mapping

treemap(business, index=c("NACE1", "NACE2"),
vSize="employees",
vColor="employees.growth",
treepalette

Description

Obtain hierarchical color palettes, either the so-called Tree Colors from the HCL color space model, or by using an existing color palette. The former method, which is recommended, is used by default in treemap (type "index") and treegraph. Use treecolors to experiment with this method.

Usage

treepalette(
  dtf,
  index = names(dtf),
  method = "HCL",
  palette = NULL,
  palette.HCL.options,
  return.parameters = TRUE,
  prepare.dat = TRUE
)

Arguments

dtf a data.frame or data.table. Required.
index the index variables of dtf
method used method: either "HCL" (recommended), which is based on the HCL color space model, or "HSV", which uses the argument palette.
palette color palette, which is only used for the HSV method
palette.HCL.options list of options to obtain Tree Colors from the HCL space (when palette="HCL"). This list contains:
  hue_start: number between 0 and 360 that determines the starting hue value (default: 30)
  hue_end: number between hue_start and hue_start + 360 that determines the ending hue value (default: 390)
  hue_perm: boolean that determines whether the colors are permuted such that adjacent levels get more distinguishable colors. If FALSE, then the colors are equally distributed from hue_start to hue_end (default: TRUE)
treepalette

**hue_rev**: boolean that determines whether the colors of even-numbered branched are reversed (to increase discrimination among branches)

**hue_fraction**: number between 0 and 1 that determines the fraction of the hue circle that is used for recursive color picking: if 1 then the full hue circle is used, which means that the hue of the colors of lower-level nodes are spread maximally. If 0, then the hue of the colors of lower-level nodes are identical of the hue of their parents. (default: .5)

**chroma**: chroma value of colors of the first-level nodes, that are determined by the first index variable (default: 60)

**luminance**: luminance value of colors of the first-level nodes, i.e. determined by the first index variable (default: 70)

**chroma_slope**: slope value for chroma of the non-first-level nodes. The chroma values for the second-level nodes are chroma+chroma_slope, for the third-level nodes chroma+2*chroma_slope, etc. (default: 5)

**luminance_slope**: slope value for luminance of the non-first-level nodes (default: -10)

For "depth" and "categorical" types, only the first two items are used. Use treecolors to experiment with these parameters.

**return.parameters**

should a data.frame with color values and parameter options be returned (TRUE), or just the vector of color values (FALSE)?

**prepare.dat**

data is by default preprocessed, except for internal use

**Value**

Either a vector of colors, or a data.frame is return (see return.parameters).
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