

# Package ‘tabplot’

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**License** GPL-3

**Title** Tableplot, a visualization of large datasets

**Type** Package

**LazyLoad** yes

**Author** Martijn Tennekes and Edwin de Jonge

**Description** A tableplot is a visualisation of a (large) dataset with a dozen of variables, both numeric and categorical. Each column represents a variable and each row bin is an aggregate of a certain number of records. Numeric variables are visualized as bar charts, and categorical variables as stacked bar charts. Missing values are taken into account. Also supports large ffd datasets from the ff package.

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**VignetteBuilder** knitr

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**NeedsCompilation** no

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tabplot-package	<i>Tableplot, a visualization of large datasets</i>
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## Description

Package: tabplot  
 Type: Package  
 Version: 1.1  
 Date: 2013-02-24  
 License: GPL-3  
 LazyLoad: yes

## Details

A tableplot is a visualisation of a (large) dataset. Each column represents a variable and each row bin is an aggregate of a certain number of records. For numeric variables, a bar chart of the mean values is depicted. For categorical variables, a stacked bar chart is depicted of the proportions of categories. Missing values are taken into account. Also supports large [ffdf](#) datasets from the [ff](#) package.

The main function of the package is [tableplot](#), which is used to create a tableplot. Other useful functions are:

- [itableplot](#) to start a graphical user interface (made with the [shiny](#) package);
- [tablePrepare](#) to prepare a large dataset. Tableplotting is much faster when the returned object is passed on to [tableplot](#) rather than the dataset itself;
- [tablePalettes](#) to show all quantitative and qualitative palettes that are included;
- [tableSave](#) to save a tableplot;
- [tableChange](#) to make layout changes to a tableplot.

For a quick intro, see `vignette("tabplot-vignette")`.

### Author(s)

Martijn Tennekes <[mtennekes@gmail.com](mailto:mtennekes@gmail.com)> and Edwin de Jonge

### Examples

```
# load diamonds dataset from ggplot2
require(ggplot2)
data(diamonds)

# create tableplot
tableplot(diamonds)
```

---

-.tabplot

*Compare two tableplots (experimental)*

---

### Description

Two tableplots can be compared by subtracting two [tableplot-objects](#). The result is a [tableplot\\_compare-object](#) object in which absolute and relative differences of mean values are stored, as well as a comparison of frequency tables for categorical variables. This object can be plotted with [plot](#).

### Usage

```
## S3 method for class 'tableplot'
tp1 - tp2
```

### Arguments

tp1            the first [tableplot-object](#)  
tp2            the second [tableplot-object](#)

### Value

a [tableplot\\_compare-object](#) that contains information about the comparison `tp1-tp2`

**Examples**

```
# load diamonds dataset from ggplot2
require(ggplot2)
data(diamonds)

# calculate normalized prices to be used as sample probabilities
price.norm <- with(diamonds, price / max(diamonds$price))

# draw samples
exp.diamonds <- diamonds[sample(1:nrow(diamonds), size=10000, prob=price.norm, replace=TRUE),]
chp.diamonds <- diamonds[sample(1:nrow(diamonds), size=10000, prob=1-price.norm, replace=TRUE),]

tp1 <- tableplot(exp.diamonds)
tp2 <- tableplot(chp.diamonds)

plot(tp2 - tp1)
```

---

`bin_data`*Bin data*

---

**Description**

Working horse for tableplot, does the actual binning

**Usage**

```
bin_data(p, sortCol = 1L, cols = seq_along(p$data), from = 0, to = 1,
  nbins = 100L, decreasing = FALSE, sample = FALSE, sampleBinSize = 100)
```

**Arguments**

<code>p</code>	prepared dataset (see <a href="#">tablePrepare</a> )
<code>sortCol</code>	column on which the table will be sorted
<code>cols</code>	columns of the data that will be used.
<code>from</code>	lower boundary in quantiles
<code>to</code>	upper boundary in quantiles
<code>nbins</code>	number of bins
<code>decreasing</code>	sort decreasingly
<code>sample</code>	sample or use whole dataset?
<code>sampleBinSize</code>	sample size per bin

---

bin_hcc_data	<i>Bin high cardinality data</i>
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---

**Description**

Bin high cardinality data

**Usage**

```
bin_hcc_data(bd, max_levels)
```

**Arguments**

bd	binned dataset (result of <a href="#">bin_data</a> )
max_levels	maximum number of levels. Each column in bd that has more than max_levels categories is rebinned to max_levels categories.

---

datetime2fac	<i>Transform a date-time vector to a factor</i>
--------------	---

---

**Description**

Transform a date-time vector from class [POSIXt](#) or [Date](#) to a factor.

**Usage**

```
datetime2fac(p, rng = range(p, na.rm = TRUE))
```

**Arguments**

p	date-time vector
rng	range of the factor.

**Details**

The range rng is cut according to different pretty rounded time periods. The cut with the number of levels that is closest to 6 is chosen. Vector p is cut accordingly. Values of p outside rng are translated to NA.

**Value**

A factor vector.

**Note**

This function is still in development stage, and can be improved and optimized. `ff` vectors are not implemented yet

**See Also**

[num2fac](#)

**Examples**

```
d <- as.Date("2012-12-21") + sample.int(500, 1000, replace=TRUE)
d2 <- datetime2fac(d)
levels(d2)

t <- as.POSIXlt(Sys.time(), "GMT") + sample.int(1e5, 1000, replace=TRUE)
t2 <- datetime2fac(t)
levels(t2)
```

---

itableplot

*Graphical User Interface to create tableplots*

---

**Description**

This graphical user interface is developed with the [shiny](#) package. All datasets that are loaded in the global workspace ([data.frame](#), [ffdf](#), or [prepared](#)) are passed on to the GUI.

**Usage**

```
itableplot()
```

**Details**

This function replaces the old `tabplotGTK` package, since it only requires an up-to-date browser (and not software like GTK). Furthermore, maintenance is a lot easier.

**Examples**

```
## Not run:
require(ggplot2)
data(diamonds)

# load other datasets
data(iris)
data(cars)

itableplot()

## End(Not run)
```

---

num2fac	<i>Transform a numerical vector to a factor</i>
---------	---

---

## Description

Transform a numerical vector from class [POSIXt](#) or [Date](#) to a factor.

## Usage

```
num2fac(num, method = "pretty", num_scale = "auto", n = 0, brks = NA)
```

## Arguments

num	numeric vector
method	<ul style="list-style-type: none"><li>• "pretty" intervals are determined by the base function <a href="#">pretty</a></li><li>• "kmeans" the method intervals are determined by the method <a href="#">kmeans</a> where n clusters (i.e. intervals) are found</li><li>• "fixed" determines the intervals by the argument brks</li><li>• "discrete" the unique values in num are mapped one to one to the levels of the new factor vector)</li></ul>
num_scale	<ul style="list-style-type: none"><li>• "auto" used scale is determined automatically</li><li>• "lin" num is directly fed to the method <a href="#">pretty</a> or <a href="#">kmeans</a></li><li>• "log" a logarithmic transformation of num is fed to the method <a href="#">pretty</a> or <a href="#">kmeans</a></li></ul>
n	the (desired) number of levels. n=0 means automatic
brks	breaks that determine the levels (only required when method="fixed")

## Value

A factor vector

## Note

This function is still in development stage, and can be improved and optimized. ff vectors are not implemented yet

## See Also

[datetime2fac](#)

## Examples

```
require(ggplot2)
data(diamonds)

diamonds$price2 <- num2fac(diamonds$price)

tableplot(diamonds)
```

---

plot.tabplot      *Plot a [tabplot-object](#)*

---

## Description

Plot a [tabplot-object](#)

## Usage

```
## S3 method for class 'tabplot'
plot(x, fontsize = 10, legend.lines = 8,
     max_print_levels = 15, text_NA = "missing", title = NULL,
     showTitle = NULL, fontsize.title = 14, showNumAxes = TRUE,
     relative = FALSE, vp = NULL, ...)

## S3 method for class 'tabplot_compare'
plot(x, ...)
```

## Arguments

x	<a href="#">tabplot-object</a> or <a href="#">tabplot_compare-object</a>
fontsize	the (maximum) fontsize
legend.lines	the number of lines preserved for the legend
max_print_levels	maximum number of printed category labels in the legend
text_NA	text printed for the missing values category in the legend
title	title of the plot (shown if showTitle==TRUE)
showTitle	show the title. By default FALSE, unless a title is given.
fontsize.title	the fontsize of the title
showNumAxes	plots an x-axis for each numerical variable, along with grid lines (TRUE by default).
relative	boolean that determines whether relative scales are used for relative tableplots. If TRUE, then <code>mean.diff.rel&lt;-(mean2-mean1)/mean1*100</code> are used. If FALSE, then the absolute difference is taken: <code>mean &lt;- mean2-mean</code> .
vp	<a href="#">viewport</a> to draw plot in (for instance useful to stack multiple tableplots)
...	other arguments are not used

## Examples

```
# load diamonds dataset from ggplot2
require(ggplot2)
data(diamonds)

tab <- tableplot(diamonds)
```



```
plot(tab, title="Shine on you Crazy Diamond!!!",
      fontsize=12,
      legend.lines=7,
      fontsize.title=16)
```

---

print.tabplot      *Print a [tabplot-object](#)*

---

### Description

Print a [tabplot-object](#)

### Usage

```
## S3 method for class 'tabplot'
print(x, ...)
```

### Arguments

x	tabplot object
...	arguments passed to other methods

---

summary.tabplot      *Summarize a [tabplot-object](#)*

---

### Description

Summarize a [tabplot-object](#)

### Usage

```
## S3 method for class 'tabplot'
summary(object, digits = max(3, getOption("digits") - 3),
        ...)
```

### Arguments

object	tabplot object
digits	integer, used for number formatting with <a href="#">format</a>
...	arguments passed to other methods

---

tableChange	<i>Change a <a href="#">tabplot-object</a></i>
-------------	--

---

### Description

Make layout changes in a [tabplot-object](#), such as the order of columns, and color palettes.

### Usage

```
tableChange(tab, select = NULL, select_string = tab$select,
  decreasing = NULL, pals = list(), colorNA = NULL, numPals = NULL)
```

### Arguments

tab	<a href="#">tabplot-object</a>
select	index vector of the desired columns (column names are not supported)
select_string	vector of names of the desired columns
decreasing	determines whether the dataset is sorted decreasingly (TRUE) of increasingly (FALSE).
pals	list of color palettes. Each list item is one of the following: <ul style="list-style-type: none"> <li>• a palette name in <a href="#">tablePalettes</a>, optionally with the starting color between brackets.</li> <li>• a palette vector</li> </ul> <p>If the list items are unnamed, they are applied to all selected categorical variables (recycled if necessary). The list items can be assigned to specific categorical variables, by naming them accordingly.</p>
colorNA	color for missing values
numPals	name(s) of the palette(s) that is(are) used for numeric variables ("Blues", "Greys", or "Greens"). Recycled if necessary.

### Value

[tabplot-object](#)

### Examples

```
# load diamonds dataset from ggplot2
require(ggplot2)
data(diamonds)

# assign tableplot as tabplot object
tab <- tableplot(diamonds)

# modify the tabplot object: reverse order of columns and customize palette
tab <- tableChange(tab, select_string=rev(names(diamonds)),
```

```

pals=list(clarity=gray(seq(0,1,length.out=8)))

# plot modified tabplot object
plot(tab)

```

---

tablePalettes	<i>Show / get all palettes of the tabplot package</i>
---------------	---

---

## Description

All color palettes are shown and/or returned that can be used for tableplots.

## Usage

```
tablePalettes(plot = TRUE)
```

## Arguments

`plot` Boolean that determines whether the palettes are plot.

## Details

Sequential palettes (for numeric variables): "Blues", "Greens", "Greys", "Oranges", and "Purples". These palettes are taken from ColorBrewer (Brewer et al., 2003).

Qualitative palattes (for categorical variables): "Set1", "Set2", "Set3", "Set4", "Set5", "Set6", "Set7", "Set8", "Paired", "HCL1", "HCL2", and "HCL3". The default palette, "Set1", is a colorblind-friendly palette (see Okabe and Ito, 2002). Palettes "Set2" to "Set6" and "Paired" are based on ColorBrewer palettes (Brewer et al., 2003). Palette "Set7", is a colorblind-friendly palette from the dichromat package (see Thomas Lumley , 2012). Palette "Set8" is a palette created by Wijffelaars (2008). The "HCL" Palettes are based on the Hue-Chroma-Luminance color space model (see Zeileis et al., 2009). The color red has been removed from the original palettes, since it is occupied by missing values.

## Value

list with palettes (silent output)

## References

Brewer, Cynthia A., Geoffrey W. Hatchard and Mark A. Harrower, 2003, ColorBrewer in Print: A Catalog of Color Schemes for Maps, *Cartography and Geographic Information Science* 30(1): 5-32.

Okabe, M. and Ito, K. Color Universal Design (CUD) - How to make figures and presentations that are friendly to Colorblind people, 2002

Wijffelaars, M. Synthesis of Color Palettes. Master's thesis. Supervisors Wijk, J. van, and Vliegen, R. 2008

Thomas Lumley (2012). dichromat: Color schemes for dichromats. R package version 1.2-4. <http://CRAN.R-project.org/package=dichromat>

Zeileis, A., Hornik, K., and Murrell, P. Escaping RGBland: Selecting colors for statistical graphics. In *Proceedings of Computational Statistics & Data Analysis*. 2009, 3259-3270.

---



*Create a tableplot*


---

## Description

A tableplot is a visualisation of (large) multivariate datasets. Each column represents a variable and each row bin is an aggregate of a certain number of records. For numeric variables, a bar chart of the mean values is depicted. For categorical variables, a stacked bar chart is depicted of the proportions of categories. Missing values are taken into account. Also supports large `ffdf` datasets from the `ff` package. For a quick intro, see `vignette("tableplot-vignette")`.

## Usage

```
tableplot(dat, select, subset = NULL, sortCol = 1, decreasing = TRUE,
  nBins = 100, from = 0, to = 100, nCols = ncol(dat), sample = FALSE,
  sampleBinSize = 1000, scales = "auto", max_levels = 50,
  pals = list("Set1", "Set2", "Set3", "Set4"), change_palette_type_at = 20,
  colorNA = "#FF1414", numPals = "Blues", limitsX = NULL,
  bias_brokenX = 0.8, IQR_bias = 5, select_string = NULL,
  subset_string = NULL, colNames = NULL, filter = NULL, plot = TRUE,
  ...)
```

## Arguments

<code>dat</code>	a <code>data.frame</code> , an <code>ffdf</code> object, or an object created by <code>tablePrepare</code> (see details below). Required.
<code>select</code>	expression indicating the columns of <code>dat</code> that are visualized in the tableplot. Also column indices are supported. By default, all columns are visualized. Use <code>select_string</code> for character strings instead of expressions.
<code>subset</code>	logical expression indicating which rows to select in <code>dat</code> (as in <code>subset</code> ). It is also possible to provide the name of a categorical variable: then, a tableplot for each category is generated. Use <code>subset_string</code> for character strings instead of an expressions.
<code>sortCol</code>	column name on which the dataset is sorted. It can be either an index or an expression name. Also a character string can be used, but this is discouraged for programming purposes (use an index instead).
<code>decreasing</code>	boolean that determines whether the dataset is sorted decreasingly ( <code>TRUE</code> ) or increasingly ( <code>FALSE</code> ).
<code>nBins</code>	number of row bins
<code>from</code>	percentage from which the sorted data is shown
<code>to</code>	percentage to which the sorted data is shown

nCols	the maximum number of columns per tableplot. If this number is smaller than the number of columns selected in <code>datNames</code> , multiple tableplots are generated, where each of them contains the sorted column(s).
sample	boolean that determines whether to sample or use the whole data. Only useful when <code>tablePrepare</code> is used.
sampleBinSize	the number of sampled objects per bin, if <code>sample</code> is TRUE.
scales	determines the horizontal axes of the numeric variables in <code>colNames</code> . Options: "lin", "log", and "auto" for automatic detection. Either <code>scale</code> is a named vector, where the names correspond to numerical variable names, or <code>scale</code> is unnamed, where the values are applied to all numeric variables (recycled if necessary).
max_levels	maximum number of levels for categorical variables. Categorical variables with more levels will be rebinned into <code>max_levels</code> levels. Either a positive number or -1, which means that categorical variables are never rebinned.
pals	list of color palettes. Each list item is one of the following: <ul style="list-style-type: none"> <li>• a palette name of <code>tablePalettes</code>, optionally with the starting color between brackets.</li> <li>• a color vector</li> </ul> <p>If the list items are unnamed, they are applied to all selected categorical variables (recycled if necessary). The list items can be assigned to specific categorical variables, by naming them accordingly.</p>
change_palette_type_at	number at which the type of categorical palettes is changed. For categorical variables with less than <code>change_palette_type_at</code> levels, the palette is recycled if necessary. For categorical variables with <code>change_palette_type_at</code> levels or more, a new palette of interpolated colors is derived (like a rainbow palette).
colorNA	color for missing values
numPals	vector of palette names that are used for numeric variables. These names are chosen from the sequential palette names in <code>tablePalettes</code> . Either <code>numPals</code> is a named vector, where the names correspond to the numerical variable names, or an unnamed vector (recycled if necessary).
limitsX	a list of vectors of length two, where each vector contains a lower and an upper limit value. Either the names of <code>limitsX</code> correspond to numerical variable names, or <code>limitsX</code> is an unnamed list (recycled if necessary).
bias_brokenX	parameter between 0 and 1 that determines when the x-axis of a numeric variable is broken. If minimum value is at least <code>bias_brokenX</code> times the maximum value, then X axis is broken. To turn off broken x-axes, set <code>bias_brokenX</code> =1.
IQR_bias	parameter that determines when a logarithmic scale is used when <code>scales</code> is set to "auto". The argument <code>IQR_bias</code> is multiplied by the interquartile range as a test.
select_string	character equivalent of the <code>select</code> argument (particularly useful for programming purposes)
subset_string	character equivalent of the <code>subset</code> argument (particularly useful for programming purposes)

colNames	deprecated; used in older versions of tabplot (prior to 0.12): use <code>select_string</code> instead
filter	deprecated; used in older versions of tabplot (prior to 0.12): use <code>subset_string</code> instead
plot	boolean, to plot or not to plot a tableplot
...	layout arguments, such as <code>fontsize</code> and <code>title</code> , are passed on to <code>plot</code>

### Details

For large dataset, we recommend to use `tablePrepare` which does all the necessary preprocessing that are needed to make any tableplot of the particular dataset. The resulting object of this function is passed on to `tableplot` (argument `dat`). Now tableplotting is very fast, and even faster with sampling enabled (`sample=TRUE`).

### Value

`tableplot-object` (silent output). If multiple tableplots are generated (which can be done by either setting `subset` to a categorical column name, or by restricting the number of columns with `nCols`), then a list of `tableplot-objects` is silently returned.

### Note

In early development versions of `tabplot` (prior to version 1.0) it was possible to sort datasets on multiple columns. To increase to `tableplot` creation speed, this feature is dropped. For multiple sorting purposes, we recommend to use the `subset` parameter instead.

### See Also

[itableplot](#)

### Examples

```
# load diamonds dataset from ggplot2
require(ggplot2)
data(diamonds)

# default tableplot
tableplot(diamonds)

# most expensive diamonds
tableplot(diamonds,
  select=c(carat, cut, color, clarity, price),
  sortCol=price,
  from=0,
  to=5)

# for large datasets, we recommend to preprocess the data with tablePrepare:
p <- tablePrepare(diamonds)

# specific subsetting
```

```

tableplot(p, subset=price < 5000 & cut=='Ideal')

# change palettes
tableplot(p,
  pals=list(cut="Set4", color="Paired", clarity=grey(seq(0, 1,length.out=7))),
  numPals=c(carat="Greens", price="Purples"))

# create a tableplot cut category, and fix scale limits of carat, table, and price
tabs <- tableplot(p, subset=cut,
  limitsX=list(carat=c(0,4), table=c(55, 65), price=c(0, 20000)), plot=FALSE)
plot(tabs[[3]], title="Very good cut diamonds")

```

---

tablePrepare	<i>Prepares a dataset for tableplotting</i>
--------------	---

---

## Description

Tableplots from a large dataset can be generated very fast when the preprocessing stage is done only once. This function preprocesses the dataset, and returns an object that can be passed to `tableplot`. From this stage, tableplots are generated very fast, no matter on which column the data is sorted or how many row bins are chosen.

## Usage

```
tablePrepare(x, name = NULL, ...)
```

## Arguments

x	<code>data.frame</code> or <code>ffdf</code> , will be transformed into an <code>ffdf</code> object.
name	name of the dataset
...	at the moment not used

## Details

The function `bin_data` needs a prepared `data.frame`. `tablePrepare` transforms the supplied data into an `ffdf` object and calculates the order of each of its columns. Knowing the order of the columns speeds up the binning process considerably. For large `ffdf` objects this may be a time consuming step so it can be wise to call `tablePrepare` before making a `tableplot`.

## Value

a prepared object, including the data and order of each of the columns

**Examples**

```
# load diamonds dataset from ggplot2
require(ggplot2)
data(diamonds)

p <- tablePrepare(diamonds)

tableplot(p, nBins=200, sortCol=depth)
tableplot(p, nBins=50, sortCol=price)
```

---

tableSave

*Save a tableplot*


---

**Description**

Save a tableplot in pdf, eps, svg, wmf, png, jpg, bmp, or tiff format.

**Usage**

```
tableSave(tab, filename = paste(tab$dataset, ".pdf", sep = ""),
  device = default_device(filename), path = NULL, scale = 1,
  width = par("din")[1], height = par("din")[2], dpi = 300,
  onePage = TRUE, ...)
```

**Arguments**

tab	a <b>tabplot-object</b> , or a list of <b>tabplot-objects</b> , which are either stacked horizontally or put on multiple pages (for pdf only)
filename	filename with extension (pdf, eps, svg, wmf, png, jpg, bmp, or tiff)
device	device, automatically extracted from filename extension
path	path to save to
scale	scaling factor
width	width (in inches)
height	height (in inches)
dpi	dpi to use for raster graphics
onePage	if true, multiple tab objects are stacked horizontally, else they are printed on multiple pages
...	other arguments passed to <a href="#">plot</a> or the used graphics device



**Examples**

```
## Not run:
require(ggplot2)
data(diamonds)

# default tableplot
tab <- tableplot(diamonds)

# save tableplot
tableSave(tab, filename="diamonds.png", title="Shine on you Crazy Diamond!!!")

## End(Not run)
```

---

tabplot-object	<i>Object that contains the information to plot a tableplot</i>
----------------	---

---

**Description**

An object of class `tabplot` contains the information to plot a tableplot without the steps that may be time-consuming, such as sorting and aggregating. The function `tableplot` silently returns a `tabplot-object` (use `plot=FALSE` to suppress that the tableplot is plotted). The function `tableChange` can be used to change a `tabplot-object`. The generic functions `plot` and `summary` are used to plot and summarize a `tabplot-object`.

---

tabplot_compare-object	<i>Object that contains the information to plot the difference of two tableplots (experimental)</i>
------------------------	---

---

**Description**

A `tabplot_compare` is created by subtracting two tableplots (see `-.tabplot`). For numeric variables, both absolute and relative difference of the mean values are computed. For categorical variables, the frequency tables are compared.

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