

Package ‘colorBlindness’

March 16, 2020

Type Package

Title Safe Color Set for Color Blindness

Version 0.1.6

Description Provide the safe color set for color blindness, the simulator of protanopia, deuteranopia. The color sets are collected from: Wong, B. (2011) <doi:10.1038/nmeth.1618>, <<http://mkweb.bcgsc.ca/biovis2012/>>, and <http://geog.uoregon.edu/datagraphics/color_scales.htm>. The simulations of the appearance of the colors to color-deficient viewers were based on algorithms in Vienot, F., Brettel, H. and Mollon, J.D. (1999) <doi:10.1002/(SICI)1520-6378(199908)24:4%3C243::AID-COL5%3E3.0.CO;2-3>. The cvdPlot() function to generate 'ggplot' grobs of simulations were modified from <<https://github.com/clauswilke/colorblindr>>.

Depends R(>= 3.6)

Imports ggplot2, grDevices, methods, cowplot, colorspace, graphics, gridGraphics, gtable, grid

Suggests knitr, reshape2, stats, png

biocViews Visualization

License GPL (>= 2)

Encoding UTF-8

LazyData true

VignetteBuilder knitr

RoxygenNote 7.0.2

NeedsCompilation no

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Repository CRAN

Date/Publication 2020-03-16 12:10:05 UTC

R topics documented:

availableColors	2
availablePalette	3
BLACK	3
colorNames	4
cvdPlot	4
cvdSimulator	5
displayAvailablePalette	6
displayColors	6
grobify	7
paletteMartin	7
replacePlotColor	10
setPDFopt	10
Index	12

availableColors	<i>available colors</i>
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Description

export available colors

Usage

```
availableColors()
```

Value

a character vector contain safe colors.

Examples

```
availableColors()
```

availablePalette	<i>Available color palette</i>
------------------	--------------------------------

Description

List all the available color palettes.

Usage

```
availablePalette()
```

Value

a character vector contain available color palettes.

Examples

```
availablePalette()
```

BLACK	<i>safe colors</i>
-------	--------------------

Description

color blindness safe colors

Usage

```
BLACK
```

```
ORANGE
```

```
SKY_BLUE
```

```
BLUIISH_GREEN
```

```
YELLOW
```

```
BLUE
```

```
VERMILLION
```

```
REDDISH_PURPLE
```

```
safeColors
```

Format

An object of class character of length 1.

References

Wong, B. (2011) <doi:10.1038/nmeth.1642> Wong, B. (2011) <doi:10.1038/nmeth.1618>

Examples

```
safeColors
```

colorNames	<i>available color variable</i>
------------	---------------------------------

Description

export available color names

Usage

```
colorNames()
```

Value

a character vector contain safe colors.

Examples

```
colorNames()
```

cvdPlot	<i>Show color-deficiency simulations of a plot</i>
---------	--

Description

Plot the color-deficiency simulations for ggplot grob.

Usage

```
cvdPlot(
  plot = last_plot(),
  layout = c("origin", "deutanope", "protanope", "desaturate")
)
```

Arguments

plot The grob to be plotted.
layout The sub-figure types. the choices are origin, deuteranope, protanope, desaturate, and enhanced, enhanced.deuteranope, enhanced.protanope, enhanced.desaturate.

Details

This function is modified from <https://github.com/clauswilke/colorblindr>

Value

An object of ggplot.

Examples

```
cvdPlot(displayColors(safeColors))  
cvdPlot(displayColors(paletteMartin))
```

cvdSimulator *simulate color vision deficiency*

Description

Transformation of R colors by simulating color vision deficiencies.

Usage

```
cvdSimulator(col, type = "deuteranope")
```

Arguments

col character. A vector of colors.
type Deficiency type, "protanope" or "deuteranope"

Details

Here use Vienot's methods but not Gustavo's methods (implemented in `colorspace::simulate_cvd`).

Value

colors.

References

Vienot, F., Brettel, H. and Mollon, J.D. (1999) <doi:10.1002/(SICI)1520-6378(199908)24:4 Sharma, G., Wu, W. and Dalal, E.N. (2005) <doi:10.1002/col.20070>

Examples

```
cvdSimulator(safeColors)
```

```
displayAvailablePalette
```

Display available palette

Description

Display all the available color palettes.

Usage

```
displayAvailablePalette(...)
```

Arguments

... parameters could be used by [geom_tile](#).

Value

an [ggplot](#) object

Examples

```
displayAvailablePalette()
```

```
displayColors
```

display colors

Description

Display the given colors

Usage

```
displayColors(col, ...)
```

```
displayAllColors(col, types = c("deuteranope", "protanope", "desaturate"), ...)
```

Arguments

col color set to display

... parameters could be used by [geom_tile](#).

types the type of color vision deficiency.

Value

an `ggplot` object

Examples

```
displayColors(safeColors)
displayColors(paletteMartin)
displayAllColors(safeColors, color="white")
displayAllColors(paletteMartin, color="white")
```

<code>grobify</code>	<i>convert plot to grob</i>
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Description

use `grid.grabExpr` or `plot_to_gtable` to convert plot to grob

Usage

```
grobify(plot)
```

Arguments

`plot` `plots`

Value

grob object.

<code>paletteMartin</code>	<i>Palette for color blindness</i>
----------------------------	------------------------------------

Description

The palette could be used for heatmap or pie graph

Usage

`paletteMartin`

`Green2Magenta16Steps`

`Blue2DarkRed12Steps`

`Blue2DarkRed18Steps`

`Blue2OrangeRed14Steps`

`Blue2DarkOrange12Steps`

`Blue2DarkOrange18Steps`

`Blue2Green14Steps`

`Brown2Blue10Steps`

`Brown2Blue12Steps`

`Blue2Gray8Steps`

`Blue2Orange8Steps`

`Blue2Orange10Steps`

`Blue2Orange12Steps`

`ModifiedSpectralScheme11Steps`

`LightBlue2DarkBlue7Steps`

`LightBlue2DarkBlue10Steps`

`PairedColor12Steps`

`SteppedSequential5Steps`

Format

An object of class character of length 15.

Details

The names of the palette is approximal color name.

`Green2Magenta16Steps`: Useful for generic diverging data.

Blue2DarkRed12/18Steps: Useful for temperature-like data, with a subjective interpretation (blue=cold, red=hot) Blue2OrangeRed14Steps: Useful as an alternative to the red/blue temperature scale.

Blue2DarkOrange12/18Steps: Useful for data without a specific subjective color association.

Blue2Green14Steps: Useful for data with a winter (blue) vs. summer (green) association.

Brown2Blue10/12Steps: Useful for data with a dry (brown) vs. wet (blue) association.

Blue2Gray8Steps: Useful in particular for diverging data like cloudiness anomalies.

Blue2Orange8/10/12Steps: Useful for data like sea-level pressure, with an subjective association (blue=low, wet, orange=high, dry)

ModifiedSpectralScheme11Steps: An alternative to the spectral scheme (no green)

LightBlue2DarkBlue7/10Steps: Useful for precipitation-like data.

PairedColor12Steps: Attempt at a categorical color scale with colors that may be distinguishable to all viewers

SteppedSequential5Steps: Useful for portraying levels-within-categories

Source

<<http://mkweb.bcgsc.ca/biovis2012/>>

<http://geog.uoregon.edu/datagraphics/color_scales.htm>

References

Light A, Bartlein PJ (2004). "The End of the Rainbow? Color Schemes for Improved Data Graphics." EOS Transactions of the American Geophysical Union, 85(40), 385.

Examples

paletteMartin
 Green2Magenta16Steps
 Blue2DarkRed12Steps
 Blue2DarkRed18Steps
 Blue2OrangeRed14Steps
 Blue2DarkOrange12Steps
 Blue2DarkOrange18Steps
 Blue2Green14Steps
 Brown2Blue10Steps
 Brown2Blue12Steps
 Blue2Gray8Steps
 Blue2Orange8Steps
 Blue2Orange10Steps
 Blue2Orange12Steps
 ModifiedSpectralScheme11Steps
 LightBlue2DarkBlue7Steps
 LightBlue2DarkBlue10Steps
 PairedColor12Steps
 SteppedSequential15Steps

replacePlotColor	<i>replace the colors for plots</i>
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Description

replace the colors of plots to meet the requirement of publication. Replacing red with magenta or green with turquoise. Replacing all the colored symbols in the legend.

Usage

```
replacePlotColor(plot)
```

Arguments

plot	The grob to be plotted.
------	-------------------------

Value

an object of gtable.

Examples

```
replacePlotColor(displayColors(c("Red", "Green", "blue")))
```

setPDFopt	<i>Auxiliary function to set width of pdf for journals</i>
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Description

Set the pdf width and height for journals.
Pre-sets of width for figures.

Usage

```
setPDFopt(
  width = c("1col", "1.5col", "0.5col", "2col"),
  presets = PRESETS$science
)
```

PRESETS

Arguments

width	columns.
presets	The pre-setting of width,height,family,font for pdf. Available choices: 0.5col, 1col, 1.5col, 2col.

Format

An object of class `list` of length 4.

Details

The family will be Helvetica. The font will be 8. The width and height will be same.

science: 0.5col=1.78 inches (4.52 cm.); 1col=3.54 inches (9 cm.); 1.5col=5 inches (12.7 cm.); 2col=7.25 inches (18.4 cm.). nature: 0.5col=2.28 inches (5.8 cm.); 1col=3.39 inches (8.6 cm.); 1.5col=4.76 inches (12.1 cm.); 2col=7 inches (17.8 cm.). cell: 0.5col=1.78 inches (4.52 cm.); 1col=3.35 inches (8.5 cm.); 1.5col=4.49 inches (11.4 cm.); 2col=6.85 inches (17.4 cm.). CA: A Cancer Journal for Clinicians 0.5col=1.62 inches (4.1 cm.); 1col=3.25 inches (8.25 cm.); 1.5col=3.87 inches (9.8 cm.); 2col=6.75 inches (17.1 cm.).

Value

A named list of all the defaults. If any arguments are supplied the return values are the old values and the result has the visibility flag turned off.

References

<http://www.sciencemag.org/sites/default/files/Figure_prep_guide.pdf>

<<https://images.nature.com/full/nature-assets/aj/artworkguidelines.pdf>>

Examples

```
op <- setPDFopt("1col")
```

Index

*Topic **datasets**

- BLACK, 3
- paletteMartin, 7
- setPDFopt, 10

- availableColors, 2
- availablePalette, 3

- BLACK, 3
- BLUE (BLACK), 3
- Blue2DarkOrange12Steps (paletteMartin), 7
- Blue2DarkOrange18Steps (paletteMartin), 7
- Blue2DarkRed12Steps (paletteMartin), 7
- Blue2DarkRed18Steps (paletteMartin), 7
- Blue2Gray8Steps (paletteMartin), 7
- Blue2Green14Steps (paletteMartin), 7
- Blue2Orange10Steps (paletteMartin), 7
- Blue2Orange12Steps (paletteMartin), 7
- Blue2Orange8Steps (paletteMartin), 7
- Blue2OrangeRed14Steps (paletteMartin), 7
- BLUISH_GREEN (BLACK), 3
- Brown2Blue10Steps (paletteMartin), 7
- Brown2Blue12Steps (paletteMartin), 7

- colorNames, 4
- cvdPlot, 4
- cvdSimulator, 5

- displayAllColors (displayColors), 6
- displayAvailablePalette, 6
- displayColors, 6

- geom_tile, 6
- ggplot, 6, 7
- Green2Magenta16Steps (paletteMartin), 7
- grobify, 7

- LightBlue2DarkBlue10Steps (paletteMartin), 7
- LightBlue2DarkBlue7Steps (paletteMartin), 7
- ModifiedSpectralScheme11Steps (paletteMartin), 7

- ORANGE (BLACK), 3

- PairedColor12Steps (paletteMartin), 7
- paletteMartin, 7
- PRESETS (setPDFopt), 10

- REDDISH_PURPLE (BLACK), 3
- replacePlotColor, 10

- safeColors (BLACK), 3
- setPDFopt, 10
- SKY_BLUE (BLACK), 3
- SteppedSequential5Steps (paletteMartin), 7

- VERMILLION (BLACK), 3

- YELLOW (BLACK), 3