

Package ‘cols4all’

August 28, 2025

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Title Colors for all

Type Package

LazyLoad yes

Description Color palettes for all people, including those with color vision deficiency. Popular color palette series have been organized by type and have been scored on several properties such as color-blind-friendliness and fairness (i.e. do colors stand out equally?). Own palettes can also be loaded and analysed. Besides the common palette types (categorical, sequential, and diverging) it also includes cyclic and bivariate color palettes. Furthermore, a color for missing values is assigned to each palette.

Version 0.9

Encoding UTF-8

Depends R (>= 4.1.0),

Imports methods, grDevices, stats, abind, png, stringdist, colorspace
(>= 2.1), spacesXYZ

Suggests colorblindcheck, kableExtra, knitr, shiny, shinyjs, ggplot2,
scales, rmarkdown, bookdown, bibtex, plotly

URL <https://cols4all.github.io/cols4all-R/>,
<https://github.com/cols4all/cols4all-R>

BugReports <https://github.com/cols4all/cols4all-R/issues>

RoxygenNote 7.3.2

NeedsCompilation no

Author Martijn Tennekkes [aut, cre],
Marco Puts [ctb],
Achim Zeileis [ctb],
Jakub Nowosad [ctb],
Robin Lovelace [ctb],
Helgasoft [ctb],
Matthew Petroff [ctb],
Olivier Roy [ctb]

Maintainer Martijn Tennekes <mtennekes@gmail.com>
Repository CRAN
Date/Publication 2025-08-28 07:50:01 UTC

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cols4all-package	<i>cols4all overview</i>
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Description

cols4all stands for: color palettes for all people, including those with color vision deficiency. Popular color palette series, such as ColorBrewer, have been organized by type and have been scored on several properties such as color-blind-friendliness and fairness (i.e. do colors stand out equally?). Own palettes can also be loaded and analysed. Besides the common palette types (categorical, sequential, and diverging) it also includes bivariate color palettes. ggplot2 scales are included.

Details

This page provides a brief overview of all package functions.

Main functions

- [c4a_gui](#) Dashboard for analyzing the palettes
- [c4a](#) Get the colors from a palette ([c4a_na](#) for the associated color for missing values)
- [c4a_plot](#) Plot a color palette

Palette names and properties

<code>c4a_palettes</code>	Get available palette names
<code>c4a_series</code>	Get available series names
<code>c4a_overview</code>	Get an overview of palettes per series x type
<code>c4a_citation</code>	Show how to cite palettes (with bibtex code)
<code>c4a_info</code>	Get information from a palette, such as type and maximum number of colors
<code>.P</code>	Environment via which palette names can be browsed with auto-completion (using <code>\$</code>)

Importing and exporting palettes

<code>c4a_data</code>	Build color palette data
<code>c4a_load</code>	Load color palette data
<code>c4a_sysdata_import</code>	Import system data
<code>c4a_sysdata_export</code>	Export system data

Author(s)

Maintainer: Martijn Tennekes <mtennekes@gmail.com>

Other contributors:

- Marco Puts <mputs@acm.org> [contributor]
- Achim Zeileis <Achim.Zeileis@R-project.org> [contributor]
- Jakub Nowosad <nowosad.jakub@gmail.com> [contributor]
- Robin Lovelace <rob00x@gmail.com> [contributor]
- Helgasoft <contact@helgasoft.com> [contributor]
- Matthew Petroff <matthew@mpetroff.net> [contributor]
- Olivier Roy [contributor]

See Also

Useful links:

- <https://cols4all.github.io/cols4all-R/>
- <https://github.com/cols4all/cols4all-R>
- Report bugs at <https://github.com/cols4all/cols4all-R/issues>

c4a

*Get a cols4all color palette***Description**

Get a cols4all color palette: c4a returns the colors of the specified palette, c4a_na returns the color for missing value that is associated with the specified palette, and c4a_ramp returns a color ramp function. Run [c4a_gui](#) to see all available palettes, which are also listed with [c4a_palettes](#).

Usage

```
c4a(
  palette = NULL,
  n = NA,
  m = NA,
  type = c("cat", "seq", "div", "cyc", "bivs", "bivc", "bivd", "bivg"),
  reverse = FALSE,
  diag_flip = FALSE,
  order = NULL,
  range = NA,
  colorsort = "orig",
  format = c("hex", "rgb", "hcl", "RGB", "XYZ", "HSV", "HLS", "LAB", "polarLAB", "LUV",
    "polarLUV"),
  nm_invalid = c("error", "repeat", "interpolate"),
  verbose = TRUE
)

c4a_ramp(..., space = c("rgb", "Lab"), interpolate = c("linear", "spline"))

c4a_na(
  palette = NULL,
  type = c("cat", "seq", "div", "cyc", "bivs", "bivc", "bivd", "bivg"),
  verbose = TRUE
)
```

Arguments

palette	name of the palette. See c4a_palettes for available palettes. If omitted, the default palette is provided by c4a_default_palette. The palette name can be prefixed with a "-" symbol, which will reverse the palette (this can also be done with the reverse argument). For bivariate palettes, a "-" means reversed horizontally (columns), a " " means reversed vertically (row), and a "+" means reversed in both directions. In addition, a "/" or "\" will flip the palette diagonally. This can be used in combination with "-", " ", or "+". E.g. "-/" will reverse the columns and flip the palette diagonally.
---------	---

n	number of colors. If omitted then: for type "cat" the maximum number of colors is returned, for types "seq", "div", and "cyc", 7, 9, and 9 colors respectively. For bivariate palettes n is the number of columns.
m	number of rows in case type is bivariate, so one of "bivs", "bivc", "bivd" or "bivg" (see c4a_types for descriptions)
type	type of color palette, in case palette is not specified: one of "cat", "seq", "div", "cyc", "bivs", "bivc", "bivd", "bivg". Run c4a_types for descriptions.
reverse	should the palette be reversed? In case of a bivariate palette, a vector of two: the first indicates the horizontal direction (columns) and the second the vertical (rows).
diag_flip	should a bivariate palette be flipped diagonally?
order	order of colors. Only applicable for "cat" palettes
range	a vector of two numbers between 0 and 1 that determine the range that is used for sequential and diverging palettes. The first number determines where the palette begins, and the second number where it ends. For sequential "seq" palettes, 0 means the leftmost (normally lightest) color, and 1 the rightmost (often darkest) color. For diverging "seq" palettes, 0 means the middle color, and 1 both extremes. If only one number is provided, this number is interpreted as the endpoint (with 0 taken as the start).
colorsort	Sort the colors. Options: "orig" (original order), "Hx" (hue, where x is a starting number from 0 to 360), "C" (chroma), "L" (luminance). All these options are available for "cat" palettes, only the last one for "seq", and none for the other palette types.
format	format of the colors. One of: "hex" character vector of hex color values, "rgb" 3 column matrix of RGB values, "hcl" 3-column matrix of HCL values, or one of the color classes from colorspace
nm_invalid	what should be done in case n or m is larger than the maximum number of colors or smaller than the minimum number? Options are "error" (an error is returned), "repeat", the palette is repeated, "interpolate" colors are interpolated. For categorical "cat" palettes only.
verbose	should messages be printed?
...	passed on to c4a.
space	a character string; interpolation in RGB or CIE Lab color spaces
interpolate	use spline or linear interpolation

Value

A vector of colors (c4a) and a color (c4a_na)

Examples

```
# get the colors from brewer.set3 and plot them
set3 <- c4a("brewer.set3")
c4a_plot_hex(set3, nrow = 1)
```

```

c4a("hcl.set2", n = 36) |> c4a_plot_hex()
c4a("-hcl.set2", n = 12) |> c4a_plot_hex()

# how to know which palettes are available?
# 1) Via the interactive tool:
## Not run:
c4a_gui()

## End(Not run)

# 2) Via the overview function:
c4a_palettes(type = "cat")
c4a_palettes(series = "brewer")
c4a_palettes(type = "cat", series = "brewer")
# Run c4a_overview() to see which are available

# 3) Via .P
.P$brewer$cat$set3

# each palette contains a color for missing values

c4a("carto.safe", 7)
c4a_na("carto.safe")

c4a_plot_hex("carto.safe", n = 7, include.na = TRUE)

c4a_plot_hex("carto.safe", n = 7, include.na = TRUE)
# same (but shorter) as
# c4a_plot_hex(c(c4a("carto.safe", 7), c4a_na("carto.safe")), include.na = TRUE)

# color ramp
c4a("viridis", 100) |> c4a_plot()
c4a_ramp("viridis")(100) |> c4a_plot()

```

c4a_citation

Show how to cite palettes

Description

Show how to cite palettes

Usage

```
c4a_citation(name, verbose = TRUE)
```

Arguments

name	name of a palette or series
verbose	should text be printed (if FALSE only a <code>utils::bibentry</code> object is returned)

Value

`utils::bibentry` object

Examples

```
c4a_citation("hcl")

c4a_citation("poly.glasbey")
```

c4a_data	<i>Build and load palette data</i>
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Description

Build palette data. Both `c4a_data` and `c4a_data_as_is` build data palette. The difference is that the former may restructure the palette colors (see details) whereas the latter takes the palette colors as they are. Data can subsequently be loaded into `cols4all` via [c4a_load](#). The `c4a_data` function can also be used to read `c4a_info` objects, which contain data for a single palette.

Usage

```
c4a_data(
  x,
  xNA = NA,
  types = "cat",
  series = "x",
  nmin = NA,
  nmax = NA,
  ndef = NA,
  mmin = NA,
  mmax = NA,
  mdef = NA,
  format.palette.name = TRUE,
  remove.blacks = NA,
  remove.whites = NA,
  take.gray.for.NA = FALSE,
  remove.other.grays = FALSE,
  light.to.dark = FALSE,
  remove.names = TRUE,
  biv.method = "byrow",
  space = "rgb",
  range_matrix_args = list(NULL),
```

```

    bib = NA,
    description = NA
  )

c4a_load(data, overwrite = FALSE)

c4a_data_as_is(
  ...,
  format.palette.name = FALSE,
  remove.blacks = FALSE,
  remove.whites = FALSE,
  take.gray.for.NA = FALSE,
  remove.other.grays = FALSE,
  light.to.dark = FALSE,
  remove.names = FALSE
)
```

Arguments

<code>x</code>	either a named list of color palettes or a <code>c4a_info</code> object. For the first case: see details for indexing. The second case will bypass the other arguments.
<code>xNA</code>	colors for missing values. Vector of the same length as <code>x</code> (or length 1). For NA values, the color for missing values is automatically determined (preferable a light grayscale color, but if it is indistinguishable by color blind people, a light color with a low chroma value is selected)
<code>types</code>	character vector of the same length as <code>x</code> (or length 1), which determines the type of palette: "cat", "seq", "div", "cyc", "bivs", "bivc", "bivd", or "bivg". See details.
<code>series</code>	a character vector of the same length as <code>x</code> (or length 1), which determines the series.
<code>nmin, nmax, ndef</code>	minimum / maximum / default number of colors for the palette. By default: <code>nmin = 1</code> , for "cat" <code>nmax</code> and <code>ndef</code> the number of supplied colors. For the other types, <code>nmax</code> is Inf. <code>ndef</code> is 7 for "seq", 9. For diverging palettes, these numbers refer to the number of columns. (See <code>mmin</code> , <code>mmax</code> , <code>mdef</code> for the rows)
<code>mmin, mmax, mdef</code>	minimum / maximum / default number of rows for bivariate palettes.
<code>format.palette.name</code>	should palette names be formatted to lowercase/underscore format?
<code>remove.blacks, remove.whites, take.gray.for.NA, remove.other.grays</code>	These arguments determine the processing of grayscale colors for categorical "cat" palettes: if <code>remove.blacks</code> and there are (near) blacks, these are removed first. Next, if <code>take.gray.for.NA</code> , <code>xNA</code> is NA, and a palette contains at least one grayscale color (which can also be white), this is used as color for missing values. In case there are more than one grayscale color, the lightest is taken. <code>remove.other.grays</code> determines what happens with the other grays.
<code>light.to.dark</code>	should sequential "seq" palettes be automatically ordered from light to dark?
<code>remove.names</code>	should individual color names be removed?

biv.method	method to create a bivariate palette. Options are "byrow" means that the colors are wrapped row-wise to a color matrix where the number of rows and columns is automatically determined, "byrowX" the same but with X (integer between 2 and 9) columns, "bycol" and "bycolX" similar but wrapped column-wise. "div2seqseq" and "div2catseq" means that colors are extracted from a diverging palette. The former translates colors into a matrix with the neutral color in the diagonal, while the latter places the neutral color in the middle column. "seq2uncseq"
space	color space in which interpolated colors are determined. Options: "rgb" (RGB) and "Lab" (CIE Lab).
range_matrix_args	list of lists, one for each palette. Each such list specifies the range of sequential and diverging palettes, in case they are not indexed. See details.
bib	bibtex reference in the form of a <code>utils::bibentry</code> object.
description	description of the series. If series contains multiple series (rather than one value), please specify a vector of the same length as series. See c4a_series for the descriptions of the currently loaded series.
data	cols4all data created with <code>c4a_data</code>
overwrite	in case the palettes already exist (i.e. the full names), should the old names be overwritten?
...	passed on to <code>c4a_data</code>

Details

In `cols4all`, palettes are organized by series and by type. The **series** or 'family' specifies where the palettes belong to. For instance "brewer" stands for the color palettes from ColorBrewer. Run [c4a_series](#) to get an overview of loaded series. The **type** specifies what kind of palette it is; see [c4a_types](#) for a description of the implemented ones.

This function structures the palette data, such that it is consistent with the other palette data. This includes:

- Palette names are made consistent. We use the convention "my_series.my_palette", so all lower case, a period to separate the series name from the palette name, and underscores to separate words.
- (Only for `c4a_data`, bypassed for `c4a_data_as_is`) Categorical palettes: black is removed from categorical palettes, and a grayscale color is assigned to be used for missing values (other grayscale colors are removed). Sequential palettes are sorted from light to dark.

Indexing: for a categorical "cat" palette, an optional "index" attribute determines which colors to use for which lengths: if the palette consists of k colors, index should be a list of k , where the i -th element is an integer vector of length i with values $1, 2, \dots, k$. See `c4a_info("rainbow")` and for an example.

Range: sequential and diverging palettes are usually defined for 9+ colors. The optional "range_matrix" attribute determines that range is used for less colors. It is a $n \times 2$ matrix where row i defines the applied range of a palette of length i . For sequential palettes a range $c(0, 1)$ means that the palette is generated (via a color ramp) between the two outermost colors. For diverging palettes, a range

`c(x, y)` means that both sides of the palette are generated (via a color ramp) from `x`, which is the distance to the center color, to `y` which represents both outermost colors.

The range is automatically set for sequential and diverging palettes that have no "index" or "range_matrix" attribute via the parameter `range_matrix_args`, which is a list per palette. The arguments for a sequential palette are: `nmin` the minimum number of colors for which the range is reduced, `nmax`, the number of colors for which the range is set to `c(0,1)`, `slope_min` and `slope_max` determine the slopes of range reduction from a palette of length `nmax` to `nmin`, and `space` sets the color space for which the color ramp is applied ("rgb" or "Lab"). The arguments for a diverging palette are the same, but only one slope is used (namely for the outermost colors).

It may take some time to process, especially large categorical palettes, because of calculations of the color blind checks.

Value

`c4a_data` object, which is a list of four items: `data`, `s`, `citation`, and `description`

Examples

```
# palettes extracted Pink Floyd albums
pf = list(piper = c("#391C1C", "#C6C6AA", "#713939", "#C6391C",
  "#C6E3C6", "#AA7155", "#AA8E71", "#C68E71"),
  saucerful = c("#000000", "#1C1C1C", "#393939", "#FFFFFF",
  "#555555", "#8E8E71", "#E3C6AA", "#715539"),
  atom = c("#C6E3FF", "#397139", "#557139", "#E3E3C6",
  "#1C1C1C", "#1C551C", "#AAAA8E", "#8EC6E3"),
  meddle = c("#715539", "#553939", "#8E7155", "#71AAAA",
  "#8E8E71", "#1CAAE3", "#55C6E3", "#AA7155"),
  obscured = c("#000000", "#1C1C1C", "#393939", "#717155",
  "#8E8E71", "#715539", "#C6AA8E", "#E3C6AA"),
  moon = c("#000000", "#FF0000", "#FF9224", "#FFFF00",
  "#71C600", "#00C6FF", "#8E398E", "#FFFFFF"),
  wish = c("#FFFFFF", "#AAC6E3", "#8E8E8E", "#717155",
  "#555539", "#8E8E71", "#555555", "#8E7155"),
  animals = c("#391C39", "#393955", "#E3C671", "#718E8E",
  "#AAAA8E", "#C67139", "#AA5539", "#E3AA39"),
  wall = c("#FFFFFF", "#E3E3E3", "#C6C6C6", "#AAAC6",
  "#1C1C1C", "#000000", "#8E8E8E", "#E3C6E3"),
  cut = c("#000000", "#E30000", "#AA0000", "#391C55",
  "#FFE3E3", "#1C1C00", "#FFAA55", "#8E8E55"),
  lapse = c("#000000", "#8E8EC6", "#8E8E71", "#7171AA",
  "#39391C", "#717171", "#AAAAAA", "#E3E3E3"),
  division = c("#000000", "#FFFC6", "#00398E", "#AA8E55",
  "#39558E", "#C6AA71", "#39391C", "#555571"),
  more = c("#0055AA", "#FFAA1C", "#1C71AA", "#003971",
  "#E38E55", "#E3AAAA", "#718EAA", "#71718E"),
  umma = c("#AA8E71", "#555539", "#39391C", "#1C1C1C",
  "#E3E3C6", "#715539", "#391C1C", "#8E7155"),
  relics = c("#3955AA", "#1C3971", "#5571C6", "#715555",
  "#8E7155", "#E3AA71", "#8E8EAA", "#E3FFFF"),
  river = c("#393939", "#555555", "#39558E", "#C6C6C6",
  "#718EAA", "#1C1C1C", "#717171", "#E3C68E"))
```

```

if (requireNamespace("colorblindcheck", quietly = TRUE)) {
  pfdata = c4a_data_as_is(pf, series = "pinkfloyd",
    description = "Palettes extracted from Pink Floyd album covers")
  c4a_load(pfdata)

  c4a_series()
  c4a_overview()

  if (requireNamespace("shiny") &&
    requireNamespace("shinyjs") &&
    requireNamespace("kableExtra") &&
    requireNamespace("colorblindcheck") &&
    requireNamespace("plotly") &&
    interactive()) {
    c4a_gui(series = "pinkfloyd", n = 8)
  }
}

```

c4a_gui

Graphical user interface to analyse palettes

Description

Graphical user interface to analyse palettes. `c4a_table` shows a table that can be opened in the browser. `c4a_gui` is a graphical user interface (shiny app) around this table.

Usage

```

c4a_gui(type = "cat", n = NA, series = "all")

c4a_table(
  type = c("cat", "seq", "div", "cyc", "bivs", "bivc", "bivd", "bivg"),
  n = NULL,
  m = NULL,
  continuous = FALSE,
  filters = character(0),
  cvd.sim = c("none", "deutan", "protan", "tritan"),
  sort = "name",
  text.format = "hex",
  text.col = "same",
  series = "all",
  range = NA,
  colorsort = "orig",
  include.na = FALSE,
  show.scores = FALSE,
  columns = NA,
  verbose = TRUE
)

```

Arguments

type	type of palette. Run c4a_types to see the implemented types and their description. For <code>c4a_gui</code> it only determines which type is shown initially.
n, m	n is the number of displayed colors. For bivariate palettes "biv", n and m are the number of columns and rows respectively. If omitted: for "cat" the full palette is displayed, for "seq", "div" and "cyc", 7, 9, and 9 colors respectively, and for "bivs"/"bivc"/"bivd"/"bivg" 4 columns and rows. For <code>c4a_gui</code> it only determines which number of colors initially.
series	Series of palettes to show. See c4a_series for options. By default, "all", which means all series. For <code>c4a_gui</code> it only determines which series are shown initially.
continuous	should the palettes as continuous instead of discrete. Only applicable for "seq", "div", and "cyc".
filters	filters to be applied. A character vector with a subset from: "nmax" (only palettes where n = nmax, which is only applicable for categorical palettes), "cbf" (colorblind-friendly), "fair" (fairness), "naming" (nameability), "crW" (sufficient contrast ratio with white), and "crB" (sufficient contrast ratio with black). By default an empty vector, so no filters are applied.
cvd.sim	color vision deficiency simulation: one of "none", "deutan", "protan", "tritan"
sort	column name to sort the data. The available column names depend on the arguments type and show.scores. They are listed in the warning message. Use a "-" prefix to reverse the order.
text.format	The format of the text of the colors. One of "hex", "RGB" or "HCL".
text.col	The text color of the colors. By default "same", which means that they are the same as the colors themselves (so invisible, but available for selection). "auto" means automatic: black for light colors and white for dark colors.
range	vector of two numbers that determine the range that is used for sequential and diverging palettes. Both numbers should be between 0 and 1. The first number determines where the palette begins, and the second number where it ends. For sequential palettes, 0 means the leftmost (normally lightest) color, and 1 the rightmost (often darkest) color. For diverging palettes, 0 means the middle color, and 1 both extremes. If only one number is provided, this number is interpreted as the endpoint (with 0 taken as the start). By default, it is set automatically, based on n.
colorsort	Sort the colors ("cat" only). Options: "orig" (original order), "Hx" (hue, where x is a starting number from 0 to 360), "C" (chroma), "L" (luminance)
include.na	should color for missing values be shown? FALSE by default
show.scores	should scores of the quality indicators be printed? See details for a description of those indicators.
columns	number of columns. By default equal to n or, if not specified, 12. Cannot be higher than the palette lengths.
verbose	should messages and warnings be printed?

Details

See vignette how the properties are calculated. Parameters, such as threshold values which determined when palettes are classified as "colorblind-friendly", can be specified via [c4a_options](#). Also the nameability score function (which is in development) can be specified there. See the examples of [c4a_options](#) for both use cases.

Value

An HMTL table (kableExtra object)

See Also

References of the palettes: [cols4all-package](#).

Examples

```
if (requireNamespace("shiny") &&
    requireNamespace("shinyjs") &&
    requireNamespace("kableExtra") &&
    requireNamespace("colorblindcheck") &&
    interactive()) {

  c4a_gui()

  # categorical palettes with maximum number of colors
  c4a_table(type = "cat")

  # sort sequential palettes by hue
  c4a_table(type = "seq", n = 7, sort = "H")

  # sort sequential palettes by hue type (how many hues are used)
  c4a_table(type = "seq", n = 5, sort = "hues")
}
```

c4a_info

Get information from a cols4all palette

Description

Get information from a cols4all palette

Usage

```
c4a_info(palette, no.match = c("message", "error", "null"), verbose = TRUE)
```

Arguments

palette	name of the palette
no.match	what happens is no match is found? Options: "message": a message is thrown with suggestions, "error": an error is thrown, "null": NULL is returned
verbose	should messages be printed?

Value

list with the following items: name, series, fullname, type, palette (colors), na (color), nmax, and reverse. The latter is TRUE when there is a "-" prefix before the palette name.

c4a_modify

Edit cols4all palettes (in development)

Description

Edit cols4all palettes. c4a_duplicate duplicates an existing cols4all palette, and c4a_modify is used to change the colors. Use c4a_data to craete palettes from scratch.

Usage

```
c4a_modify(palette, x = NULL, xNA = NULL)
```

```
c4a_duplicate(palette, name = NA)
```

Arguments

palette	name of the palette
x	vector of the new colors. It should either the same length, or a named vector, where the names correspond to the index numbers. E.g. c("3" = "#AABBCC") will replace the third color with the color "#AABBCC".
xNA	the new color for missing values.
name	name of new palette

See Also

[c4a_data\(\)](#)

Examples

```
c4a_duplicate("brewer.set2", "set2_mod")
c4a_modify("set2_mod", c("4" = "#EA8AB8"))
```

c4a_options

*Set cols4all options***Description**

Get or set global options for c4a. Works similar as the base function options

Usage

```
c4a_options(...)
```

Arguments

... Use character values to retrieve options. To set options, either use named arguments (where the names refer to the options), a list that consists of those options.

Details**Option****Description**

defaults

Default palettes per type

CBF_th

Parameters that label a palette as color blind friendly

CBVF_th

Parameters that label a palette as very color blind friendly

CBU_th

Parameters that label a palette as color blind unfriendly

CrangeFair

Maximum chroma range for which a palette is considered harmonic

CrangeUnfair

Minimum chroma range for which a palette is considered disharmonic

LrangeFair

Maximum luminance range for which a palette is considered harmonic

LrangeUnfair

Minimum luminance range for which a palette is considered disharmonic

Cintense

Chroma of colors that are considered intense

Cpastel

Chroma of colors that are considered 'pastel'

HwidthDivRainbow

A diverging palette is labeled as 'rainbow hue' if HwidthL or HwidthR are at least HwidthDivRainbow

HwidthDivSingle

A diverging palette is labeled as 'single hue' if HwidthL and HwidthR are at most HwidthDivSingle

HwidthSeqRainbow

A sequential palette is labeled as 'rainbow hue' if Hwidth is at least HwidthSeqRainbow

HwidthSeqSingle

A sequential palette is labeled as 'single hue' if Hwidth is at most HwidthSeqSingle

naming_fun

Function that returns a distance matrix with the naming_colors (see examples)

naming_fun_args

List of arguments for naming_fun

naming_colors

Vector of prototype colors for the color names (see examples)

naming_softmax

List of parameters for the softmax function applied to the distance matrix

Value

A list of options

Examples

```

# Example how to lower the color-blind friendly threshold
# for categorical palettes (so more smileys in the GUI!)
# CBF_th: one smiley
# CBVF_th: two smileys

# current table
## Not run:
c4a_table(n = 9, sort = "cbfriendly")

opts = c4a_options("CBF_th", "CBVF_th")
opts$CBF_th$cat["min_dist"] = 7
opts$CBVF_th$cat["min_dist"] = 10

old = c4a_options(opts)

# more smileys :-) :-)
c4a_table(n = 9, sort = "cbfriendly")

# set the old settings back
c4a_options(old)

## End(Not run)

# Example how to use own nameability function
#
# This function should:
# - have an argument "pal" (vector of colors)
# - optionally have other arguments
# - return a distance matrix of n rows (length of pal) and k columns (classes).
#   It should have column names that correspond to the naming colors (see below).
naming_RGB = function(pal) {
  cols = colorspace::hex2RGB(pal)
  coords = cols@coords

  cls = apply(coords, MARGIN = 1, which.max)
  mx = apply(coords, MARGIN = 1, max)
  dominance = ((mx + 0.001) / (rowSums(coords) + 0.001))
  cls[dominance < 0.4] = 4L

  m = matrix(0, nrow = length(pal), ncol = 4,
    dimnames = list(NULL, c("Red", "Green", "Blue", "Other")))
  for (i in 1:nrow(m)) {
    m[i, cls[i]] = 1
  }

  -m
}

# testing this function...
naming_RGB(c4a("brewer.set1")) #fair enough

```



```
# This vector should contain the 'prototype' colors, and have names that correspond
# to the column names of the returned matrices by the function above.
names_RGB =
  c("Red" = "#FF0000",
    "Green" = "#00FF00",
    "Blue" = "#0000FF",
    "Other" = "#AAAAAA")

# Set the options (may take a while because if calculated the nameability scores)
## Not run:
c4a_options(naming_fun = naming_RGB,
naming_fun_args = list(),
naming_colors = names_RGB)

## End(Not run)
```

c4a_palettes

Get available palette names and series

Description

c4a_palettes lists all available cols4all color palettes. Palettes are organized by series. The available series are listed with c4a_series. Palettes are also organized per functional type, where we currently support: categorical "cat", sequential "seq", diverging "div", cyclic "cyc", and bivariate (seq x seq "bivs", seq x cat "bivc", seq x div "bivd", seq x desaturated "bivg") palette types. The function c4a_types lists all available types. The function c4a_overview gives an overview table of the number of palette per series and type. In an IDE with auto-completion (such as RStudio) it is possible to browse through the palette names with .P (using \$ like in lists).

Usage

```
c4a_palettes(
  type = c("all", "cat", "seq", "div", "cyc", "bivs", "bivc", "bivd", "bivg"),
  series = NULL,
  full.names = TRUE
)

c4a_series(type = c("all", "cat", "seq", "div", "cyc"), as.data.frame = TRUE)

c4a_types(series = NULL, as.data.frame = TRUE)

c4a_overview(return.matrix = FALSE, zero.count.as.NA = FALSE)

.P
```

Arguments

<code>type</code>	type of color palette: one of "all" (all palettes), "cat", "seq", "div", "cyc", "bivs", "bivc", "bivd", or "bivg". See c4a_types for descriptions.
<code>series</code>	series to list the palettes from. Run <code>c4a_series</code> to see the options.
<code>full.names</code>	should full names, i.e. with the prefix "series."? By default TRUE.
<code>as.data.frame</code>	should <code>c4a_series</code> and <code>c4a_types</code> return the result as a data.frame, with description included as a column?
<code>return.matrix</code>	should only a matrix be returned with numbers per palette and type? If FALSE a data.frame is returned with additional information
<code>zero.count.as.NA</code>	should zeros counted in the table be returned as 0 (FALSE, default) or as NA (TRUE)?

Format

An object of class environment of length 20.

Value

names of the loaded color palettes

See Also

References of the palettes: [cols4all-package](#).

Examples

```
c4a_series()

c4a_types()

c4a_overview()

c4a_palettes(type = "cat", series = "tol")

c4a_palettes(type = "seq", series = "kovesi")

# handy when auto-completion is available:
.P$kovesi$seq$linear_terrain
```

c4a_plot	<i>Plot a color palette</i>
----------	-----------------------------

Description

Plot a color palette, either a cols4all palette, or a color vector. `c4a_plot_cvd` is a shortcut to include color-blind simulated colors, `c4a_plot_hex` is a shortcut to print hex codes instead of labels.

Usage

```
c4a_plot(
  palette,
  ...,
  dark = FALSE,
  include.na = FALSE,
  hex = FALSE,
  include.cvd = FALSE,
  nrows = NA,
  ncols = NA
)

c4a_plot_cvd(...)

c4a_plot_hex(...)
```

Arguments

<code>palette</code>	Palette name (see c4a) or a color vector
<code>...</code>	arguments passed on to c4a
<code>dark</code>	dark (black) background?
<code>include.na</code>	should a color for missing values be included?
<code>hex</code>	should hex codes be printed instead of color labels (or numbers)?
<code>include.cvd</code>	should color deficiency simulated colors be included?
<code>nrows, ncols</code>	Number of rows and columns. Ignored if <code>include.cvd = TRUE</code> (in that case, rows are used for the simulated colors). By default automatically calculated based on aspect ratio of the device.

Value

Besides the plot, a [gTree](#) is returned silently

Examples

```

c4a_plot("brewer.set1", nrow=1)

c4a_plot_hex("brewer.set1", nrow=1)

c4a_plot_cvd("brewer.set1")

c4a_plot_cvd("greens")

c4a_plot_cvd("tol.pu_gn")

c4a_plot(.P$cols4all$bivs$pu_gn_bivs, n = 5)

c4a_plot(.P$met$bivc$monet)

c4a_plot(.P$cols4all$bivd$pu_gn_bivd, n = 5)

c4a_plot(.P$cols4all$bivg$gn_bivg, n = 5)

```

c4a_scores

Get information from a cols4all palette

Description

Get information from a cols4all palette

Usage

```

c4a_scores(
  palette = NULL,
  type = NULL,
  series = NULL,
  n = NA,
  no.match = c("message", "error", "null"),
  verbose = TRUE
)

```

Arguments

palette	name of the palette
type	type of palettes (in case palette is not specified)
series	series name (in case palette is not specified)
n	number of colors
no.match	what happens if no match is found? Options: "message": a message is thrown with suggestions, "error": an error is thrown, "null": NULL is returned
verbose	should messages be printed?

Value

list with the following items: name, series, fullname, type, palette (colors), na (color), nmax, and reverse. The latter is TRUE when there is a "-" prefix before the palette name.

Examples

```
c4a_scores("blues3")

pals = c4a_palettes(type = "cat")
scores_cat7 = t(sapply(pals, c4a_scores, n = 7))

head(scores_cat7)
```

c4a_sysdata_import	<i>Import and export system data</i>
--------------------	--------------------------------------

Description

Import and export system data. `c4a_sysdata_import` will import system data and overwrite the current system data, `c4a_sysdata_export` will export the current system data, and `c4a_sysdata_remove` (partly) removes system data.

Usage

```
c4a_sysdata_import(data)

c4a_sysdata_export()

c4a_sysdata_remove(fullnames = NULL, series = NULL, are.you.sure = NA)
```

Arguments

<code>data</code>	cols4all data (see <code>c4a_data</code>)
<code>fullnames</code>	full palette names (so in the format <code>series.palette_name</code>)
<code>series</code>	a character vector of series names that should be removed (use "all" to remove all).
<code>are.you.sure</code>	are you sure you want to remove series?

Value

`c4a_sysdata_export` returns the system data (a list)

Examples

```
x = c4a_sysdata_export()
c4a_sysdata_import(x)
y = c4a_sysdata_export()
identical(x, y)
```

```
scale_color_discrete_c4a_cat
```

col4all scales for ggplot2

Description

col4all scales for ggplot2. The scale functions are organized as `scale_<aesthetic>_<mapping>_c4a_<type>`, where the `<aesthetic>` should be either `colo(u)r` or `fill`, `<mapping>` refers to the mapping that is applied (discrete, continuous or binned), and `<type>` is the palette type: `cat`, `seq`, or `div`.

Usage

```
scale_color_discrete_c4a_cat(  
  palette = NULL,  
  reverse = FALSE,  
  order = NULL,  
  ...  
)  
  
scale_colour_discrete_c4a_cat(  
  palette = NULL,  
  reverse = FALSE,  
  order = NULL,  
  ...  
)  
  
scale_fill_discrete_c4a_cat(palette = NULL, reverse = FALSE, order = NULL, ...)  
  
scale_color_discrete_c4a_seq(  
  palette = NULL,  
  reverse = FALSE,  
  range = NULL,  
  ...  
)  
  
scale_colour_discrete_c4a_seq(  
  palette = NULL,  
  reverse = FALSE,  
  range = NULL,  
  ...  
)  
  
scale_fill_discrete_c4a_seq(palette = NULL, reverse = FALSE, range = NULL, ...)  
  
scale_color_discrete_c4a_div(  
  palette = NULL,  
  reverse = FALSE,
```

```
    range = NULL,  
    ...  
  )  
  
scale_colour_discrete_c4a_div(  
  palette = NULL,  
  reverse = FALSE,  
  range = NULL,  
  ...  
)  
  
scale_fill_discrete_c4a_div(palette = NULL, reverse = FALSE, range = NULL, ...)  
  
scale_color_continuous_c4a_seq(  
  palette = NULL,  
  reverse = FALSE,  
  range = NULL,  
  mid = 0,  
  n_interp = 11,  
  ...  
)  
  
scale_colour_continuous_c4a_seq(  
  palette = NULL,  
  reverse = FALSE,  
  range = NULL,  
  mid = 0,  
  n_interp = 11,  
  ...  
)  
  
scale_fill_continuous_c4a_seq(  
  palette = NULL,  
  reverse = FALSE,  
  range = NULL,  
  mid = 0,  
  n_interp = 11,  
  ...  
)  
  
scale_color_continuous_c4a_div(  
  palette = NULL,  
  reverse = FALSE,  
  range = NULL,  
  mid = 0,  
  n_interp = 11,  
  ...  
)
```

```
scale_colour_continuous_c4a_div(  
  palette = NULL,  
  reverse = FALSE,  
  range = NULL,  
  mid = 0,  
  n_interp = 11,  
  ...  
)
```

```
scale_fill_continuous_c4a_div(  
  palette = NULL,  
  reverse = FALSE,  
  range = NULL,  
  mid = 0,  
  n_interp = 11,  
  ...  
)
```

```
scale_color_binned_c4a_seq(  
  palette = NULL,  
  reverse = FALSE,  
  range = NULL,  
  mid = 0,  
  n_interp = 11,  
  ...  
)
```

```
scale_colour_binned_c4a_seq(  
  palette = NULL,  
  reverse = FALSE,  
  range = NULL,  
  mid = 0,  
  n_interp = 11,  
  ...  
)
```

```
scale_fill_binned_c4a_seq(  
  palette = NULL,  
  reverse = FALSE,  
  range = NULL,  
  mid = 0,  
  n_interp = 11,  
  ...  
)
```

```
scale_color_binned_c4a_div(  
  palette = NULL,
```



```

    reverse = FALSE,
    range = NULL,
    mid = 0,
    n_interp = 11,
    ...
  )

scale_colour_binned_c4a_div(
  palette = NULL,
  reverse = FALSE,
  range = NULL,
  mid = 0,
  n_interp = 11,
  ...
)

scale_fill_binned_c4a_div(
  palette = NULL,
  reverse = FALSE,
  range = NULL,
  mid = 0,
  n_interp = 11,
  ...
)

```

Arguments

palette, reverse, order, range	See c4a .
...	parameters passed on to the underlying scale functions: discrete_scale , continuous_scale , and binned_scale .
mid	data value that should be mapped to the mid-point of the diverging color scale. By default 0, which is useful for many use cases. However, if the data has only positive (or only negative) values, it may be worthwhile to specify this. Use NA to set it to the middle of the value range.
n_interp	number of discrete colors that should be used to interpolate the continuous color scale. Recommended to use an odd number to include the midpoint

Value

A ggplot2 component that defines the scale

Examples

```

if (require("ggplot2")) {
  data("diamonds")
  diam_exp = diamonds[diamonds$price >= 15000, ]
  diam_exp$clarity[1:500] = NA
}

```

```
# discrete categorical scale
ggplot(diam_exp, aes(x = carat, y = price, color = color)) +
  geom_point(size = 2) +
  scale_color_discrete_c4a_cat("carto.safe") +
  theme_light()

# missing values
c4a_plot("tol.muted", 8)
ggplot(diam_exp, aes(x = carat, y = price, fill = clarity)) +
  geom_point(size = 2, shape = 21) +
  scale_fill_discrete_c4a_cat("tol.muted") +
  theme_light()

# discrete sequential scale
ggplot(diam_exp, aes(x = carat, y = price, color = cut)) +
  geom_point(size = 2) +
  scale_color_discrete_c4a_seq("hcl.blues2") +
  theme_light()

# continuous sequential scale
ggplot(diam_exp, aes(x = carat, y = price, color = depth)) +
  geom_point(size = 2) +
  scale_color_continuous_c4a_seq("hcl.blues2", range = c(0.4, 1)) +
  theme_light()

# continuous diverging scale
ggplot(diam_exp, aes(x = carat, y = depth, color = price)) +
  geom_point(size = 2) +
  scale_color_continuous_c4a_div("wes.zissou1", mid = NA) +
  theme_light()

# binned sequential scale
ggplot(diam_exp, aes(x = carat, y = price, color = depth)) +
  geom_point(size = 2) +
  scale_color_binned_c4a_seq("scico.batlow", range = c(0.4, 1)) +
  theme_light()
}
```

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