

# Package: cols4all (via r-universe)

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**License** GPL-3

**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.

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**URL** <https://mtennekes.github.io/cols4all/>,  
<https://github.com/mtennekes/cols4all>

**BugReports** <https://github.com/mtennekes/cols4all/issues>

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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

<a href="#">c4a_gui</a>	Dashboard for analyzing the palettes
<a href="#">c4a</a>	Get the colors from a palette ( <a href="#">c4a_na</a> for the associated color for missing values)
<a href="#">c4a_plot</a>	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 \$)

### 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

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### See Also

Useful links:

- <https://mtennekes.github.io/cols4all/>
- <https://github.com/mtennekes/cols4all>
- Report bugs at <https://github.com/mtennekes/cols4all/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,
  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**

<code>palette</code>	name of the palette. See <code>c4a_palettes</code> for available palettes. If omitted, the default palette is provided by <code>c4a_default_palette</code> . The palette name can be prefixed with a "-" symbol, which will reverse the palette (this can also be done with the <code>reverse</code> argument).
<code>n</code>	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.
<code>m</code>	number of rows in case type is bivariate, so one of "bivs", "bivc", "bivd" or "bivg" (see <code>c4a_types</code> for descriptions)

type	type of color palette, in case palette is not specified: one of "cat", "seq", "div", "cyc", "bivs", "bivc", "bivd", "bivg". Run <a href="#">c4a_types</a> for descriptions.
reverse	should the palette be reversed?
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).
colsort	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 <a href="#">colorspace</a>
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, nrows = 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

*Build and load palette data*

---

## 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, mmax, 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?
<code>biv.method</code>	method to a create 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 <code>series</code> contains multiple series (rather than one value), please specify a vector of the same length as <code>series</code> . See <a href="#">c4a_series</a> 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", "#1CAA E3", "#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", "#FFFFFFC6", "#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(),
  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 <a href="#">c4a_types</a> 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 <a href="#">c4a_series</a> 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 <code>type</code> and <code>show.scores</code> . 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 HTML 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	<i>Edit cols4all palettes (in development)</i>
------------	--

---

**Description**

Edit cols4all palettes. `c4a_duplicate` duplicates an existing cols4all palette, and `c4a_modify` is used to change the colors. Use `c4a_data` to create 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 be the same length, or a named vector, where the names correspond to the index numbers. E.g. <code>c("3" = "#AABCC")</code> will replace the third color with the color <code>"#AABCC"</code> .
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	<i>Set cols4all options</i>
-------------	-----------------------------

---

**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**

<b>Option</b>	<b>Description</b>
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

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

### Format

An object of class environment of length 17.

**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

*Plot a color palette*


---

**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**

palette	Palette name (see <a href="#">c4a</a> ) or a color vector
...	arguments passed on to <a href="#">c4a</a>
dark	dark (black) background?
include.na	should a color for missing values be included?
hex	should hex codes be printed instead of color labels (or numbers)?
include.cvd	should color deficiency simulated colors be included?
nrows, ncols	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", nrows=1)

c4a_plot_hex("brewer.set1", nrows=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 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.

**Examples**

```
c4a_scores("blues3")

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

head(scores_cat7)
```

---

c4a\_sysdata\_import      *Import and export system data*

---

**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**

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

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 = mean(diam_exp$price)) +  
  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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