Package ‘cols4all’

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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 analyzed. Besides the common palette types (categorical, sequential, and diverging) it also includes bivariate color palettes. Furthermore, a color for missing values is assigned to each palette.

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Author Martijn Tennekes [aut, cre],
Marco Puts [ctb],
Achim Zeileis [ctb],
Jakub Nowosad [ctb],
Robin Lovelace [ctb],
Helgasoft [ctb],
Matthew Petroff [ctb]
**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**
c4a

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

**Importing and exporting palettes**

- `c4a_data`: Build color palette data
- `c4a_load`: Load color palette data
- `c4a_sysdata_import`: Import system data
- `c4a_sysdata_export`: Export system data

**Author(s)**

Martijn Tennekes <mtennekes@gmail.com>

**See Also**

https://github.com/mtennekes/cols4all

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**c4a**

*Get a cols4all color palette*

**Description**

Get a cols4all color palette: `c4a` returns the colors of the specified palette, and `c4a_na` returns the color for missing value that is associated with the specified palette. Run `c4a_gui` to see all available palettes, which are also listed with `c4a_palettes`.

**Usage**

```r
library(cols4all)

c4a(
  palette = NULL,
  n = NA,
  m = NA,
  type = c("cat", "seq", "div", "bivs", "bivc", "bivd", "bivg"),
  reverse = FALSE,
  order = NULL,
)
```

range = NA,
format = c("hex", "RGB", "HCL"),
nm_invalid = c("error", "repeat", "interpolate"),
verbose = TRUE
)
c4a_na(palette = NULL, type = c("cat", "seq", "div"), verbose = TRUE)

Arguments

palette name of the palette. See c4a_palettes for options. 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).

n number of colors. If omitted then: for type "cat" the maximum number of colors is returned, for types "seq" and "div", 9 colors.

m number of rows in case type is "bivs", "bivc", "bivd" or "bivg" (which stand for respectively sequential, categorical, diverging and desaturated (g for 'grayscale')).

type type of color palette, in case palette is not specified: one of "cat" (categorical/qualitative palette), "seq" (sequential palette), "div" (diverging palette), and "bivs"/"bivc"/"bivd"/"bivg" (bivariate: respectively seq-seq seq-cat, seq-div, and seq-desaturated).

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).

format format of the colors. One of: "hex" character vector of hex color values, "RGB" 3 column matrix of RGB values, or "HCL" 3-column matrix of HCL values

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?

Value

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

c4a_palettes("div")

c4a(type = "cat")

(pal = c4a("tol.sunset", n = 7, range = c(0, .6)))

c4a_plot(pal)

c4a("set2")

c4a("hcl.set2")

c4a("hcl.set2", n = 8)

# reversed palette
c4a("hcl.set2", reverse = TRUE, n = 8)

# handy shortcut
c4a("-hcl.set2", n = 8)

# the color for missing values is white:
c4a_na("hcl.set2")

---

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 utils::bibentry 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`.

Usage

```r
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 = TRUE,
  take.gray.for.NA = TRUE,
  remove.other.grays = FALSE,
  light.to.dark = TRUE,
  remove.names = TRUE,
  biv.method = "byrow",
  space = "rgb",
  range_matrix_args = list(NULL),
  bib = NA,
  description = NA
)
```

c4a_load(data)

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

x named list of color palettes. See details for indexing.

xNA colors for missing values. Vector of the same length as x (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).

types character vector of the same length as x (or length 1), which determines the type of palette: "cat", "seq", "div", "bivs", "bivc", "bivd", or "bivg". See details.

series a character vector of the same length as x (or length 1), which determines the series.

nmin, nmax, ndef minimum / maximum / default number of colors for the palette. By default: nmin = 1, for "cat" nmax and ndef the number of supplied colors. For the other types, nmax is Inf. ndef is 7 for "seq", 9. For diverging palettes, these numbers refer to the number of columns. (See mmin, mmax, mdef for the rows)

mmin, mmax, mdef minimum / maximum / default number of rows for bivariate palettes.

format.palette.name should palette names be formatted to lowercase/underscore format?

remove.blacks, take.gray.for.NA, remove.other.grays

These arguments determine the processing of grayscale colors for categorical "cat" palettes: if remove.blacks and there are (near) blacks, these are removed first. Next, if take.gray.for.NA, xNA 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.

remove.other.grays determines what happens with the other grays.

light.to.dark should sequential "seq" palettes be automatically ordered from light to dark?

remove.names should individual color names be removed?

biv.method 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 utils::bibentry 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 `c4a_data`... passed on to `c4a_data`

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,...,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 x 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", "#E366AA", "#715539"),
   atom = c("#C6E3FF", "#397139", "#557139", "#E3E3C6", "#1C1C1C", "#1C5555", "#AAAAAE", "#E3C6E3"),
   meddle = c("#715539", "#553939", "#8E7155", "#71AAA", "#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", "#555555", "#8E8E71", "#555555", "#8E7155"),
   wall = c("#FFFFFF", "#E3E3E3", "#C6C6C6", "#AAAC6", "#1C1C1C", "#000000", "#8E8E8E", "#E3C6E3"),
   cut = c("#000000", "#391C39", "#393955", "#E3C6E3", "#1C1C00", "#FFAA55", "#8E8E55"),
   lapse = c("#000000", "#E3E3C6", "#8E8E71", "#7171AA", "#39391C", "#717171", "#AAAAAA", "#E3E3E3"),
   division = c("#000000", "#FFFFFF", "#00398E", "#AA8E55", "#39558E", "#C6AA71", "#39391C", "#555571"),
   more = c("#0055AA", "#FFA1A1", "#1C71AA", "#003971", "#E38E55", "#E3AA8A", "#718EAA", "#71718E"),
   umma = c("#AA8E71", "#555539", "#39391C", "#1C1C1C", "#E3E3C6", "#715539", "#391C1C", "#8E7155"),
   relics = c("#3955AA", "#1C1C71", "#5571C6", "#715555", "#8E7155", "#E3AA71", "#8E8EAA", "#E3FFFF"),
   river = c("#393939", "#555555", "#39558E", "#C6C6C6", "#718EAA", "#1C1C1C", "#717171", "#E3C6E8"))

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") &&
interactive()) {
c4a_gui(series = "pinkfloyd", n = 8)
}
Graphical user interface to analyse palettes

Usage

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

c4a_table(
  type = c("cat", "seq", "div", "bivs", "bivc", "bivd", "bivg"),
  n = NULL,
  m = NULL,
  cvd.sim = c("none", "deutan", "protan", "tritan"),
  sort = "name",
  text.format = "hex",
  text.col = "same",
  series = "all",
  range = NA,
  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 c4a_gui 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" and "div", 9 colors, and for "bivs"/"bivc"/"bivd"/"bivg" 4 columns and rows. For c4a_gui 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 c4a_gui it only determines which series are shown initially.

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".
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.

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?

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 = "hueType")
}
**c4a_info**

*Get information from a cols4all palette*

**Description**

Get information from a cols4all palette

**Usage**

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

**Arguments**

- `palette` name of the palette
- `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.

---

**c4a_options**

*Set cols4all options*

**Description**

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

**Usage**

```r
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

<table>
<thead>
<tr>
<th>Option</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>defaults</td>
<td>Default palettes per type</td>
</tr>
<tr>
<td>CBF_th</td>
<td>Parameters that label a palette as color blind friendly</td>
</tr>
<tr>
<td>CBU_th</td>
<td>Parameters that label a palette as color blind unfriendly</td>
</tr>
<tr>
<td>CrangeFair</td>
<td>Maximum chroma range for which a palette is considered harmonic</td>
</tr>
<tr>
<td>CrangeUnfair</td>
<td>Minimum chroma range for which a palette is considered disharmonic</td>
</tr>
<tr>
<td>LrangeFair</td>
<td>Maximum luminance range for which a palette is considered harmonic</td>
</tr>
<tr>
<td>LrangeUnfair</td>
<td>Minimum luminance range for which a palette is considered disharmonic</td>
</tr>
<tr>
<td>Cintense</td>
<td>Chroma of colors that are considered intense</td>
</tr>
<tr>
<td>Cpastel</td>
<td>Chroma of colors that are considered 'pastel'</td>
</tr>
<tr>
<td>HwidthDivRainbow</td>
<td>A diverging palette is labeled as 'rainbow hue' if HwidthL or HwidthR are at least HwidthDivRainbow</td>
</tr>
<tr>
<td>HwidthDivSingle</td>
<td>A diverging palette is labeled as 'single hue' if HwidthL and HwidthR are at most HwidthDivSingle</td>
</tr>
<tr>
<td>HwidthSeqRainbow</td>
<td>A sequential palette is labeled as 'rainbow hue' if Hwidth is at least HwidthSeqRainbow</td>
</tr>
<tr>
<td>HwidthSeqSingle</td>
<td>A sequential palette is labeled as 'single hue' if Hwidth is at most HwidthSeqSingle</td>
</tr>
</tbody>
</table>

Value

A list of options

---

`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", and diverging "div" 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**

```r
# Get available palette names and series

c4a_palettes(
  type = c("all", "cat", "seq", "div"),
  series = NULL,
  full.names = TRUE
)

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

c4a_types(series = NULL, as.data.frame = TRUE)
```
c4a_overview()

Arguments

- **type**: type of color palette: one of "all" (all palettes), "cat" (categorical/qualitative palettes), "seq" (sequential palettes) and "div" (diverging palettes).
- **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?

Format

An object of class `environment` of length 16.

Value

names of the loaded color palettes

See Also

References of the palettes: `cols4all-package`.

Examples

```r
  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.

**Usage**
c4a_plot(palette, ..., include.na = FALSE)

**Arguments**
- **palette**  
  Palette name (see c4a) or a color vector
- **...**  
  arguments passed on to c4a
- **include.na**  
  should a color for missing values be included?

**Value**
Besides the plot, a gTree is returned silently

**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

```r
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 `color` 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

```r
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_color_discrete_c4a_cat

    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_discrete_c4a_cat

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(
scale_color_discrete_c4a_cat

`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

```r
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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