

# Package ‘scales’

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**Title** Scale Functions for Visualization

**Version** 1.1.1

**Description** Graphical scales map data to aesthetics, and provide methods for automatically determining breaks and labels for axes and legends.

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**URL** <https://scales.r-lib.org>, <https://github.com/r-lib/scales>

**BugReports** <https://github.com/r-lib/scales/issues>

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lifecycle,  
munsell (>= 0.5),  
R6,  
RColorBrewer,  
viridisLite

**Suggests** bit64,  
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*alpha* *Modify colour transparency*

**Description**

Vectorised in both colour and alpha.

**Usage**

```
alpha(colour, alpha = NA)
```

**Arguments**

colour	colour
alpha	new alpha level in [0,1]. If alpha is NA, existing alpha values are preserved.

**Examples**

```
alpha("red", 0.1)
alpha(colours(), 0.5)
alpha("red", seq(0, 1, length.out = 10))
alpha(c("first" = "gold", "second" = "lightgray", "third" = "#cd7f32"), .5)
```

*area\_pal* *Area palettes (continuous)*

**Description**

Area palettes (continuous)

**Usage**

```
area_pal(range = c(1, 6))
abs_area(max)
```

**Arguments**

range	Numeric vector of length two, giving range of possible sizes. Should be greater than 0.
max	A number representing the maximum size.

asn\_trans

*Arc-sin square root transformation*

---

**Description**

This is the variance stabilising transformation for the binomial distribution.

**Usage**

```
asn_trans()
```

**Examples**

```
plot(asn_trans(), xlim = c(0, 1))
```

---

atanh\_trans

*Arc-tangent transformation*

---

**Description**

Arc-tangent transformation

**Usage**

```
atanh_trans()
```

**Examples**

```
plot(atanh_trans(), xlim = c(-1, 1))
```

---

boxcox\_trans

*Box-Cox & modulus transformations*

---

**Description**

The Box-Cox transformation is a flexible transformation, often used to transform data towards normality. The modulus transformation generalises Box-Cox to also work with negative values.

**Usage**

```
boxcox_trans(p, offset = 0)
```

```
modulus_trans(p, offset = 1)
```

**Arguments**

**p** Transformation exponent,  $\lambda$ .

**offset** Constant offset. 0 for Box-Cox type 1, otherwise any non-negative constant (Box-Cox type 2). `modulus_trans()` sets the default to 1.

## Details

The Box-Cox power transformation (type 1) requires strictly positive values and takes the following form for  $y > 0$ :

$$y^{(\lambda)} = \frac{y^\lambda - 1}{\lambda}$$

When  $y = 0$ , the natural log transform is used.

The modulus transformation implements a generalisation of the Box-Cox transformation that works for data with both positive and negative values. The equation takes the following forms, when  $y \neq 0$ :

$$y^{(\lambda)} = \text{sign}(y) * \frac{(|y| + 1)^\lambda - 1}{\lambda}$$

and when  $y = 0$ :

$$y^{(\lambda)} = \text{sign}(y) * \ln(|y| + 1)$$

## References

Box, G. E., & Cox, D. R. (1964). An analysis of transformations. *Journal of the Royal Statistical Society. Series B (Methodological)*, 211-252. <https://www.jstor.org/stable/2984418>

John, J. A., & Draper, N. R. (1980). An alternative family of transformations. *Applied Statistics*, 190-197. <http://www.jstor.org/stable/2986305>

## See Also

[yj\\_trans\(\)](#)

## Examples

```
plot(boxcox_trans(-1), xlim = c(0, 10))
plot(boxcox_trans(0), xlim = c(0, 10))
plot(boxcox_trans(1), xlim = c(0, 10))
plot(boxcox_trans(2), xlim = c(0, 10))

plot(modulus_trans(-1), xlim = c(-10, 10))
plot(modulus_trans(0), xlim = c(-10, 10))
plot(modulus_trans(1), xlim = c(-10, 10))
plot(modulus_trans(2), xlim = c(-10, 10))
```

---

breaks\_extended

*Automatic breaks for numeric axes*

---

## Description

Uses Wilkinson's extended breaks algorithm as implemented in the **labeling** package.

## Usage

```
breaks_extended(n = 5, ...)
```

**Arguments**

n	Desired number of breaks. You may get slightly more or fewer breaks than requested.
...	other arguments passed on to <code>labeling::extended()</code>

**References**

Talbot, J., Lin, S., Hanrahan, P. (2010) An Extension of Wilkinson's Algorithm for Positioning Tick Labels on Axes, InfoVis 2010 <http://vis.stanford.edu/files/2010-TickLabels-InfoVis.pdf>.

**Examples**

```
demo_continuous(c(0, 10))
demo_continuous(c(0, 10), breaks = breaks_extended(3))
demo_continuous(c(0, 10), breaks = breaks_extended(10))
```

---

breaks\_log

*Breaks for log axes*


---

**Description**

This algorithm starts by looking for integer powers of base. If that doesn't provide enough breaks, it then looks for additional intermediate breaks which are integer multiples of integer powers of base. If that fails (which it can for very small ranges), we fall back to `extended_breaks()`

**Usage**

```
breaks_log(n = 5, base = 10)
```

**Arguments**

n	desired number of breaks
base	base of logarithm to use

**Details**

The algorithm starts by looking for a set of integer powers of base that cover the range of the data. If that does not generate at least  $n - 2$  breaks, we look for an integer between 1 and base that splits the interval approximately in half. For example, in the case of base = 10, this integer is 3 because  $\log_{10}(3) = 0.477$ . This leaves 2 intervals:  $c(1, 3)$  and  $c(3, 10)$ . If we still need more breaks, we look for another integer that splits the largest remaining interval (on the log-scale) approximately in half. For base = 10, this is 5 because  $\log_{10}(5) = 0.699$ .

The generic algorithm starts with a set of integers steps containing only 1 and a set of candidate integers containing all integers larger than 1 and smaller than base. Then for each remaining candidate integer  $x$ , the smallest interval (on the log-scale) in the vector `sort(c(x, steps, base))` is calculated. The candidate  $x$  which yields the largest minimal interval is added to steps and removed from the candidate set. This is repeated until either a sufficient number of breaks,  $\geq n - 2$ , are returned or all candidates have been used.

## Examples

```
demo_log10(c(1, 1e5))
demo_log10(c(1, 1e6))

# Request more breaks by setting n
demo_log10(c(1, 1e6), breaks = breaks_log(6))

# Some tricky ranges
demo_log10(c(2000, 9000))
demo_log10(c(2000, 14000))
demo_log10(c(2000, 85000), expand = c(0, 0))

# An even smaller range that requires falling back to linear breaks
demo_log10(c(1800, 2000))
```

---

breaks_pretty	<i>Pretty breaks for date/times</i>
---------------	-------------------------------------

---

## Description

Uses default R break algorithm as implemented in `pretty()`. This is primarily useful for date/times, as `extended_breaks()` should do a slightly better job for numeric scales.

## Usage

```
breaks_pretty(n = 5, ...)
```

## Arguments

<code>n</code>	Desired number of breaks. You may get slightly more or fewer breaks than requested.
<code>...</code>	other arguments passed on to <code>pretty()</code>

## Details

`pretty_breaks()` is retired; use `breaks_pretty()` instead.

## Examples

```
one_month <- as.POSIXct(c("2020-05-01", "2020-06-01"))
demo_datetime(one_month)
demo_datetime(one_month, breaks = breaks_pretty(2))
demo_datetime(one_month, breaks = breaks_pretty(4))

# Tightly spaced date breaks often need custom labels too
demo_datetime(one_month, breaks = breaks_pretty(12))
demo_datetime(one_month,
  breaks = breaks_pretty(12),
  labels = label_date_short()
)
```

---

breaks_width	<i>Equally spaced breaks</i>
--------------	------------------------------

---

### Description

Useful for numeric, date, and date-time scales.

### Usage

```
breaks_width(width, offset = 0)
```

### Arguments

width	Distance between each break. Either a number, or for date/times, a single string of the form "n unit", e.g. "1 month", "5 days". Unit can be of one "sec", "min", "hour", "day", "week", "month", "year".
offset	Use if you don't want breaks to start at zero

### Examples

```
demo_continuous(c(0, 100))
demo_continuous(c(0, 100), breaks = breaks_width(10))
demo_continuous(c(0, 100), breaks = breaks_width(20, -4))
demo_continuous(c(0, 100), breaks = breaks_width(20, 4))

# This is also useful for dates
one_month <- as.POSIXct(c("2020-05-01", "2020-06-01"))
demo_datetime(one_month)
demo_datetime(one_month, breaks = breaks_width("1 week"))
demo_datetime(one_month, breaks = breaks_width("5 days"))
# This is so useful that scale_x_datetime() has a shorthand:
demo_datetime(one_month, date_breaks = "5 days")

# hms times also work
one_hour <- hms::hms(hours = 0:1)
demo_time(one_hour)
demo_time(one_hour, breaks = breaks_width("15 min"))
demo_time(one_hour, breaks = breaks_width("600 sec"))
```

---

brewer_pal	<i>Colour Brewer palette (discrete)</i>
------------	---

---

### Description

Colour Brewer palette (discrete)

### Usage

```
brewer_pal(type = "seq", palette = 1, direction = 1)
```

**Arguments**

type	One of seq (sequential), div (diverging) or qual (qualitative)
palette	If a string, will use that named palette. If a number, will index into the list of palettes of appropriate type
direction	Sets the order of colours in the scale. If 1, the default, colours are as output by <code>RColorBrewer::brewer.pal()</code> . If -1, the order of colours is reversed.

**References**

<http://colorbrewer2.org>

**Examples**

```
show_col(brewer_pal()(10))
show_col(brewer_pal("div")(5))
show_col(brewer_pal(palette = "Greens")(5))

# Can use with gradient_n to create a continuous gradient
cols <- brewer_pal("div")(5)
show_col(gradient_n_pal(cols)(seq(0, 1, length.out = 30)))
```

---

col2hcl

---

*Modify standard R colour in hcl colour space.*


---

**Description**

Transforms rgb to hcl, sets non-missing arguments and then backtransforms to rgb.

**Usage**

```
col2hcl(colour, h = NULL, c = NULL, l = NULL, alpha = NULL)
```

**Arguments**

colour	character vector of colours to be modified
h	Hue, [0, 360]
c	Chroma, [0, 100]
l	Luminance, [0, 100]
alpha	Alpha, [0, 1].

**Examples**

```
reds <- rep("red", 6)
show_col(col2hcl(reds, h = seq(0, 180, length = 6)))
show_col(col2hcl(reds, c = seq(0, 80, length = 6)))
show_col(col2hcl(reds, l = seq(0, 100, length = 6)))
show_col(col2hcl(reds, alpha = seq(0, 1, length = 6)))
```

---

`colour_ramp`*Fast colour interpolation*

---

## Description

Returns a function that maps the interval [0,1] to a set of colours. Interpolation is performed in the CIELAB colour space. Similar to `colorRamp(space = 'Lab')`, but hundreds of times faster, and provides results in `"#RRGGBB"` (or `"#RRGGBBAA"`) character form instead of RGB colour matrices.

## Usage

```
colour_ramp(colors, na.color = NA, alpha = TRUE)
```

## Arguments

- |                       |   |
|-----------------------|---|
| <code>colors</code>   | Colours to interpolate; must be a valid argument to <code>grDevices::col2rgb()</code> . This can be a character vector of <code>"#RRGGBB"</code> or <code>"#RRGGBBAA"</code> , colour names from <code>grDevices::colors()</code> , or a positive integer that indexes into <code>grDevices::palette()</code> .   |
| <code>na.color</code> | The colour to map to NA values (for example, <code>"#606060"</code> for dark grey, or <code>"#00000000"</code> for transparent) and values outside of [0,1]. Can itself be NA, which will simply cause an NA to be inserted into the output.  |
| <code>alpha</code>    | Whether to include alpha transparency channels in interpolation. If TRUE then the alpha information is included in the interpolation. The returned colours will be provided in <code>"#RRGGBBAA"</code> format when needed, i.e., in cases where the colour is not fully opaque, so that the <code>"AA"</code> part is not equal to <code>"FF"</code> . Fully opaque colours will be returned in <code>"#RRGGBB"</code> format. If FALSE, the alpha information is discarded before interpolation and colours are always returned as <code>"#RRGGBB"</code> . |

## Value

A function that takes a numeric vector and returns a character vector of the same length with RGB or RGBA hex colours.

## See Also

[colorRamp](#)

## Examples

```
ramp <- colour_ramp(c("red", "green", "blue"))
show_col(ramp(seq(0, 1, length = 12)))
```

---

col_numeric	<i>Colour mapping</i>
-------------	-----------------------

---

## Description

Conveniently maps data values (numeric or factor/character) to colours according to a given palette, which can be provided in a variety of formats.

## Usage

```
col_numeric(  
  palette,  
  domain,  
  na.color = "#808080",  
  alpha = FALSE,  
  reverse = FALSE  
)  
  
col_bin(  
  palette,  
  domain,  
  bins = 7,  
  pretty = TRUE,  
  na.color = "#808080",  
  alpha = FALSE,  
  reverse = FALSE,  
  right = FALSE  
)  
  
col_quantile(  
  palette,  
  domain,  
  n = 4,  
  probs = seq(0, 1, length.out = n + 1),  
  na.color = "#808080",  
  alpha = FALSE,  
  reverse = FALSE,  
  right = FALSE  
)  
  
col_factor(  
  palette,  
  domain,  
  levels = NULL,  
  ordered = FALSE,  
  na.color = "#808080",  
  alpha = FALSE,  
  reverse = FALSE  
)
```

**Arguments**

palette	The colours or colour function that values will be mapped to
domain	The possible values that can be mapped. For <code>col_numeric</code> and <code>col_bin</code> , this can be a simple numeric range (e.g. <code>c(0, 100)</code> ); <code>col_quantile</code> needs representative numeric data; and <code>col_factor</code> needs categorical data. If <code>NULL</code> , then whenever the resulting colour function is called, the <code>x</code> value will represent the domain. This implies that if the function is invoked multiple times, the encoding between values and colours may not be consistent; if consistency is needed, you must provide a non- <code>NULL</code> domain.
na.color	The colour to return for NA values. Note that <code>na.color = NA</code> is valid.
alpha	Whether alpha channels should be respected or ignored. If <code>TRUE</code> then colors without explicit alpha information will be treated as fully opaque.
reverse	Whether the colors (or color function) in <code>palette</code> should be used in reverse order. For example, if the default order of a palette goes from blue to green, then <code>reverse = TRUE</code> will result in the colors going from green to blue.
bins	Either a numeric vector of two or more unique cut points or a single number (greater than or equal to 2) giving the number of intervals into which the domain values are to be cut.
pretty	Whether to use the function <code>pretty()</code> to generate the bins when the argument <code>bins</code> is a single number. When <code>pretty = TRUE</code> , the actual number of bins may not be the number of bins you specified. When <code>pretty = FALSE</code> , <code>seq()</code> is used to generate the bins and the breaks may not be "pretty".
right	parameter supplied to <code>base::cut()</code> . See Details
n	Number of equal-size quantiles desired. For more precise control, use the <code>probs</code> argument instead.
probs	See <code>stats::quantile()</code> . If provided, the <code>n</code> argument is ignored.
levels	An alternate way of specifying levels; if specified, <code>domain</code> is ignored
ordered	If <code>TRUE</code> and <code>domain</code> needs to be coerced to a factor, treat it as already in the correct order

**Details**

`col_numeric` is a simple linear mapping from continuous numeric data to an interpolated palette.

`col_bin` also maps continuous numeric data, but performs binning based on value (see the `base::cut()` function). `col_bin` defaults for the cut function are `include.lowest = TRUE` and `right = FALSE`.

`col_quantile` similarly bins numeric data, but via the `stats::quantile()` function.

`col_factor` maps factors to colours. If the palette is discrete and has a different number of colours than the number of factors, interpolation is used.

The `palette` argument can be any of the following:

1. A character vector of RGB or named colours. Examples: `palette()`, `c("#000000", "#0000FF", "#FFFFFF")`, `topo.colors(10)`
2. The name of an RColorBrewer palette, e.g. "BuPu" or "Greens".
3. The full name of a viridis palette: "viridis", "magma", "inferno", or "plasma".
4. A function that receives a single value between 0 and 1 and returns a colour. Examples: `colorRamp(c("#000000", "#FFFFFF"), interpolate="spline")`.

**Value**

A function that takes a single parameter `x`; when called with a vector of numbers (except for `col_factor`, which expects factors/characters), #RRGGBB colour strings are returned (unless `alpha = TRUE` in which case #RRGGBBAA may also be possible).

**Examples**

```
pal <- col_bin("Greens", domain = 0:100)
show_col(pal(sort(runif(10, 60, 100))))

# Exponential distribution, mapped continuously
show_col(col_numeric("Blues", domain = NULL)(sort(rexp(16))))
# Exponential distribution, mapped by interval
show_col(col_bin("Blues", domain = NULL, bins = 4)(sort(rexp(16))))
# Exponential distribution, mapped by quantile
show_col(col_quantile("Blues", domain = NULL)(sort(rexp(16))))

# Categorical data; by default, the values being coloured span the gamut...
show_col(col_factor("RdYlBu", domain = NULL)(LETTERS[1:5]))
# ...unless the data is a factor, without droplevels...
show_col(col_factor("RdYlBu", domain = NULL)(factor(LETTERS[1:5], levels=LETTERS)))
# ...or the domain is stated explicitly.
show_col(col_factor("RdYlBu", levels = LETTERS)(LETTERS[1:5]))
```

---

cscale

*Continuous scale*


---

**Description**

Continuous scale

**Usage**

```
cscale(x, palette, na.value = NA_real_, trans = identity_trans())
```

**Arguments**

<code>x</code>	vector of continuous values to scale
<code>palette</code>	palette to use. Built in palettes: <a href="#">area_pal</a> , <a href="#">brewer_pal</a> , <a href="#">dichromat_pal</a> , <a href="#">div_gradient_pal</a> , <a href="#">gradient_n_pal</a> , <a href="#">grey_pal</a> , <a href="#">hue_pal</a> , <a href="#">identity_pal</a> , <a href="#">linetype_pal</a> , <a href="#">manual_pal</a> , <a href="#">rescale_pal</a> , <a href="#">seq_gradient_pal</a> , <a href="#">shape_pal</a> , <a href="#">viridis_pal</a>
<code>na.value</code>	value to use for missing values
<code>trans</code>	transformation object describing the how to transform the raw data prior to scaling. Defaults to the identity transformation which leaves the data unchanged. Built in transformations: <a href="#">asn_trans</a> , <a href="#">atanh_trans</a> , <a href="#">boxcox_trans</a> , <a href="#">date_trans</a> , <a href="#">exp_trans</a> , <a href="#">hms_trans</a> , <a href="#">identity_trans</a> , <a href="#">log10_trans</a> , <a href="#">log1p_trans</a> , <a href="#">log2_trans</a> , <a href="#">log_trans</a> , <a href="#">logit_trans</a> , <a href="#">modulus_trans</a> , <a href="#">probability_trans</a> , <a href="#">probit_trans</a> , <a href="#">pseudo_log_trans</a> , <a href="#">reciprocal_trans</a> , <a href="#">reverse_trans</a> , <a href="#">sqrt_trans</a> , <a href="#">time_trans</a> , <a href="#">yj_trans</a> .

**Examples**

```
with(mtcars, plot(displ, mpg, cex = cscale(hp, rescale_pal()))
with(mtcars, plot(displ, mpg, cex = cscale(hp, rescale_pal(),
      trans = sqrt_trans()))
with(mtcars, plot(displ, mpg, cex = cscale(hp, area_pal()))
with(mtcars, plot(displ, mpg, pch = 20, cex = 5,
      col = cscale(hp, seq_gradient_pal("grey80", "black"))))
```

---

date\_trans

*Transformation for dates (class Date)*


---

**Description**

Transformation for dates (class Date)

**Usage**

```
date_trans()
```

**Examples**

```
years <- seq(as.Date("1910/1/1"), as.Date("1999/1/1"), "years")
t <- date_trans()
t$transform(years)
t$inverse(t$transform(years))
t$format(t$breaks(range(years)))
```

---

dichromat\_pal

*Dichromat (colour-blind) palette (discrete)*


---

**Description**

Dichromat (colour-blind) palette (discrete)

**Usage**

```
dichromat_pal(name)
```

**Arguments**

name Name of colour palette. One of: BrownToBlue.10, BrownToBlue.12, BlueToDarkOrange.12, BlueToDarkOrange.18, DarkRedToBlue.12, DarkRedToBlue.18, BlueToGreen.14, BlueToGray.8, BlueToOrangeRed.14, BlueToOrange.10, BlueToOrange.12, BlueToOrange.8, LightBlueToDarkBlue.10, LightBlueToDarkBlue.7, Categorical.12, GreenToMagenta.16, SteppedSequential.5

**Examples**

```

if (requireNamespace("dichromat", quietly = TRUE)) {
  show_col(dichromat_pal("BluetoOrange.10")(10))
  show_col(dichromat_pal("BluetoOrange.10")(5))

  # Can use with gradient_n to create a continuous gradient
  cols <- dichromat_pal("DarkRedtoBlue.12")(12)
  show_col(gradient_n_pal(cols)(seq(0, 1, length.out = 30)))
}

```

---

div\_gradient\_pal      *Diverging colour gradient (continuous).*

---

**Description**

Diverging colour gradient (continuous).

**Usage**

```

div_gradient_pal(
  low = mns1("10B 4/6"),
  mid = mns1("N 8/0"),
  high = mns1("10R 4/6"),
  space = "Lab"
)

```

**Arguments**

low	colour for low end of gradient.
mid	colour for mid point
high	colour for high end of gradient.
space	colour space in which to calculate gradient. Must be "Lab" - other values are deprecated.

**Examples**

```

x <- seq(-1, 1, length.out = 100)
r <- sqrt(outer(x^2, x^2, "+"))
image(r, col = div_gradient_pal()(seq(0, 1, length.out = 12)))
image(r, col = div_gradient_pal()(seq(0, 1, length.out = 30)))
image(r, col = div_gradient_pal()(seq(0, 1, length.out = 100)))

library(munsell)
image(r, col = div_gradient_pal(low =
  mns1(complement("10R 4/6"), fix = TRUE))(seq(0, 1, length = 100)))

```

---

dscale	<i>Discrete scale</i>
--------	-----------------------

---

**Description**

Discrete scale

**Usage**

```
dscale(x, palette, na.value = NA)
```

**Arguments**

x	vector of discrete values to scale
palette	aesthetic palette to use
na.value	aesthetic to use for missing values

**Examples**

```
with(mtcars, plot(displ, mpg, pch = 20, cex = 3,  
  col = dscale(factor(cyl), brewer_pal())))
```

---

expand_range	<i>Expand a range with a multiplicative or additive constant</i>
--------------	--

---

**Description**

Expand a range with a multiplicative or additive constant

**Usage**

```
expand_range(range, mul = 0, add = 0, zero_width = 1)
```

**Arguments**

range	range of data, numeric vector of length 2
mul	multiplicative constant
add	additive constant
zero_width	distance to use if range has zero width

---

exp_trans	<i>Exponential transformation (inverse of log transformation)</i>
-----------	---

---

**Description**

Exponential transformation (inverse of log transformation)

**Usage**

```
exp_trans(base = exp(1))
```

**Arguments**

base	Base of logarithm
------	-------------------

**Examples**

```
plot(exp_trans(0.5), xlim = c(-2, 2))
plot(exp_trans(1), xlim = c(-2, 2))
plot(exp_trans(2), xlim = c(-2, 2))
plot(exp_trans(), xlim = c(-2, 2))
```

---

gradient_n_pal	<i>Arbitrary colour gradient palette (continuous)</i>
----------------	---

---

**Description**

Arbitrary colour gradient palette (continuous)

**Usage**

```
gradient_n_pal(colours, values = NULL, space = "Lab")
```

**Arguments**

colours	vector of colours
values	if colours should not be evenly positioned along the gradient this vector gives the position (between 0 and 1) for each colour in the colours vector. See <a href="#">rescale()</a> for a convenience function to map an arbitrary range to between 0 and 1.
space	colour space in which to calculate gradient. Must be "Lab" - other values are deprecated.

---

grey_pal	<i>Grey scale palette (discrete)</i>
----------	--------------------------------------

---

**Description**

Grey scale palette (discrete)

**Usage**

```
grey_pal(start = 0.2, end = 0.8)
```

**Arguments**

start	grey value at low end of palette
end	grey value at high end of palette

**See Also**

[seq\\_gradient\\_pal\(\)](#) for continuous version

**Examples**

```
show_col(grey_pal()(25))  
show_col(grey_pal(0, 1)(25))
```

---

hms_trans	<i>Transformation for times (class hms)</i>
-----------	---

---

**Description**

Transformation for times (class hms)

**Usage**

```
hms_trans()
```

**Examples**

```
if (require("hms")) {  
  hms <- round(runif(10) * 86400)  
  t <- hms_trans()  
  t$transform(hms)  
  t$inverse(t$transform(hms))  
  t$breaks(hms)  
}
```

---

hue_pal	<i>Hue palette (discrete)</i>
---------	-------------------------------

---

**Description**

Hue palette (discrete)

**Usage**

```
hue_pal(h = c(0, 360) + 15, c = 100, l = 65, h.start = 0, direction = 1)
```

**Arguments**

h	range of hues to use, in [0, 360]
c	chroma (intensity of colour), maximum value varies depending on combination of hue and luminance.
l	luminance (lightness), in [0, 100]
h.start	hue to start at
direction	direction to travel around the colour wheel, 1 = clockwise, -1 = counter-clockwise

**Examples**

```
show_col(hue_pal()(4))
show_col(hue_pal()(9))
show_col(hue_pal(l = 90)(9))
show_col(hue_pal(l = 30)(9))

show_col(hue_pal()(9))
show_col(hue_pal(direction = -1)(9))

show_col(hue_pal()(9))
show_col(hue_pal(h = c(0, 90))(9))
show_col(hue_pal(h = c(90, 180))(9))
show_col(hue_pal(h = c(180, 270))(9))
show_col(hue_pal(h = c(270, 360))(9))
```

---

identity_pal	<i>Identity palette</i>
--------------	-------------------------

---

**Description**

Leaves values unchanged - useful when the data is already scaled.

**Usage**

```
identity_pal()
```

---

identity_trans	<i>Identity transformation (do nothing)</i>
----------------	---

---

**Description**

Identity transformation (do nothing)

**Usage**

```
identity_trans()
```

**Examples**

```
plot(identity_trans(), xlim = c(-1, 1))
```

---

label_bytes	<i>Label bytes (1 kb, 2 MB, etc)</i>
-------------	--------------------------------------

---

**Description**

Scale bytes into human friendly units. Can use either SI units (e.g. kB = 1000 bytes) or binary units (e.g. kiB = 1024 bytes). See [Units of Information](#) on Wikipedia for more details.

**Usage**

```
label_bytes(units = "auto_si", accuracy = 1, ...)
```

**Arguments**

units	Unit to use. Should either one of: <ul style="list-style-type: none"> <li>• "kB", "MB", "GB", "TB", "PB", "EB", "ZB", and "YB" for SI units (base 1000).</li> <li>• "kiB", "MiB", "GiB", "TiB", "PiB", "EiB", "ZiB", and "YiB" for binary units (base 1024).</li> <li>• auto_si or auto_binary to automatically pick the most appropriate unit for each value.</li> </ul>
accuracy	A number to round to. Use (e.g.) 0.01 to show 2 decimal places of precision. If NULL, the default, uses a heuristic that should ensure breaks have the minimum number of digits needed to show the difference between adjacent values. Applied to rescaled data.
...	Other arguments passed on to <a href="#">number()</a>

**Value**

A labeller function that takes a numeric vector of breaks and returns a character vector of labels.

**See Also**

Other labels for continuous scales: [label\\_dollar\(\)](#), [label\\_number\\_auto\(\)](#), [label\\_number\\_si\(\)](#), [label\\_ordinal\(\)](#), [label\\_parse\(\)](#), [label\\_percent\(\)](#), [label\\_pvalue\(\)](#), [label\\_scientific\(\)](#)

Other labels for log scales: [label\\_number\\_si\(\)](#), [label\\_scientific\(\)](#)

**Examples**

```
demo_continuous(c(1, 1e6))
demo_continuous(c(1, 1e6), label = label_bytes())
# Force all to use same units
demo_continuous(c(1, 1e6), label = label_bytes("kB"))

# Auto units are particularly nice on log scales
demo_log10(c(1, 1e6))
demo_log10(c(1, 1e7), label = label_bytes())

# You can also use binary units where a megabyte is defined as
# (1024) ^ 2 bytes rather than (1000) ^ 2. You'll need to override
# the default breaks to make this more informative.
demo_continuous(c(1, 1024^2), label = label_bytes("auto_binary"))
demo_continuous(c(1, 1024^2),
  breaks = breaks_width(250 * 1024),
  label = label_bytes("auto_binary")
)
```

---

label\_date

*Label date/times*


---

**Description**

`label_date()` and `label_time()` label date/times using date/time format strings. `label_date_short()` automatically constructs a short format string sufficient to uniquely identify labels. It's inspired by matplotlib's [ConciseDateFormatter](#), but uses a slightly different approach: `ConciseDateFormatter` formats "firsts" (e.g. first day of month, first day of day) specially; `date_short()` formats changes (e.g. new month, new year) specially.

**Usage**

```
label_date(format = "%Y-%m-%d", tz = "UTC")
```

```
label_date_short(format = c("%Y", "%b", "%d", "%H:%M"), sep = "\n")
```

```
label_time(format = "%H:%M:%S", tz = "UTC")
```

```
date_format(format = "%Y-%m-%d", tz = "UTC")
```

```
time_format(format = "%H:%M:%S", tz = "UTC")
```

**Arguments**

format	For <code>date_format()</code> and <code>time_format()</code> a date/time format string using standard POSIX specification. See <code>strptime()</code> for details. For <code>date_short()</code> a character vector of length 4 giving the format components to use for year, month, day, and hour respectively.
tz	a time zone name, see <code>timezones()</code> . Defaults to UTC
sep	Separator to use when combining date formats into a single string.

**Value**

All `label_()` functions return a "labelling" function, i.e. a function that takes a vector `x` and returns a character vector of `length(x)` giving a label for each input value.

Labelling functions are designed to be used with the `labels` argument of `ggplot2` scales. The examples demonstrate their use with `x` scales, but they work similarly for all scales, including those that generate legends rather than axes.

**Old interface**

`date_format()` and `time_format()` are retired; please use `label_date()` and `label_time()` instead.

**Examples**

```
date_range <- function(start, days) {
  start <- as.POSIXct(start)
  c(start, start + days * 24 * 60 * 60)
}

two_months <- date_range("2020-05-01", 60)
demo_datetime(two_months)
demo_datetime(two_months, labels = date_format("%m/%d"))
# ggplot2 provides a short-hand:
demo_datetime(two_months, date_labels = "%m/%d")

# An alternative labelling system is label_date_short()
demo_datetime(two_months, date_breaks = "7 days", labels = label_date_short())
# This is particularly effective for dense labels
one_year <- date_range("2020-05-01", 365)
demo_datetime(one_year, date_breaks = "month")
demo_datetime(one_year, date_breaks = "month", labels = label_date_short())
```

---

label\_dollar

*Label currencies (\$100, \$2.50, etc)*


---

**Description**

Format numbers as currency, rounding values to dollars or cents using a convenient heuristic.

**Usage**

```
label_dollar(
  accuracy = NULL,
  scale = 1,
  prefix = "$",
  suffix = "",
  big.mark = ",",
  decimal.mark = ".",
  trim = TRUE,
  largest_with_cents = 1e+05,
  negative_parens = FALSE,
  ...
)
```

```
dollar_format(
  accuracy = NULL,
  scale = 1,
  prefix = "$",
  suffix = "",
  big.mark = ",",
  decimal.mark = ".",
  trim = TRUE,
  largest_with_cents = 1e+05,
  negative_parens = FALSE,
  ...
)
```

```
dollar(
  x,
  accuracy = NULL,
  scale = 1,
  prefix = "$",
  suffix = "",
  big.mark = ",",
  decimal.mark = ".",
  trim = TRUE,
  largest_with_cents = 1e+05,
  negative_parens = FALSE,
  ...
)
```

**Arguments**

accuracy, largest_with_cents	Number to round to. If NULL, the default, values will be rounded to the nearest integer, unless any of the values has non-zero fractional component (e.g. cents) and the largest value is less than largest_with_cents which by default is 100,000.
scale	A scaling factor: x will be multiplied by scale before formatting. This is useful if the underlying data is very small or very large.
prefix, suffix	Symbols to display before and after value.
big.mark	Character used between every 3 digits to separate thousands.

decimal.mark	The character to be used to indicate the numeric decimal point.
trim	Logical, if FALSE, values are right-justified to a common width (see <code>base::format()</code> ).
negative_parens	Display negative using parentheses?
...	Other arguments passed on to <code>base::format()</code> .
x	A numeric vector

### Value

All `label_()` functions return a "labelling" function, i.e. a function that takes a vector `x` and returns a character vector of `length(x)` giving a label for each input value.

Labelling functions are designed to be used with the `labels` argument of `ggplot2` scales. The examples demonstrate their use with `x` scales, but they work similarly for all scales, including those that generate legends rather than axes.

### Old interface

`dollar()` and `format_dollar()` are retired; please use `label_dollar()` instead.

### See Also

Other labels for continuous scales: `label_bytes()`, `label_number_auto()`, `label_number_si()`, `label_ordinal()`, `label_parse()`, `label_percent()`, `label_pvalue()`, `label_scientific()`

### Examples

```
demo_continuous(c(0, 1), labels = label_dollar())
demo_continuous(c(1, 100), labels = label_dollar())

# Customise currency display with prefix and suffix
demo_continuous(c(1, 100), labels = label_dollar(prefix = "USD "))
euro <- dollar_format(
  prefix = "",
  suffix = "\u20ac",
  big.mark = ".",
  decimal.mark = ",",
)
demo_continuous(c(1000, 1100), labels = euro)

# Use negative_parens = TRUE for finance style display
demo_continuous(c(-100, 100), labels = label_dollar(negative_parens = TRUE))
```

---

label\_number

*Label numbers in decimal format (e.g. 0.12, 1,234)*

---

### Description

Use `label_number()` force decimal display of numbers (i.e. don't use `scientific` notation). `label_comma()` is a special case that inserts a comma every three digits.

**Usage**

```
label_number(  
  accuracy = NULL,  
  scale = 1,  
  prefix = "",  
  suffix = "",  
  big.mark = " ",  
  decimal.mark = ".",  
  trim = TRUE,  
  ...  
)
```

```
label_comma(  
  accuracy = NULL,  
  scale = 1,  
  prefix = "",  
  suffix = "",  
  big.mark = ",",  
  decimal.mark = ".",  
  trim = TRUE,  
  digits,  
  ...  
)
```

```
comma(  
  x,  
  accuracy = NULL,  
  scale = 1,  
  prefix = "",  
  suffix = "",  
  big.mark = ",",  
  decimal.mark = ".",  
  trim = TRUE,  
  digits,  
  ...  
)
```

```
number_format(  
  accuracy = NULL,  
  scale = 1,  
  prefix = "",  
  suffix = "",  
  big.mark = " ",  
  decimal.mark = ".",  
  trim = TRUE,  
  ...  
)
```

```
comma_format(  
  accuracy = NULL,  
  scale = 1,  
  prefix = "",
```

```

    suffix = "",
    big.mark = ",",
    decimal.mark = ".",
    trim = TRUE,
    digits,
    ...
  )

```

### Arguments

accuracy	A number to round to. Use (e.g.) 0.01 to show 2 decimal places of precision. If NULL, the default, uses a heuristic that should ensure breaks have the minimum number of digits needed to show the difference between adjacent values. Applied to rescaled data.
scale	A scaling factor: x will be multiplied by scale before formatting. This is useful if the underlying data is very small or very large.
prefix, suffix	Symbols to display before and after value.
big.mark	Character used between every 3 digits to separate thousands.
decimal.mark	The character to be used to indicate the numeric decimal point.
trim	Logical, if FALSE, values are right-justified to a common width (see <code>base::format()</code> ).
...	Other arguments passed on to <code>base::format()</code> .
digits	Deprecated, use accuracy instead.
x	A numeric vector to format.

### Value

All `label_()` functions return a "labelling" function, i.e. a function that takes a vector x and returns a character vector of `length(x)` giving a label for each input value.

Labelling functions are designed to be used with the `labels` argument of `ggplot2` scales. The examples demonstrate their use with x scales, but they work similarly for all scales, including those that generate legends rather than axes.

### Old interface

`number_format()`, `comma_format()`, and `comma()` are retired; please use `label_number()` and `label_comma()` instead.

### Examples

```

demo_continuous(c(-1e6, 1e6))
demo_continuous(c(-1e6, 1e6), labels = label_number())
demo_continuous(c(-1e6, 1e6), labels = label_comma())

# Use scale to rescale very small or large numbers to generate
# more readable labels
demo_continuous(c(0, 1e6), labels = label_number())
demo_continuous(c(0, 1e6), labels = label_number(scale = 1 / 1e3))
demo_continuous(c(0, 1e-6), labels = label_number())
demo_continuous(c(0, 1e-6), labels = label_number(scale = 1e6))

# You can use prefix and suffix for other types of display
demo_continuous(c(32, 212), label = label_number(suffix = "\u00b0F"))
demo_continuous(c(0, 100), label = label_number(suffix = "\u00b0C"))

```

---

label_number_auto	<i>Label numbers, avoiding scientific notation where possible</i>
-------------------	---

---

### Description

Switches between `number_format()` and `scientific_format()` based on a set of heuristics designed to automatically generate useful labels across a wide range of inputs

### Usage

```
label_number_auto()
```

### See Also

Other labels for continuous scales: `label_bytes()`, `label_dollar()`, `label_number_si()`, `label_ordinal()`, `label_parse()`, `label_percent()`, `label_pvalue()`, `label_scientific()`

### Examples

```
# Very small and very large numbers get scientific notation
demo_continuous(c(0, 1e-6), labels = label_number_auto())
demo_continuous(c(0, 1e9), labels = label_number_auto())

# Other ranges get the numbers printed in full
demo_continuous(c(0, 1e-3), labels = label_number_auto())
demo_continuous(c(0, 1), labels = label_number_auto())
demo_continuous(c(0, 1e3), labels = label_number_auto())
demo_continuous(c(0, 1e6), labels = label_number_auto())

# Transformation is applied individually so you get as little
# scientific notation as possible
demo_log10(c(1, 1e7), labels = label_number_auto())
```

---

label_number_si	<i>Label numbers with SI prefixes (2k, 1M, 5T etc)</i>
-----------------	--

---

### Description

`number_si()` automatically scales and labels with the best SI prefix, "K" for values  $\geq 10e3$ , "M" for  $\geq 10e6$ , "B" for  $\geq 10e9$ , and "T" for  $\geq 10e12$ .

### Usage

```
label_number_si(accuracy = 1, unit = NULL, sep = NULL, ...)
```

**Arguments**

accuracy	A number to round to. Use (e.g.) 0.01 to show 2 decimal places of precision. If NULL, the default, uses a heuristic that should ensure breaks have the minimum number of digits needed to show the difference between adjacent values. Applied to rescaled data.
unit	Optional units specifier.
sep	Separator between number and SI unit. Defaults to " " if units is supplied, and "" if not.
...	Other arguments passed on to <code>base::format()</code> .

**Value**

All `label_()` functions return a "labelling" function, i.e. a function that takes a vector `x` and returns a character vector of `length(x)` giving a label for each input value.

Labelling functions are designed to be used with the `labels` argument of `ggplot2` scales. The examples demonstrate their use with `x` scales, but they work similarly for all scales, including those that generate legends rather than axes.

**See Also**

Other labels for continuous scales: `label_bytes()`, `label_dollar()`, `label_number_auto()`, `label_ordinal()`, `label_parse()`, `label_percent()`, `label_pvalue()`, `label_scientific()`

Other labels for log scales: `label_bytes()`, `label_scientific()`

**Examples**

```
demo_continuous(c(1, 1e9), label = label_number_si())
demo_continuous(c(1, 5000), label = label_number_si(unit = "g"))
demo_continuous(c(1, 1000), label = label_number_si(unit = "m"))

demo_log10(c(1, 1e9), breaks = log_breaks(10), labels = label_number_si())
```

---

label\_ordinal

*Label ordinal numbers (1st, 2nd, 3rd, etc)*

---

**Description**

Round values to integers and then display as ordinal values (e.g. 1st, 2nd, 3rd). Built-in rules are provided for English, French, and Spanish.

**Usage**

```
label_ordinal(
  prefix = "",
  suffix = "",
  big.mark = " ",
  rules = ordinal_english(),
  ...
)
```

```

ordinal_english()

ordinal_french(gender = c("masculin", "feminin"), plural = FALSE)

ordinal_spanish()

ordinal_format(
  prefix = "",
  suffix = "",
  big.mark = " ",
  rules = ordinal_english(),
  ...
)

ordinal(
  x,
  prefix = "",
  suffix = "",
  big.mark = " ",
  rules = ordinal_english(),
  ...
)

```

### Arguments

prefix, suffix	Symbols to display before and after value.
big.mark	Character used between every 3 digits to separate thousands.
rules	Named list of regular expressions, matched in order. Name gives suffix, and value specifies which numbers to match.
...	Other arguments passed on to <code>base::format()</code> .
gender	Masculin or feminin gender for French ordinal.
plural	Plural or singular for French ordinal.
x	A numeric vector to format.

### Value

All `label_()` functions return a "labelling" function, i.e. a function that takes a vector `x` and returns a character vector of `length(x)` giving a label for each input value.

Labelling functions are designed to be used with the `labels` argument of `ggplot2` scales. The examples demonstrate their use with `x` scales, but they work similarly for all scales, including those that generate legends rather than axes.

### Old interface

`ordinal()` and `format_ordinal()` are retired; please use `label_ordinal()` instead.

### See Also

Other labels for continuous scales: [label\\_bytes\(\)](#), [label\\_dollar\(\)](#), [label\\_number\\_auto\(\)](#), [label\\_number\\_si\(\)](#), [label\\_parse\(\)](#), [label\\_percent\(\)](#), [label\\_pvalue\(\)](#), [label\\_scientific\(\)](#)

**Examples**

```

demo_continuous(c(1, 5))
demo_continuous(c(1, 5), labels = label_ordinal())
demo_continuous(c(1, 5), labels = label_ordinal(rules = ordinal_french()))

# The rules are just a set of regular expressions that are applied in turn
ordinal_french()
ordinal_english()

# Note that ordinal rounds values, so you may need to adjust the breaks too
demo_continuous(c(1, 10))
demo_continuous(c(1, 10), labels = label_ordinal())
demo_continuous(c(1, 10),
  labels = label_ordinal(),
  breaks = breaks_width(2)
)

```

---

label\_parse

*Label with mathematical annotations*


---

**Description**

label\_parse() produces expression from strings by parsing them; label\_math() constructs expressions by replacing the pronoun .x with each string.

**Usage**

```

label_parse()

label_math(expr = 10^.x, format = force)

parse_format()

math_format(expr = 10^.x, format = force)

```

**Arguments**

expr	expression to use
format	another format function to apply prior to mathematical transformation - this makes it easier to use floating point numbers in mathematical expressions.

**Value**

All label\_() functions return a "labelling" function, i.e. a function that takes a vector x and returns a character vector of length(x) giving a label for each input value.

Labelling functions are designed to be used with the labels argument of ggplot2 scales. The examples demonstrate their use with x scales, but they work similarly for all scales, including those that generate legends rather than axes.

**Old interface**

parse\_format() and math\_format() was retired; please use label\_parse() and label\_math() instead.

**See Also**

[plotmath](#) for the details of mathematical formatting in R.

Other labels for continuous scales: [label\\_bytes\(\)](#), [label\\_dollar\(\)](#), [label\\_number\\_auto\(\)](#), [label\\_number\\_si\(\)](#), [label\\_ordinal\(\)](#), [label\\_percent\(\)](#), [label\\_pvalue\(\)](#), [label\\_scientific\(\)](#)

Other labels for discrete scales: [label\\_wrap\(\)](#)

**Examples**

```
# Use label_parse() with discrete scales
greek <- c("alpha", "beta", "gamma")
demo_discrete(greek)
demo_discrete(greek, labels = label_parse())

# Use label_math() with continuous scales
demo_continuous(c(1, 5))
demo_continuous(c(1, 5), labels = label_math(alpha[.x]))
```

---

label_percent	<i>Label percentages (2.5%, 50%, etc)</i>
---------------	---

---

**Description**

Label percentages (2.5%, 50%, etc)

**Usage**

```
label_percent(
  accuracy = NULL,
  scale = 100,
  prefix = "",
  suffix = "%",
  big.mark = " ",
  decimal.mark = ".",
  trim = TRUE,
  ...
)

percent_format(
  accuracy = NULL,
  scale = 100,
  prefix = "",
  suffix = "%",
  big.mark = " ",
  decimal.mark = ".",
  trim = TRUE,
  ...
)

percent(
  x,
  accuracy = NULL,
```

```

scale = 100,
prefix = "",
suffix = "%",
big.mark = " ",
decimal.mark = ".",
trim = TRUE,
...
)

```

### Arguments

accuracy	A number to round to. Use (e.g.) 0.01 to show 2 decimal places of precision. If NULL, the default, uses a heuristic that should ensure breaks have the minimum number of digits needed to show the difference between adjacent values. Applied to rescaled data.
scale	A scaling factor: x will be multiplied by scale before formatting. This is useful if the underlying data is very small or very large.
prefix	Symbols to display before and after value.
suffix	Symbols to display before and after value.
big.mark	Character used between every 3 digits to separate thousands.
decimal.mark	The character to be used to indicate the numeric decimal point.
trim	Logical, if FALSE, values are right-justified to a common width (see <code>base::format()</code> ).
...	Other arguments passed on to <code>base::format()</code> .
x	A numeric vector to format.

### Value

All `label_()` functions return a "labelling" function, i.e. a function that takes a vector x and returns a character vector of `length(x)` giving a label for each input value.

Labelling functions are designed to be used with the `labels` argument of `ggplot2` scales. The examples demonstrate their use with x scales, but they work similarly for all scales, including those that generate legends rather than axes.

### Old interface

`percent()` and `percent_format()` are retired; please use `label_percent()` instead.

### See Also

Other labels for continuous scales: `label_bytes()`, `label_dollar()`, `label_number_auto()`, `label_number_si()`, `label_ordinal()`, `label_parse()`, `label_pvalue()`, `label_scientific()`

### Examples

```

demo_continuous(c(0, 1))
demo_continuous(c(0, 1), labels = label_percent())

# Use prefix and suffix to create your own variants
french_percent <- label_percent(
  decimal.mark = ",",
  suffix = " %"
)
demo_continuous(c(0, .01), labels = french_percent)

```

---

label_pvalue	<i>Label p-values (e.g. &lt;0.001, 0.25, p &gt;= 0.99)</i>
--------------	--

---

### Description

Formatter for p-values, using "<" and ">" for p-values close to 0 and 1.

### Usage

```
label_pvalue(
  accuracy = 0.001,
  decimal.mark = ".",
  prefix = NULL,
  add_p = FALSE
)
```

```
pvalue_format(
  accuracy = 0.001,
  decimal.mark = ".",
  prefix = NULL,
  add_p = FALSE
)
```

```
pvalue(x, accuracy = 0.001, decimal.mark = ".", prefix = NULL, add_p = FALSE)
```

### Arguments

accuracy	A number to round to. Use (e.g.) 0.01 to show 2 decimal places of precision. If NULL, the default, uses a heuristic that should ensure breaks have the minimum number of digits needed to show the difference between adjacent values. Applied to rescaled data.
decimal.mark	The character to be used to indicate the numeric decimal point.
prefix	A character vector of length 3 giving the prefixes to put in front of numbers. The default values are c("<", "", ">") if add_p is TRUE and c("p<", "p=", "p>") if FALSE.
add_p	Add "p=" before the value?
x	A numeric vector to format.

### Value

All label\_() functions return a "labelling" function, i.e. a function that takes a vector x and returns a character vector of length(x) giving a label for each input value.

Labelling functions are designed to be used with the labels argument of ggplot2 scales. The examples demonstrate their use with x scales, but they work similarly for all scales, including those that generate legends rather than axes.

### Old interface

pvalue() and pvalue\_dollar() are retired; please use label\_pvalue() instead.

**See Also**

Other labels for continuous scales: [label\\_bytes\(\)](#), [label\\_dollar\(\)](#), [label\\_number\\_auto\(\)](#), [label\\_number\\_si\(\)](#), [label\\_ordinal\(\)](#), [label\\_parse\(\)](#), [label\\_percent\(\)](#), [label\\_scientific\(\)](#)

**Examples**

```
demo_continuous(c(0, 1))
demo_continuous(c(0, 1), labels = label_pvalue())
demo_continuous(c(0, 1), labels = label_pvalue(accuracy = 0.1))
demo_continuous(c(0, 1), labels = label_pvalue(add_p = TRUE))

# Or provide your own prefixes
prefix <- c("p < ", "p = ", "p > ")
demo_continuous(c(0, 1), labels = label_pvalue(prefix = prefix))
```

---

label_scientific	<i>Label numbers with scientific notation (e.g. 1e05, 1.5e-02)</i>
------------------	--

---

**Description**

Label numbers with scientific notation (e.g. 1e05, 1.5e-02)

**Usage**

```
label_scientific(
  digits = 3,
  scale = 1,
  prefix = "",
  suffix = "",
  decimal.mark = ".",
  trim = TRUE,
  ...
)
```

```
scientific_format(
  digits = 3,
  scale = 1,
  prefix = "",
  suffix = "",
  decimal.mark = ".",
  trim = TRUE,
  ...
)
```

```
scientific(
  x,
  digits = 3,
  scale = 1,
  prefix = "",
  suffix = "",
  decimal.mark = ".",
  ...
)
```

```

    trim = TRUE,
    ...
  )

```

### Arguments

digits	Number of digits to show before exponent.
scale	A scaling factor: $x$ will be multiplied by <code>scale</code> before formatting. This is useful if the underlying data is very small or very large.
prefix, suffix	Symbols to display before and after value.
decimal.mark	The character to be used to indicate the numeric decimal point.
trim	Logical, if FALSE, values are right-justified to a common width (see <code>base::format()</code> ).
...	Other arguments passed on to <code>base::format()</code> .
x	A numeric vector to format.

### Value

All `label_()` functions return a "labelling" function, i.e. a function that takes a vector  $x$  and returns a character vector of length( $x$ ) giving a label for each input value.

Labelling functions are designed to be used with the `labels` argument of `ggplot2` scales. The examples demonstrate their use with  $x$  scales, but they work similarly for all scales, including those that generate legends rather than axes.

### Old interface

`scientific_format()` and `scientific()` are retired; please use `label_scientific()`.

### See Also

Other labels for continuous scales: `label_bytes()`, `label_dollar()`, `label_number_auto()`, `label_number_si()`, `label_ordinal()`, `label_parse()`, `label_percent()`, `label_pvalue()`

Other labels for log scales: `label_bytes()`, `label_number_si()`

### Examples

```

demo_continuous(c(1, 10))
demo_continuous(c(1, 10), labels = label_scientific())
demo_continuous(c(1, 10), labels = label_scientific(digits = 3))

demo_log10(c(1, 1e9))

```

---

label\_wrap

*Label strings by wrapping across multiple lines*

---

### Description

Uses `strwrap()` to split long labels across multiple lines.

### Usage

```
label_wrap(width)
```

```
wrap_format(width)
```

### Arguments

`width`            Number of characters per line.

### Value

All `label_()` functions return a "labelling" function, i.e. a function that takes a vector `x` and returns a character vector of `length(x)` giving a label for each input value.

Labelling functions are designed to be used with the `labels` argument of `ggplot2` scales. The examples demonstrate their use with `x` scales, but they work similarly for all scales, including those that generate legends rather than axes.

### Old interface

`wrap_format()` is retired; please use `label_format()` instead.

### See Also

Other labels for discrete scales: [label\\_parse\(\)](#)

### Examples

```
x <- c(
  "this is a long label",
  "this is another long label",
  "this a label this is even longer"
)
demo_discrete(x)
demo_discrete(x, labels = label_wrap(10))
demo_discrete(x, labels = label_wrap(20))
```

---

linetype_pal	<i>Line type palette (discrete)</i>
--------------	-------------------------------------

---

**Description**

Based on a set supplied by Richard Pearson, University of Manchester

**Usage**

```
linetype_pal()
```

---

log_trans	<i>Log transformations</i>
-----------	----------------------------

---

**Description**

- `log_trans()`:  $\log(x)$
- `log1p()`:  $\log(x + 1)$
- `pseudo_log_trans()`: smoothly transition to linear scale around 0.

**Usage**

```
log_trans(base = exp(1))
log10_trans()
log2_trans()
log1p_trans()
pseudo_log_trans(sigma = 1, base = exp(1))
```

**Arguments**

base	base of logarithm
sigma	Scaling factor for the linear part of pseudo-log transformation.

**Examples**

```
plot(log2_trans(), xlim = c(0, 5))
plot(log_trans(), xlim = c(0, 5))
plot(log10_trans(), xlim = c(0, 5))

plot(log_trans(), xlim = c(0, 2))
plot(log1p_trans(), xlim = c(-1, 1))

# The pseudo-log is defined for all real numbers
plot(pseudo_log_trans(), xlim = c(-5, 5))
lines(log_trans(), xlim = c(0, 5), col = "red")
```

```
# For large positives numbrs it's very close to log
plot(pseudo_log_trans(), xlim = c(1, 20))
lines(log_trans(), xlim = c(1, 20), col = "red")
```

---

manual_pal	<i>Manual palette (discrete)</i>
------------	----------------------------------

---

### Description

Manual palette (discrete)

### Usage

```
manual_pal(values)
```

### Arguments

values            vector of values to be used as a palette.

---

minor_breaks_width	<i>Minor breaks</i>
--------------------	---------------------

---

### Description

Generate minor breaks between major breaks either spaced with a fixed width, or having a fixed number.

### Usage

```
minor_breaks_width(width, offset)
```

```
minor_breaks_n(n)
```

### Arguments

width	Distance between each break. Either a number, or for date/times, a single string of the form "n unit", e.g. "1 month", "5 days". Unit can be of one "sec", "min", "hour", "day", "week", "month", "year".
offset	Use if you don't want breaks to start at zero
n	number of breaks

### Examples

```
demo_log10(c(1, 1e6))
if (FALSE) {
  # Requires https://github.com/tidyverse/ggplot2/pull/3591
  demo_log10(c(1, 1e6), minor_breaks = minor_breaks_n(10))
}
```

---

muted	<i>Mute standard colour</i>
-------	-----------------------------

---

**Description**

Mute standard colour

**Usage**

```
muted(colour, l = 30, c = 70)
```

**Arguments**

colour	character vector of colours to modify
l	new luminance
c	new chroma

**Examples**

```
muted("red")
muted("blue")
show_col(c("red", "blue", muted("red"), muted("blue")))
```

---

oob	<i>Out of bounds handling</i>
-----	-------------------------------

---

**Description**

This set of functions modify data values outside a given range. The `oob_*`() functions are designed to be passed as the `oob` argument of `ggplot2` continuous and binned scales, with `oob_discard` being an exception.

These functions affect out of bounds values in the following ways:

- `oob_censor()` replaces out of bounds values with NAs. This is the default `oob` argument for continuous scales.
- `oob_censor_any()` acts like `oob_censor()`, but also replaces infinite values with NAs.
- `oob_squish()` replaces out of bounds values with the nearest limit. This is the default `oob` argument for binned scales.
- `oob_squish_any()` acts like `oob_squish()`, but also replaces infinite values with the nearest limit.
- `oob_squish_infinite()` only replaces infinite values by the nearest limit.
- `oob_keep()` does not adjust out of bounds values. In position scales, behaves as zooming limits without data removal.
- `oob_discard()` removes out of bounds values from the input. Not suitable for `ggplot2` scales.

## Usage

```
oob_censor(x, range = c(0, 1), only.finite = TRUE)
oob_censor_any(x, range = c(0, 1))
oob_discard(x, range = c(0, 1))
oob_squish(x, range = c(0, 1), only.finite = TRUE)
oob_squish_any(x, range = c(0, 1))
oob_squish_infinite(x, range = c(0, 1))
oob_keep(x, range = c(0, 1))
censor(x, range = c(0, 1), only.finite = TRUE)
discard(x, range = c(0, 1))
squish(x, range = c(0, 1), only.finite = TRUE)
squish_infinite(x, range = c(0, 1))
```

## Arguments

<code>x</code>	A numeric vector of values to modify.
<code>range</code>	A numeric vector of length two giving the minimum and maximum limit of the desired output range respectively.
<code>only.finite</code>	A logical of length one. When TRUE, only finite values are altered. When FALSE, also infinite values are altered.

## Details

The `oob_censor_any()` and `oob_squish_any()` functions are the same as `oob_censor()` and `oob_squish()` with the `only.finite` argument set to FALSE.

Replacing position values with NAs, as `oob_censor()` does, will typically lead to removal of those datapoints in ggplot.

Setting ggplot coordinate limits is equivalent to using `oob_keep()` in position scales.

## Value

Most `oob_()` functions return a vector of numerical values of the same length as the `x` argument, wherein out of bounds values have been modified. Only `oob_discard()` returns a vector of less than or of equal length to the `x` argument.

## Old interface

`censor()`, `squish()`, `squish_infinite()` and `discard()` are no longer recommended; please use `oob_censor()`, `oob_squish()`, `oob_squish_infinite()` and `oob_discard()` instead.

**Author(s)**

oob\_squish(): Homer Strong [homer.strong@gmail.com](mailto:homer.strong@gmail.com)

**Examples**

```
# Censoring replaces out of bounds values with NAs
oob_censor(c(-Inf, -1, 0.5, 1, 2, NA, Inf))
oob_censor_any(c(-Inf, -1, 0.5, 1, 2, NA, Inf))

# Squishing replaces out of bounds values with the nearest range limit
oob_squish(c(-Inf, -1, 0.5, 1, 2, NA, Inf))
oob_squish_any(c(-Inf, -1, 0.5, 1, 2, NA, Inf))
oob_squish_infinite(c(-Inf, -1, 0.5, 1, 2, NA, Inf))

# Keeping does not alter values
oob_keep(c(-Inf, -1, 0.5, 1, 2, NA, Inf))

# Discarding will remove out of bounds values
oob_discard(c(-Inf, -1, 0.5, 1, 2, NA, Inf))
```

---

probability\_trans      *Probability transformation*

---

**Description**

Probability transformation

**Usage**

```
probability_trans(distribution, ...)

logit_trans()

probit_trans()
```

**Arguments**

**distribution**      probability distribution. Should be standard R abbreviation so that "p" + distribution is a valid probability density function, and "q" + distribution is a valid quantile function.

**...**              other arguments passed on to distribution and quantile functions

**Examples**

```
plot(logit_trans(), xlim = c(0, 1))
plot(probit_trans(), xlim = c(0, 1))
```

---

Range	<i>Mutable ranges</i>
-------	-----------------------

---

**Description**

Mutable ranges have a two methods (train and reset), and make it possible to build up complete ranges with multiple passes.

---

reciprocal_trans	<i>Reciprocal transformation</i>
------------------	----------------------------------

---

**Description**

Reciprocal transformation

**Usage**

```
reciprocal_trans()
```

**Examples**

```
plot(reciprocal_trans(), xlim = c(0, 1))
```

---

rescale	<i>Rescale continuous vector to have specified minimum and maximum</i>
---------	--

---

**Description**

Rescale continuous vector to have specified minimum and maximum

**Usage**

```
rescale(x, to, from, ...)

## S3 method for class 'numeric'
rescale(x, to = c(0, 1), from = range(x, na.rm = TRUE, finite = TRUE), ...)

## S3 method for class 'dist'
rescale(x, to = c(0, 1), from = range(x, na.rm = TRUE, finite = TRUE), ...)

## S3 method for class 'logical'
rescale(x, to = c(0, 1), from = range(x, na.rm = TRUE, finite = TRUE), ...)

## S3 method for class 'POSIXt'
rescale(x, to = c(0, 1), from = range(x, na.rm = TRUE, finite = TRUE), ...)

## S3 method for class 'Date'
rescale(x, to = c(0, 1), from = range(x, na.rm = TRUE, finite = TRUE), ...)

## S3 method for class 'integer64'
rescale(x, to = c(0, 1), from = range(x, na.rm = TRUE), ...)
```

**Arguments**

x	continuous vector of values to manipulate.
to	output range (numeric vector of length two)
from	input range (vector of length two). If not given, is calculated from the range of x
...	other arguments passed on to methods

**Examples**

```
rescale(1:100)
rescale(runif(50))
rescale(1)
```

---

rescale_max	<i>Rescale numeric vector to have specified maximum</i>
-------------	---

---

**Description**

Rescale numeric vector to have specified maximum

**Usage**

```
rescale_max(x, to = c(0, 1), from = range(x, na.rm = TRUE))
```

**Arguments**

x	numeric vector of values to manipulate.
to	output range (numeric vector of length two)
from	input range (numeric vector of length two). If not given, is calculated from the range of x

**Examples**

```
rescale_max(1:100)
rescale_max(runif(50))
rescale_max(1)
```

---

`rescale_mid`*Rescale vector to have specified minimum, midpoint, and maximum*

---

**Description**

Rescale vector to have specified minimum, midpoint, and maximum

**Usage**

```
rescale_mid(x, to, from, mid, ...)  
  
## S3 method for class 'numeric'  
rescale_mid(x, to = c(0, 1), from = range(x, na.rm = TRUE), mid = 0, ...)  
  
## S3 method for class 'logical'  
rescale_mid(x, to = c(0, 1), from = range(x, na.rm = TRUE), mid = 0, ...)  
  
## S3 method for class 'dist'  
rescale_mid(x, to = c(0, 1), from = range(x, na.rm = TRUE), mid = 0, ...)  
  
## S3 method for class 'POSIXt'  
rescale_mid(x, to = c(0, 1), from = range(x, na.rm = TRUE), mid, ...)  
  
## S3 method for class 'Date'  
rescale_mid(x, to = c(0, 1), from = range(x, na.rm = TRUE), mid, ...)  
  
## S3 method for class 'integer64'  
rescale_mid(x, to = c(0, 1), from = range(x, na.rm = TRUE), mid = 0, ...)
```

**Arguments**

<code>x</code>	vector of values to manipulate.
<code>to</code>	output range (numeric vector of length two)
<code>from</code>	input range (vector of length two). If not given, is calculated from the range of <code>x</code>
<code>mid</code>	mid-point of input range
<code>...</code>	other arguments passed on to methods

**Examples**

```
rescale_mid(1:100, mid = 50.5)  
rescale_mid(runif(50), mid = 0.5)  
rescale_mid(1)
```

---

rescale_none	<i>Don't perform rescaling</i>
--------------	--------------------------------

---

**Description**

Don't perform rescaling

**Usage**

```
rescale_none(x, ...)
```

**Arguments**

x	numeric vector of values to manipulate.
...	all other arguments ignored

**Examples**

```
rescale_none(1:100)
```

---

rescale_pal	<i>Rescale palette (continuous)</i>
-------------	-------------------------------------

---

**Description**

Just rescales the input to the specific output range. Useful for alpha, size, and continuous position.

**Usage**

```
rescale_pal(range = c(0.1, 1))
```

**Arguments**

range	Numeric vector of length two, giving range of possible values. Should be between 0 and 1.
-------	---

---

reverse_trans	<i>Reverse transformation</i>
---------------	-------------------------------

---

**Description**

Reverse transformation

**Usage**

```
reverse_trans()
```

**Examples**

```
plot(reverse_trans(), xlim = c(-1, 1))
```

---

seq\_gradient\_pal      *Sequential colour gradient palette (continuous)*

---

### Description

Sequential colour gradient palette (continuous)

### Usage

```
seq_gradient_pal(low = mns1("10B 4/6"), high = mns1("10R 4/6"), space = "Lab")
```

### Arguments

low	colour for low end of gradient.
high	colour for high end of gradient.
space	colour space in which to calculate gradient. Must be "Lab" - other values are deprecated.

### Examples

```
x <- seq(0, 1, length.out = 25)
show_col(seq_gradient_pal()(x))
show_col(seq_gradient_pal("white", "black")(x))

library(munsell)
show_col(seq_gradient_pal("white", mns1("10R 4/6"))(x))
```

---

shape\_pal      *Shape palette (discrete)*

---

### Description

Shape palette (discrete)

### Usage

```
shape_pal(solid = TRUE)
```

### Arguments

solid	should shapes be solid or not?
-------	--------------------------------

---

sqrt_trans	<i>Square-root transformation</i>
------------	-----------------------------------

---

**Description**

This is the variance stabilising transformation for the Poisson distribution.

**Usage**

```
sqrt_trans()
```

**Examples**

```
plot(sqrt_trans(), xlim = c(0, 5))
```

---

time_trans	<i>Transformation for date-times (class POSIXt)</i>
------------	---

---

**Description**

Transformation for date-times (class POSIXt)

**Usage**

```
time_trans(tz = NULL)
```

**Arguments**

**tz**                      Optionally supply the time zone. If NULL, the default, the time zone will be extracted from first input with a non-null tz.

**Examples**

```
hours <- seq(ISOdate(2000,3,20, tz = ""), by = "hour", length.out = 10)
t <- time_trans()
t$transform(hours)
t$inverse(t$transform(hours))
t$format(t$breaks(range(hours)))
```

---

train_continuous	<i>Train (update) a continuous scale</i>
------------------	--

---

**Description**

Strips attributes and always returns a numeric vector

**Usage**

```
train_continuous(new, existing = NULL)
```

**Arguments**

new	New data to add to scale
existing	Optional existing scale to update

---

train_discrete	<i>Train (update) a discrete scale</i>
----------------	--

---

**Description**

Train (update) a discrete scale

**Usage**

```
train_discrete(new, existing = NULL, drop = FALSE, na.rm = FALSE)
```

**Arguments**

new	New data to add to scale
existing	Optional existing scale to update
drop	TRUE, will drop factor levels not associated with data
na.rm	If TRUE, will remove missing values

---

viridis_pal	<i>Viridis palette</i>
-------------	------------------------

---

**Description**

Viridis palette

**Usage**

```
viridis_pal(alpha = 1, begin = 0, end = 1, direction = 1, option = "D")
```

**Arguments**

alpha	The alpha transparency, a number in [0,1], see argument alpha in <a href="#">hsv</a> .
begin	The (corrected) hue in [0,1] at which the viridis colormap begins.
end	The (corrected) hue in [0,1] at which the viridis colormap ends.
direction	Sets the order of colors in the scale. If 1, the default, colors are ordered from darkest to lightest. If -1, the order of colors is reversed.
option	A character string indicating the colormap option to use. Four options are available: "magma" (or "A"), "inferno" (or "B"), "plasma" (or "C"), "viridis" (or "D", the default option) and "cividis" (or "E").

**References**

<https://bids.github.io/colormap/>

**Examples**

```
show_col(viridis_pal()(10))
show_col(viridis_pal(direction = -1)(6))
show_col(viridis_pal(begin = 0.2, end = 0.8)(4))
show_col(viridis_pal(option = "plasma")(6))
```

---

yj_trans	<i>Yeo-Johnson transformation</i>
----------	-----------------------------------

---

**Description**

The Yeo-Johnson transformation is a flexible transformation that is similar to Box-Cox, [boxcox\\_trans\(\)](#), but does not require input values to be greater than zero.

**Usage**

```
yj_trans(p)
```

**Arguments**

p	Transformation exponent, $\lambda$ .
---	--------------------------------------

**Details**

The transformation takes one of four forms depending on the values of  $y$  and  $\lambda$ .

- $y \geq 0$  and  $\lambda \neq 0$ :  $y^{(\lambda)} = \frac{(y+1)^\lambda - 1}{\lambda}$
- $y \geq 0$  and  $\lambda = 0$ :  $y^{(\lambda)} = \ln(y + 1)$
- $y < 0$  and  $\lambda \neq 2$ :  $y^{(\lambda)} = -\frac{(-y+1)^{(2-\lambda)} - 1}{2-\lambda}$
- $y < 0$  and  $\lambda = 2$ :  $y^{(\lambda)} = -\ln(-y + 1)$

**References**

Yeo, I., & Johnson, R. (2000). A New Family of Power Transformations to Improve Normality or Symmetry. *Biometrika*, 87(4), 954-959. <http://www.jstor.org/stable/2673623>

**Examples**

```
plot(yj_trans(-1), xlim = c(-10, 10))
plot(yj_trans(0), xlim = c(-10, 10))
plot(yj_trans(1), xlim = c(-10, 10))
plot(yj_trans(2), xlim = c(-10, 10))
```

---

zero\_range

*Determine if range of vector is close to zero, with a specified tolerance*

---

**Description**

The machine epsilon is the difference between 1.0 and the next number that can be represented by the machine. By default, this function uses  $\text{epsilon} * 1000$  as the tolerance. First it scales the values so that they have a mean of 1, and then it checks if the difference between them is larger than the tolerance.

**Usage**

```
zero_range(x, tol = 1000 * .Machine$double.eps)
```

**Arguments**

`x` numeric range: vector of length 2  
`tol` A value specifying the tolerance.

**Value**

logical TRUE if the relative difference of the endpoints of the range are not distinguishable from 0.

**Examples**

```
eps <- .Machine$double.eps
zero_range(c(1, 1 + eps))      # TRUE
zero_range(c(1, 1 + 99 * eps)) # TRUE
zero_range(c(1, 1 + 1001 * eps)) # FALSE - Crossed the tol threshold
zero_range(c(1, 1 + 2 * eps), tol = eps) # FALSE - Changed tol

# Scaling up or down all the values has no effect since the values
# are rescaled to 1 before checking against tol
zero_range(100000 * c(1, 1 + eps))      # TRUE
zero_range(100000 * c(1, 1 + 1001 * eps)) # FALSE
zero_range(.00001 * c(1, 1 + eps))      # TRUE
zero_range(.00001 * c(1, 1 + 1001 * eps)) # FALSE

# NA values
zero_range(c(1, NA)) # NA
zero_range(c(1, NaN)) # NA

# Infinite values
zero_range(c(1, Inf)) # FALSE
zero_range(c(-Inf, Inf)) # FALSE
zero_range(c(Inf, Inf)) # TRUE
```

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