# Package 'santoku' 

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Maintainer David Hugh-Jones [davidhughjones@gmail.com](mailto:davidhughjones@gmail.com)
Description A tool for cutting data into intervals. Allows singleton intervals. Always includes the whole range of data by default. Flexible labelling. Convenience functions for cutting by quantiles etc. Handles dates, times, units and other vectors.

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Author David Hugh-Jones [aut, cre], Daniel Possenriede [ctb]

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santoku-package A versatile cutting tool for $R$

## Description

santoku is a tool for cutting data into intervals. It provides the function chop(), which is similar to base R's cut () or Hmisc: : cut2(). chop ( $x$, breaks) takes a vector $x$ and returns a factor of the same length, coding which interval each element of $x$ falls into.

## Details

Here are some advantages of santoku:

- By default, chop() always covers the whole range of the data, so you won't get unexpected NA values.
- Unlike cut() or cut2(), chop() can handle single values as well as intervals. For example, chop $(x$, breaks $=c(1,2,2,3))$ will create a separate factor level for values exactly equal to 2 .
- Flexible and easy labelling.
- Convenience functions for creating quantile intervals, evenly-spaced intervals or equal-sized groups.
- Convenience functions to quickly tabulate chopped data.
- Can chop numbers, dates, date-times and other objects.

These advantages make santoku especially useful for exploratory analysis, where you may not know the range of your data in advance.
To get started, read the vignette:
vignette("santoku")
For more details, start with the documentation for chop().

## Author(s)

Maintainer: David Hugh-Jones [davidhughjones@gmail.com](mailto:davidhughjones@gmail.com)
Other contributors:

- Daniel Possenriede [possenriede@gmail.com](mailto:possenriede@gmail.com) [contributor]


## See Also

Useful links:

- https://github.com/hughjonesd/santoku
- https://hughjonesd.github.io/santoku/
- Report bugs at https://github.com/hughjonesd/santoku/issues

```
breaks-class Class representing a set of intervals
```


## Description

Class representing a set of intervals

## Usage

\#\# S3 method for class 'breaks'
format(x, ...)
\#\# S3 method for class 'breaks'
print(x, ...)
is.breaks(x, ...)

## Arguments

| $x$ | A breaks object |
| :--- | :--- |
| $\ldots$. | Unused |

brk_default Create a standard set of breaks

## Description

Create a standard set of breaks

## Usage

brk_default(breaks)

## Arguments

> breaks A numeric vector.

## Value

A function which returns an object of class breaks.

## Examples

chop(1:10, c(2, 5, 8))
chop(1:10, brk_default(c(2, 5, 8)))

```
brk_manual
Create a breaks object manually
```


## Description

Create a breaks object manually

## Usage

brk_manual(breaks, left_vec)

## Arguments

breaks A vector, which must be sorted.
left_vec A logical vector, the same length as breaks. Specifies whether each break is left-closed or right-closed.

## Details

All breaks must be closed on exactly one side, like . . , x) [x, ... (left-closed) or . .., x) [x, ... (right-closed).
For example, if breaks $=1: 3$ and left $=c($ TRUE, FALSE, TRUE $)$, then the resulting intervals are

```
T F T
[ 1, 2 ] ( 2, 3)
```

Singleton breaks are created by repeating a number in breaks. Singletons must be closed on both sides, so if there is a repeated number at indices $i, i+1$, left[i] must be TRUE and left[ $i+1]$ must be FALSE.

## Value

A function which returns an object of class breaks.

## Examples

lbrks <- brk_manual(1:3, rep(TRUE, 3))

```
chop(1:3, lbrks, extend = FALSE)
```

rbrks <- brk_manual(1:3, rep(FALSE, 3))
chop(1:3, rbrks, extend = FALSE)
brks_singleton <- brk_manual(
$c(1,2,2,3)$,
c(TRUE, TRUE, FALSE, TRUE))
chop(1:3, brks_singleton, extend = FALSE)

```
brk_width-for-datetime
```


## Equal-width intervals for dates or datetimes

## Description

brk_width() can be used with time interval classes from base R or the lubridate package.

## Usage

\#\# S3 method for class 'Duration'
brk_width(width, start)

## Arguments

width A scalar difftime, Period or Duration object.
start A scalar of class Date or POSIXct. Can be omitted.

## Details

If width is a Period, lubridate: :add_with_rollback() is used to calculate the widths. This can be useful for e.g. calendar months.

## Examples

```
if (requireNamespace("lubridate")) {
    year2001 <- as.Date("2001-01-01") + 0:364
    tab_width(year2001, months(1),
            labels = lbl_discrete(" to ", fmt = "%e %b %y"))
}
```

chop Cut data into intervals

## Description

chop() cuts $x$ into intervals. It returns a factor of the same length as $x$, representing which interval contains each element of $x$. kiru() is an alias for chop. $\operatorname{tab}()$ calls chop() and returns a contingency table() from the result.

## Usage

```
chop(
    x,
    breaks,
    labels = lbl_intervals(),
    extend = NULL,
    left = TRUE,
    close_end = TRUE,
    raw = NULL,
    drop = TRUE
)
kiru(
    x,
    breaks,
    labels = lbl_intervals(),
    extend = NULL,
    left = TRUE,
    close_end = TRUE,
    raw = NULL,
    drop = TRUE
)
```

tab(
chop

```
    x,
    breaks,
    labels = lbl_intervals(),
    extend = NULL,
    left = TRUE,
    close_end = TRUE,
    raw = NULL,
    drop = TRUE
    )
```


## Arguments

## x

A vector.
breaks A numeric vector of cut-points or a function to create cut-points from x .
labels A character vector of labels or a function to create labels.
extend Logical. If TRUE, always extend breaks to +/-Inf. If NULL, extend breaks to $\min (x)$ and/or $\max (x)$ only if necessary. If NULL, never extend.
left Logical. Left-closed or right-closed breaks?
close_end Logical. Close last break at right? (If left is FALSE, close first break at left?)
raw Logical. Use raw values in labels?
drop Logical. Drop unused levels from the result?

## Details

x may be a numeric vector, or more generally, any vector which can be compared with < and == (see Ops). In particular Date and date-time objects are supported. Character vectors are supported with a warning.

## Breaks:

breaks may be a vector or a function.
If it is a vector, breaks gives the break endpoints. Repeated values create singleton intervals. For example breaks $=c(1,3,3,5)$ creates 3 intervals: $[1,3),\{3\}$ and $(3,5]$.
If breaks is a function, it is called with the $x$, extend, left and close_end arguments, and should return an object of class breaks. Use brk_* functions to create a variety of data-dependent breaks.
Names of breaks may be used for labels. See "Labels" below.

## Options for breaks:

By default, left-closed intervals are created. If left is FALSE, right-closed intervals are created.
If close_end is TRUE the final break (or first break if left is FALSE) will be closed at both ends. This guarantees that all values x with min(breaks) <= $\mathrm{x}<=\max$ (breaks) are included in the intervals.
Before version 0.9.0, close_end was FALSE by default, and also behaved differently with respect to extended breaks: see "Extending intervals" below.
Using mathematical set notation:

- If left is TRUE and close_end is TRUE, breaks will look like [b1, b2), [b2, b3) ... [b_n-1, b_n].
- If left is FALSE and close_end is TRUE, breaks will look like [b1, b2], (b2, b3] ... (b_n-1, b_n].
- If left is TRUE and close_end is FALSE, all breaks will look like ...[b1, b2) ....
- If left is FALSE and close_end is FALSE, all breaks will look like ... (b1, b2] ....


## Extending intervals:

If extend is TRUE, intervals will be extended to [-Inf, min(breaks)) and (max(breaks), Inf].
If extend is NULL (the default), intervals will be extended to [min(x), min(breaks)) and (max (breaks), $\max (x)]$, only if necessary - i.e. if elements of $x$ would be below or above the unextended breaks. close_end is applied after breaks are extended, i.e. always to the very last or very first break. This is a change from previous behaviour. Up to version 0.8.0, close_end was applied to the userspecified intervals, then extend was applied. Note that if breaks are extended, then the extended break is always closed anyway.

## Labels:

labels may be a character vector. It should have the same length as the (possibly extended) number of intervals. Alternatively, labels may be a lbl_* function such as lbl_seq().
If breaks is a named vector, then non-zero-length names of breaks will be used as labels for the interval starting at the corresponding element. This overrides the labels argument (but unnamed breaks will still use labels). This feature is [Experimental].
If labels is NULL, then integer codes will be returned instead of a factor.
If raw is TRUE, labels will show the actual numbers calculated by breaks. If raw is FALSE then labels may show other objects, such as quantiles for chop_quantiles() and friends, proportions of the range for chop_proportions(), or standard deviations for chop_mean_sd().
If raw is NULL then $1 b l_{-} *$ functions will use their default (usually FALSE). Otherwise, raw argument to chop() overrides raw arguments passed into lbl_* functions directly.

## Miscellaneous:

NA values in $x$, and values which are outside the extended endpoints, return NA.
kiru() is a synonym for chop(). If you load \{tidyr\}, you can use it to avoid confusion with tidyr: : chop().
Note that chop(), like all of R, uses binary arithmetic. Thus, numbers may not be exactly equal to what you think they should be. There is an example below.

## Value

chop() returns a factor of the same length as $x$, representing the intervals containing the value of x.
$t a b()$ returns a contingency table().

## See Also

base: :cut(), non-standard-types for chopping objects that aren't numbers.
Other chopping functions: chop_equally(), chop_evenly(), chop_fn(), chop_mean_sd(), chop_n(), chop_proportions(), chop_quantiles(), chop_width(), fillet()

## Examples

```
chop(1:7, c(2, 4, 6))
chop(1:7, c(2, 4, 6), extend = FALSE)
# Repeat a number for a singleton break:
chop(1:7, c(2, 4, 4, 6))
chop(1:7, c(2, 4, 6), left = FALSE)
chop(1:7, c(2, 4, 6), close_end = FALSE)
chop(1:7, brk_quantiles(c(0.25, 0.75)))
# A single break is fine if `extend` is not `FALSE`:
chop(1:7, 4)
# Floating point inaccuracy:
chop(0.3/3, c(0, 0.1, 0.1, 1), labels = c("< 0.1", "0.1", "> 0.1"))
# -- Labels --
chop(1:7, c(Lowest = 1, Low = 2, Mid = 4, High = 6))
chop(1:7, c(2, 4, 6), labels = c("Lowest", "Low", "Mid", "High"))
chop(1:7, c(2, 4, 6), labels = lbl_dash())
# Mixing names and other labels:
chop(1:7, c("<2" = 1, 2, 4, ">=6" = 6), labels = lbl_dash())
# -- Non-standard types --
chop(as.Date("2001-01-01") + 1:7, as.Date("2001-01-04"))
suppressWarnings(chop(LETTERS[1:7], "D"))
tab(1:10, c(2, 5, 8))
```

    chop_equally Chop equal-sized groups
    
## Description

chop_equally() chops $x$ into groups with an equal number of elements.

## Usage

```
chop_equally(
        x,
        groups,
        ...,
        labels = lbl_intervals(),
        left = is.numeric(x),
        close_end = TRUE,
        raw = TRUE
    )
    brk_equally(groups)
    tab_equally(x, groups, ..., left = is.numeric(x), raw = TRUE)
```


## Arguments

x
groups Number of groups.
... Passed to chop().
labels A character vector of labels or a function to create labels.
left Logical. Left-closed or right-closed breaks?
close_end Logical. Close last break at right? (If left is FALSE, close first break at left?)
raw Logical. Use raw values in labels?

## Details

chop_equally() uses brk_quantiles() under the hood. If $x$ has duplicate elements, you may get fewer groups than requested. If so, a warning will be emitted. See the examples.

## Value

chop_* functions return a factor of the same length as $x$.
brk_* functions return a function to create breaks.
tab_* functions return a contingency table().

## See Also

Other chopping functions: chop(), chop_evenly(), chop_fn(), chop_mean_sd(), chop_n(), chop_proportions(), chop_quantiles(), chop_width(), fillet()

## Examples

```
chop_equally(1:10, 5)
# You can't always guarantee `groups` groups:
dupes <- c(1, 1, 1, 2, 3, 4, 4, 4)
quantile(dupes, 0:4/4)
chop_equally(dupes, 4)
```

```
chop_evenly Chop into equal-width intervals
```


## Description

chop_evenly () chops $x$ into intervals intervals of equal width.

## Usage

```
chop_evenly(x, intervals, ..., close_end = TRUE)
brk_evenly(intervals)
    tab_evenly(x, intervals, ...)
```


## Arguments

| $x$ | A vector. |
| :--- | :--- |
| intervals | Integer: number of intervals to create. |
| $\ldots$ | Passed to chop(). |
| close_end | Logical. Close last break at right? (If left is FALSE, close first break at left?) |

## Details

chop_evenly() sets close_end = TRUE by default.

## Value

chop_* functions return a factor of the same length as $x$.
brk_* functions return a function to create breaks.
tab_* functions return a contingency table().

## See Also

Other chopping functions: chop(), chop_equally(), chop_fn(), chop_mean_sd(), chop_n(), chop_proportions(), chop_quantiles(), chop_width(), fillet()

## Examples

```
chop_evenly(0:10, 5)
```

chop_fn Chop using an existing function

## Description

chop_fn() is a convenience wrapper: chop_fn(x,foo, ...) is the same as chop(x, foo(x, ...)).

## Usage

```
    chop_fn(
```

        x ,
        fn,
        ...,
        extend \(=\) NULL,
        left = TRUE,
        close_end = TRUE,
        raw = NULL,
        drop = TRUE
    )
    brk_fn(fn, ...)
    tab_fn(
        x ,
        fn,
        ...,
        extend \(=\) NULL,
        left = TRUE,
        close_end = TRUE,
        raw = NULL,
        drop = TRUE
    )
    
## Arguments

x
fn
...
extend
left
close_end
raw
drop

A vector.
A function which returns a numeric vector of breaks.
Further arguments to fn
Logical. If TRUE, always extend breaks to +/-Inf. If NULL, extend breaks to $\min (x)$ and/or $\max (x)$ only if necessary. If NULL, never extend.
Logical. Left-closed or right-closed breaks?
Logical. Close last break at right? (If left is FALSE, close first break at left?)
Logical. Use raw values in labels?
Logical. Drop unused levels from the result?

## Value

chop_* functions return a factor of the same length as $x$.
brk_* functions return a function to create breaks.
tab_* functions return a contingency table().

## See Also

Other chopping functions: $\operatorname{chop}()$, chop_equally(), chop_evenly (), chop_mean_sd(), chop_n(), chop_proportions(), chop_quantiles(), chop_width(), fillet()

## Examples

```
if (requireNamespace("scales")) {
    chop_fn(rlnorm(10), scales::breaks_log(5))
    # same as
    # x <- rlnorm(10)
    # chop(x, scales::breaks_log(5)(x))
}
```

chop_mean_sd Chop by standard deviations

## Description

Intervals are measured in standard deviations on either side of the mean.

## Usage

chop_mean_sd(x, sds = 1:3, ..., raw = FALSE, sd = deprecated())
brk_mean_sd(sds = 1:3, sd = deprecated())
tab_mean_sd(x, sds = 1:3, ..., raw = FALSE)

## Arguments

x
sds
... Passed to chop().
sd [Deprecated]
raw Logical. Use raw values in labels?
A vector.
Positive numeric vector of standard deviations.

## Details

In version 0.7.0, these functions changed to specifying sds as a vector. To chop 1,2 and 3 standard deviations around the mean, write chop_mean_sd(x, sds =1:3) instead of chop_mean_sd(x, sd $=3$ ).

## Value

chop_* functions return a factor of the same length as $x$.
brk_* functions return a function to create breaks.
tab_* functions return a contingency table().

## See Also

Other chopping functions: chop(), chop_equally(), chop_evenly(), chop_fn(), chop_n(), chop_proportions(), chop_quantiles(), chop_width(), fillet()

## Examples

```
    chop_mean_sd(1:10)
```

    \(\operatorname{chop}(1: 10\), brk_mean_sd())
    tab_mean_sd(1:10)
    chop_n Chop into fixed-sized groups
    
## Description

chop_n() creates intervals containing a fixed number of elements.

## Usage

```
chop_n(x, n, ..., close_end = TRUE, tail = "split")
    brk_n(n, tail = "split")
    tab_n(x, n, ..., tail = "split")
```


## Arguments

x
$\mathrm{n} \quad$ Integer. Number of elements in each interval.
... Passed to chop().
close_end Logical. Close last break at right? (If left is FALSE, close first break at left?)
tail String. What to do if the final interval has fewer than $n$ elements? "split" to keep it separate. "merge" to merge it with the neighbouring interval.

## Details

The algorithm guarantees that intervals contain no more than $n$ elements, so long as there are no duplicates in x and tail = "split". It also guarantees that intervals contain no fewer than n elements, except possibly the last interval (or first interval if left is FALSE).

To ensure that all intervals contain at least n elements (so long as there are at least n elements in x !) set tail = "merge".

If tail = "split" and there are intervals containing duplicates with more than n elements, a warning is given.

## Value

chop_* functions return a factor of the same length as $x$.
brk_* functions return a function to create breaks.
tab_* functions return a contingency table().

## See Also

Other chopping functions: chop(), chop_equally(), chop_evenly(), chop_fn(), chop_mean_sd(), chop_proportions(), chop_quantiles(), chop_width(), fillet()

## Examples

```
chop_n(1:10, 5)
chop_n(1:5, 2)
chop_n(1:5, 2, tail = "merge")
# too many duplicates
x <- rep(1:2, each = 3)
chop_n(x, 2)
tab_n(1:10, 5)
# fewer elements in one group
tab_n(1:10, 4)
```

chop_pretty Chop using pretty breakpoints

## Description

chop_pretty () uses base: : pretty() to calculate breakpoints which are 1,2 or 5 times a power of 10 . These look nice in graphs.

## Usage

chop_pretty (x, n = 5, ...)
brk_pretty (n = 5, ...)
tab_pretty (x, n = 5, ...)

## Arguments

X
$\mathrm{n} \quad$ Positive integer passed to base: :pretty (). How many intervals to chop into?
... Passed to chop() by chop_pretty() and tab_pretty(); passed to base: :pretty() by brk_pretty().

## Details

base: : pretty () tries to return $n+1$ breakpoints, i.e. $n$ intervals, but note that this is not guaranteed. There are methods for Date and POSIXct objects.

For fine-grained control over base: : pretty () parameters, use chop(x, brk_pretty(...)).

## Value

chop_* functions return a factor of the same length as $x$.
brk_* functions return a function to create breaks.
tab_* functions return a contingency table().

## Examples

chop_pretty (1:10)
chop(1:10, brk_pretty ( $\mathrm{n}=5$, high.u.bias $=0$ )
tab_pretty (1:10)
chop_proportions Chop into proportions of the range of $x$

## Description

chop_proportions() chops $x$ into proportions of its range, excluding infinite values.

## Usage

chop_proportions(x, proportions, ..., raw = TRUE)
brk_proportions(proportions)
tab_proportions(x, proportions, ..., raw = TRUE)

## Arguments

X
proportions Numeric vector between 0 and 1: proportions of x's range. If proportions has names, these will be used for labels.
... Passed to chop().
raw Logical. Use raw values in labels?

## Details

By default, labels show the raw numeric endpoints. To label intervals by the proportions, use raw $=$ FALSE.

## Value

chop_* functions return a factor of the same length as $x$.
brk_* functions return a function to create breaks.
tab_* functions return a contingency table().

## See Also

Other chopping functions: chop(), chop_equally(), chop_evenly(), chop_fn(), chop_mean_sd(), chop_n(), chop_quantiles(), chop_width(), fillet()

## Examples

chop_proportions(0:10, c(0.2, 0.8))
chop_proportions $0: 10, \mathrm{c}($ Low $=0, \mathrm{Mid}=0.2, \mathrm{High}=0.8)$ )

```
chop_quantiles Chop by quantiles
```


## Description

chop_quantiles() chops data by quantiles. chop_deciles() is a convenience function which chops into deciles.

```
Usage
    chop_quantiles(
        x,
        probs,
        ...,
        left = is.numeric(x),
        raw = FALSE,
        weights = NULL
)
```

```
chop_deciles(x, ...)
brk_quantiles(probs, ..., weights = NULL)
tab_quantiles(x, probs, ..., left = is.numeric(x), raw = FALSE)
tab_deciles(x, ...)
```


## Arguments

$x$
probs
... For chop_quantiles, passed to chop(). For brk_quantiles(), passed to stats: :quantile() or Hmisc: :wtd.quantile().
left Logical. Left-closed or right-closed breaks?
raw Logical. Use raw values in labels?
weights NULL or numeric vector of same length as $x$. If not NULL, Hmisc: : wtd. quantile() is used to calculate weighted quantiles.

## Details

For non-numeric $x$, left is set to FALSE by default. This works better for calculating "type 1" quantiles, since they round down. See stats: :quantile().
If $x$ contains duplicates, consecutive quantiles may be the same number so that some intervals get merged.

## Value

chop_* functions return a factor of the same length as $x$.
brk_* functions return a function to create breaks.
tab_* functions return a contingency table().

## See Also

Other chopping functions: chop(), chop_equally(), chop_evenly(), chop_fn(), chop_mean_sd(), chop_n(), chop_proportions(), chop_width(), fillet()

## Examples

```
chop_quantiles(1:10, 1:3/4)
chop_quantiles(1:10, c(Q1 = 0, Q2 = 0.25, Q3 = 0.5, Q4 = 0.75))
chop(1:10, brk_quantiles(1:3/4))
chop_deciles(1:10)
```

```
    # to label by the quantiles themselves:
    chop_quantiles(1:10, 1:3/4, raw = TRUE)
    # duplicates:
    tab_quantiles(c(1, 1, 1, 2, 3), 1:5/5)
    set.seed(42)
    tab_quantiles(rnorm(100), probs = 1:3/4, raw = TRUE)
```

    chop_width
    Chop into fixed-width intervals
    
## Description

chop_width() chops $x$ into intervals of fixed width.

## Usage

chop_width(x, width, start, ..., left = sign(width) > 0)
brk_width(width, start)
\#\# Default S3 method:
brk_width(width, start)
tab_width(x, width, start, ..., left = sign(width) > 0)

## Arguments

x
width
start Starting point for intervals. By default the smallest finite $\times$ (largest if width is negative).
... Passed to chop().
left
A vector.
Width of intervals.

Logical. Left-closed or right-closed breaks?

## Details

If width is negative, chop_width() sets left = FALSE and intervals will go downwards from start.

## Value

chop_* functions return a factor of the same length as $x$.
brk_* functions return a function to create breaks.
tab_* functions return a contingency table().

## See Also

brk_width-for-datetime
Other chopping functions: chop(), chop_equally(), chop_evenly(), chop_fn(), chop_mean_sd(), chop_n(), chop_proportions(), chop_quantiles(), fillet()

## Examples

```
    chop_width(1:10, 2)
    chop_width(1:10, 2, start = 0)
    chop_width(1:9, -2)
    chop(1:10, brk_width(2, 0))
    tab_width(1:10, 2, start = 0)
```

    exactly Define singleton intervals explicitly
    
## Description

exactly () duplicates its input. It lets you define singleton intervals like this: chop ( $x, \mathrm{c}(1$, exactly(2), 3)). This is the same as $\operatorname{chop}(x, c(1,2,2,3))$ but conveys your intent more clearly.

## Usage

```
exactly(x)
```


## Arguments

x
A numeric vector.

## Value

The same as rep (x, each = 2).

## Examples

```
chop(1:10, c(2, exactly(5), 8))
# same:
chop(1:10, c(2, 5, 5, 8))
```

fillet Chop data precisely (for programmers)

## Description

fillet () calls chop() with extend = FALSE and drop = FALSE. This ensures that you get only the breaks and labels you ask for. When programming, consider using fillet() instead of chop().

## Usage

```
fillet(
        x,
        breaks,
        labels = lbl_intervals(),
        left = TRUE,
        close_end = TRUE,
        raw = NULL
    )
```


## Arguments

x
breaks A numeric vector of cut-points or a function to create cut-points from x .
labels A character vector of labels or a function to create labels.
left Logical. Left-closed or right-closed breaks?
close_end Logical. Close last break at right? (If left is FALSE, close first break at left?)
raw Logical. Use raw values in labels?

## Value

fillet() returns a factor of the same length as $x$, representing the intervals containing the value of $x$.

## See Also

Other chopping functions: chop(), chop_equally(), chop_evenly(), chop_fn(), chop_mean_sd(), chop_n(), chop_proportions(), chop_quantiles(), chop_width()

## Examples

fillet (1:10, c(2, 5, 8))

## Description

This label style is user-friendly, but doesn't distinguish between left- and right-closed intervals. It's good for continuous data where you don't expect points to be exactly on the breaks.

## Usage

```
    lbl_dash(
```

        symbol = em_dash(),
        fmt = NULL,
        single = "\{l\}",
        first = NULL,
        last = NULL,
        raw = FALSE
    )
    
## Arguments

symbol String: symbol to use for the dash.
fmt String, list or function. A format for break endpoints.
single Glue string: label for singleton intervals. See lbl_glue() for details.
first Glue string: override label for the first category. Write e.g. first = "<\{r\}" to create a label like " $<18$ ". See lbl_glue() for details.
last String: override label for the last category. Write e.g. last = ">\{l\}" to create a label like ">65". See lbl_glue() for details.
raw [Deprecated]. Use the raw argument to chop() instead.

## Details

If you don't want unicode output, use lbl_dash("-").

## Value

A function that creates a vector of labels.

## Formatting endpoints

If fmt is not NULL then it is used to format the endpoints.

- If fmt is a string, then numeric endpoints will be formatted by sprintf(fmt, breaks); other endpoints, e.g. Date objects, will be formatted by format (breaks, fmt).
- If fmt is a list, then it will be used as arguments to format.
- If fmt is a function, it should take a vector of numbers (or other objects that can be used as breaks) and return a character vector. It may be helpful to use functions from the \{scales\} package, e.g. scales: : label_comma().


## See Also

Other labelling functions: lbl_discrete(), lbl_endpoints(), lbl_glue(), lbl_intervals(), lbl_manual(), lbl_midpoints(), lbl_seq()

## Examples

```
    chop(1:10, c(2, 5, 8), lbl_dash())
    chop(1:10, c(2, 5, 8), lbl_dash(" to ", fmt = "%.1f"))
    chop(1:10, c(2, 5, 8), lbl_dash(first = "<{r}"))
    pretty <- function (x) prettyNum(x, big.mark = ",", digits = 1)
    chop(runif(10) * 10000, c(3000, 7000), lbl_dash(" to ", fmt = pretty))
```

    lbl_discrete Label discrete data
    
## Description

lbl_discrete() creates labels for discrete data, such as integers. For example, breaks c (1, 3, 4, 6,7 ) are labelled: "1-2", "3", "4-5", "6-7".

## Usage

lbl_discrete(
symbol = em_dash(),
unit = 1,
fmt = NULL,
single $=$ NULL,
first = NULL,
last $=$ NULL
)

## Arguments

symbol String: symbol to use for the dash.
unit Minimum difference between distinct values of data. For integers, 1.
fmt String, list or function. A format for break endpoints.
single Glue string: label for singleton intervals. See lbl_glue() for details.
first Glue string: override label for the first category. Write e.g. first $="<\{r\}$ " to create a label like " $<18$ ". See lbl_glue() for details.
last String: override label for the last category. Write e.g. last = " $>\{1\}$ " to create a label like ">65". See lbl_glue() for details.

## Details

No check is done that the data are discrete-valued. If they are not, then these labels may be misleading. Here, discrete-valued means that if $x<y$, then $x<=y$ - unit.
Be aware that Date objects may have non-integer values. See Date.

## Value

A function that creates a vector of labels.

## Formatting endpoints

If fmt is not NULL then it is used to format the endpoints.

- If fmt is a string, then numeric endpoints will be formatted by sprintf(fmt, breaks); other endpoints, e.g. Date objects, will be formatted by format (breaks, fmt).
- If fmt is a list, then it will be used as arguments to format.
- If fmt is a function, it should take a vector of numbers (or other objects that can be used as breaks) and return a character vector. It may be helpful to use functions from the \{scales\} package, e.g. scales: :label_comma().


## See Also

Other labelling functions: lbl_dash(), lbl_endpoints(), lbl_glue(), lbl_intervals(), lbl_manual(), lbl_midpoints(), lbl_seq()

## Examples

```
tab(1:7, c(1, 3, 5), lbl_discrete())
tab(1:7, c(3, 5), lbl_discrete(first = "<= {r}"))
tab(1:7 * 1000, c(1, 3, 5) * 1000, lbl_discrete(unit = 1000))
# Misleading labels for non-integer data
chop(2.5, c(1, 3, 5), lbl_discrete())
```


## Description

This is useful when the left endpoint unambiguously indicates the interval. In other cases it may give errors due to duplicate labels.

## Usage

lbl_endpoints(
left = TRUE,
fmt = NULL,
single = NULL,
first = NULL,
last = NULL,
raw = FALSE
)
lbl_endpoint(fmt $=$ NULL, raw $=$ FALSE, left $=$ TRUE)

## Arguments

left Flag. Use left endpoint or right endpoint?
fmt String, list or function. A format for break endpoints.
single Glue string: label for singleton intervals. See lbl_glue() for details.
first Glue string: override label for the first category. Write e.g. first $=$ " $<\{r\}$ " to create a label like "<18". See lbl_glue() for details.
last String: override label for the last category. Write e.g. last $=$ " $>\{1\}$ " to create a label like ">65". See lbl_glue() for details.
raw [Deprecated]. Use the raw argument to chop() instead.

## Details

lbl_endpoint () is [Defunct] and gives an error since santoku 1.0.0.

## Value

A function that creates a vector of labels.

## Formatting endpoints

If fmt is not NULL then it is used to format the endpoints.

- If fmt is a string, then numeric endpoints will be formatted by sprintf(fmt, breaks); other endpoints, e.g. Date objects, will be formatted by format (breaks, fmt).
- If fmt is a list, then it will be used as arguments to format.
- If fmt is a function, it should take a vector of numbers (or other objects that can be used as breaks) and return a character vector. It may be helpful to use functions from the \{scales\} package, e.g. scales::label_comma().


## See Also

Other labelling functions: lbl_dash(), lbl_discrete(), lbl_glue(), lbl_intervals(), lbl_manual(), lbl_midpoints(), lbl_seq()

## Examples

```
    chop(1:10, c(2, 5, 8), lbl_endpoints(left = TRUE))
    chop(1:10, c(2, 5, 8), lbl_endpoints(left = FALSE))
    if (requireNamespace("lubridate")) {
        tab_width(
                as.Date("2000-01-01") + 0:365,
                months(1),
                labels = lbl_endpoints(fmt = "%b")
            )
    }
    ## Not run:
    # This gives breaks `[1, 2) [2, 3) {3}` which lead to
    # duplicate labels `"2", "3", "3"`:
    chop(1:3, 1:3, lbl_endpoints(left = FALSE))
    ## End(Not run)
```

lbl_glue Label chopped intervals using the glue package

## Description

Use " $\{1\}$ " and " $\{r\}$ " to show the left and right endpoints of the intervals.

## Usage

```
lbl_glue(
    label,
    fmt = NULL,
    single = NULL,
    first = NULL,
    last = NULL,
    raw = FALSE,
)
```


## Arguments

| label | A glue string passed to glue: :glue(). |
| :--- | :--- |
| fmt | String, list or function. A format for break endpoints. |
| single | Glue string: label for singleton intervals. See lbl_glue() for details. |
| first | Glue string: override label for the first category. Write e.g. first $="<\{r\} "$ to <br> create a label like " $<18 "$. See lbl_glue() for details. |
| last | String: override label for the last category. Write e.g. last $=">\{l\} "$ to create a <br> label like " $>65 " . ~ S e e ~ l b l \_g l u e() ~ f o r ~ d e t a i l s . ~$ |
| raw | [Deprecated]. Use the raw argument to chop() instead. |
| $\ldots$ | Further arguments passed to glue: :glue(). |

## Details

The following variables are available in the glue string:

- 1 is a character vector of left endpoints of intervals.
- $r$ is a character vector of right endpoints of intervals.
- l_closed is a logical vector. Elements are TRUE when the left endpoint is closed.
- $r_{\text {_ }} c l o s e d$ is a logical vector, TRUE when the right endpoint is closed.

Endpoints will be formatted by fmt before being passed to glue().

## Value

A function that creates a vector of labels.

## Formatting endpoints

If fmt is not NULL then it is used to format the endpoints.

- If fmt is a string, then numeric endpoints will be formatted by sprintf(fmt, breaks); other endpoints, e.g. Date objects, will be formatted by format (breaks, fmt).
- If fmt is a list, then it will be used as arguments to format.
- If fmt is a function, it should take a vector of numbers (or other objects that can be used as breaks) and return a character vector. It may be helpful to use functions from the \{scales\} package, e.g. scales::label_comma().


## See Also

Other labelling functions: lbl_dash(), lbl_discrete(), lbl_endpoints(), lbl_intervals(), lbl_manual(), lbl_midpoints(), lbl_seq()

## Examples

```
tab(1:10, c(1, 3, 3, 7),
    labels = lbl_glue("{l} to {r}", single = "Exactly {l}"))
tab(1:10 * 1000, c(1, 3, 5, 7) * 1000,
    labels = lbl_glue("{l}-{r}",
            fmt = function(x) prettyNum(x, big.mark=',')))
# reproducing lbl_intervals():
interval_left <- "{ifelse(l_closed, '[', '(')}"
interval_right <- "{ifelse(r_closed, ']', ')')}"
glue_string <- paste0(interval_left, "{l}", ", ", "{r}", interval_right)
tab(1:10, c(1, 3, 3, 7), labels = lbl_glue(glue_string, single = "{{{l}}}"))
```


## Description

These labels are the most exact, since they show you whether intervals are "closed" or "open", i.e. whether they include their endpoints.

```
Usage
    lbl_intervals(
        fmt = NULL,
        single = "{{{l}}}",
        first = NULL,
        last = NULL,
        raw = FALSE
    )
```


## Arguments

fmt String, list or function. A format for break endpoints.
single Glue string: label for singleton intervals. See lbl_glue() for details.
first Glue string: override label for the first category. Write e.g. first $="<\{r\}$ " to create a label like "<18". See lbl_glue() for details.
last String: override label for the last category. Write e.g. last $=$ " $>\{1\}$ " to create a label like ">65". See lbl_glue() for details.
raw [Deprecated]. Use the raw argument to chop() instead.

## Details

Mathematical set notation looks like this:

- [a, b]: all numbers x where $\mathrm{a}<=\mathrm{x}<=\mathrm{b}$;
- (a, b): all numbers where $\mathrm{a}<\mathrm{x}<\mathrm{b}$;
- [a, b): all numbers where $a<=x<b ;$
- ( $\mathrm{a}, \mathrm{b}$ ]: all numbers where $\mathrm{a}<\mathrm{x}<=\mathrm{b}$;
- \{a\}: just the number a exactly.


## Value

A function that creates a vector of labels.

## Formatting endpoints

If fmt is not NULL then it is used to format the endpoints.

- If fmt is a string, then numeric endpoints will be formatted by sprintf(fmt, breaks); other endpoints, e.g. Date objects, will be formatted by format (breaks, fmt).
- If fmt is a list, then it will be used as arguments to format.
- If fmt is a function, it should take a vector of numbers (or other objects that can be used as breaks) and return a character vector. It may be helpful to use functions from the \{scales\} package, e.g. scales: :label_comma().


## See Also

Other labelling functions: lbl_dash(), lbl_discrete(), lbl_endpoints(), lbl_glue(), lbl_manual(), lbl_midpoints(), lbl_seq()

## Examples

```
tab(-10:10, c(-3, 0, 0, 3),
    labels = lbl_intervals())
tab(-10:10, c(-3, 0, 0, 3),
    labels = lbl_intervals(fmt = list(nsmall = 1)))
tab_evenly(runif(20), 10,
    labels = lbl_intervals(fmt = percent))
```

    lbl_midpoints
    Label chopped intervals by their midpoints
    
## Description

This uses the midpoint of each interval for its label.

## Usage

```
lbl_midpoints(
    fmt = NULL,
    single = NULL,
    first = NULL,
    last = NULL,
    raw = FALSE
)
```


## Arguments

| fmt | String, list or function. A format for break endpoints. |
| :--- | :--- |
| single | Glue string: label for singleton intervals. See lbl_glue() for details. <br> Girst <br> create a label like " $<18 "$. See $1 b l \_g l u e() ~ f o r ~ d e t a i l s . ~$ |
| last | String: override label for the last category. Write e.g. last $=">\{1\} "$ to create a <br> label like " $>65 " . ~ S e e ~ l b l \_g l u e() ~ f o r ~ d e t a i l s . ~$ |
| [Deprecated]. Use the raw argument to chop() instead. |  |

## Value

A function that creates a vector of labels.

## Formatting endpoints

If fmt is not NULL then it is used to format the endpoints.

- If fmt is a string, then numeric endpoints will be formatted by sprintf(fmt, breaks); other endpoints, e.g. Date objects, will be formatted by format (breaks, fmt).
- If fmt is a list, then it will be used as arguments to format.
- If fmt is a function, it should take a vector of numbers (or other objects that can be used as breaks) and return a character vector. It may be helpful to use functions from the \{scales\} package, e.g. scales::label_comma().


## See Also

Other labelling functions: $1 b 1 \_d a s h(), 1 b l \_d i s c r e t e(), 1 b l \_e n d p o i n t s(), 1 b l \_g l u e(), l b l \_i n t e r v a l s(), ~$ lbl_manual(), lbl_seq()

## Examples

chop(1:10, c(2, 5, 8), lbl_midpoints())
lbl_seq Label chopped intervals in sequence

## Description

lbl_seq() labels intervals sequentially, using numbers or letters.

## Usage

lbl_seq(start = "a")

## Arguments

start String. A template for the sequence. See below.

## Details

start shows the first element of the sequence. It must contain exactly one character out of the set "a", "A", "i", "I" or "1". For later elements:

- "a" will be replaced by "a", "b", "c", ...
- "A" will be replaced by "A", "B", "C", ...
- "i" will be replaced by lower-case Roman numerals "i", "ii", "iii", ...
- "I" will be replaced by upper-case Roman numerals "I", "II", "III", ...
- "1" will be replaced by numbers "1", "2", "3", ...

Other characters will be retained as-is.

## Value

A function that creates a vector of labels.

## See Also

Other labelling functions: $1 b 1 \_d a s h(), 1 b l \_d i s c r e t e(), ~ l b l \_e n d p o i n t s(), ~ l b l \_g l u e(), ~ l b l \_i n t e r v a l s(), ~$ lbl_manual(), lbl_midpoints()

## Examples

```
chop(1:10, c(2, 5, 8), lbl_seq())
chop(1:10, c(2, 5, 8), lbl_seq("i."))
chop(1:10, c(2, 5, 8), lbl_seq("(A)"))
```

non-standard-types Tips for chopping non-standard types

## Description

Santoku can handle many non-standard types.

## Details

- If objects can be compared using $<,==$ etc. then they should be choppable.
- Objects which can't be converted to numeric are handled within R code, which may be slower.
- Character $x$ and breaks are chopped with a warning.
- If $x$ and breaks are not the same type, they should be able to be cast to the same type, usually using vctrs: :vec_cast_common().
- Not all chopping operations make sense, for example, chop_mean_sd() on a character vector.
- For indexed objects such as stats: :ts() objects, indices will be dropped from the result.
- If you get errors, try setting extend = FALSE (but also file a bug report).
- To request support for a type, open an issue on Github.


## See Also

brk-width-for-Datetime
percent Simple percentage formatter

## Description

percent () formats $x$ as a percentage. For a wider range of formatters, consider the scales package.

## Usage

percent ( x )

## Arguments

X
Numeric values.

## Value

$x$ formatted as a percent.

## Examples

percent(0.5)

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