Package 'santoku'

June 9, 2020

```
Type Package
Title A Versatile Cutting Tool
Version 0.4.0
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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 and times.
License MIT + file LICENSE
Encoding UTF-8
LazyData true
Roxygen list(markdown = TRUE)
RoxygenNote 7.1.0
Suggests covr,
     testthat (>= 2.1.0),
     knitr,
     lubridate,
     microbenchmark,
     purrr,
     rmarkdown,
     scales,
     withr
LinkingTo Rcpp
Imports Rcpp,
     assertthat,
     lifecycle,
     vctrs
URL https://github.com/hughjonesd/santoku, https://hughjonesd.github.io/santoku/
BugReports https://github.com/hughjonesd/santoku/issues
VignetteBuilder knitr
RdMacros lifecycle
```

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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
- 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 or date-times.

breaks-class 3

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>

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

```
## $3 method for class 'breaks'
format(x, ...)
## $3 method for class 'breaks'
print(x, ...)
is.breaks(x, ...)
```

Arguments

```
x A breaks object
```

... Unused

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brk-left-right

Left- or right-closed breaks

Description

Questioning

Usage

```
brk_left(breaks)
brk_right(breaks)
```

Arguments

breaks

A numeric vector.

Details

These functions are in the "questioning" stage because they clash with the left argument to chop() and friends.

These functions override the left argument of chop().

Value

A (function which returns an) object of class breaks.

Examples

```
chop(5:7, brk_left(5:7))
chop(5:7, brk_right(5:7))
chop(5:7, brk_left(5:7))
```

brk_default

Create a standard set of breaks

Description

Create a standard set of breaks

Usage

```
brk_default(breaks)
```

Arguments

breaks

A numeric vector.

brk_manual 5

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.

brk_width-for-datetime

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)</pre>
```

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.

A scalar of class Date or POSIXct. Can be omitted.

Usage

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

Arguments

start

width A scalar difftime, Period or Duration object.

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.

chop 7

chop Cut numeric 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.

Usage

```
chop(
  х,
  breaks,
  labels,
  extend = NULL,
  left = TRUE,
  close_end = FALSE,
  drop = TRUE
)
kiru(
  Х,
  breaks,
  labels,
  extend = NULL,
  left = TRUE,
  close_end = FALSE,
  drop = TRUE
```

Arguments

X	A numeric vector.
breaks	See below.
labels	See below.
extend	Logical. Extend breaks to +/-Inf?
left	Logical. Left-closed breaks?
close_end	Logical. Close last break at right? (If left is FALSE, close first break at left?)

drop Logical. Drop unused levels from the result?

Details

breaks may be a numeric 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 $\{3,5\}$.

By default, left-closed intervals are created. If left is FALSE, right- closed intervals are created.

If close_end is TRUE the end break will be closed at both ends, ensuring that all values y with $min(x) \le y \le max(x)$ are included in the default intervals. That is:

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- If left is TRUE and close_end is TRUE, breaks will look like [x1, x2), [x2, x3) ... [x_n-1, x_n].
- If left is FALSE and close_end is TRUE, breaks will look like [x1, x2], (x2, x3] ... (x_n-1, x_n].
- If left is TRUE and close_end is FALSE, all breaks will look like ...[x1, x2) ...
- If left is FALSE and close_end is FALSE, all breaks will look like ...(x1, x2] ...

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 in this context, to create a variety of data-dependent breaks.

labels may be a character vector. It should have the same length as the number of intervals. Alternatively, use a lbl_function such as [lbl_seq()].

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 min(x) < min(breaks) and max(x) > max(breaks) respectively.

Extending intervals, either by extend = NULL or extend = FALSE, *always* leaves the central, non-extended intervals unchanged. In particular, close_end applies to the central intervals, not to the extended ones. For example, if breaks = c(1,3,5) and close_end = TRUE, the resulting breaks will be

```
[1, 3), [3, 5]
```

and if they are extended on both ends the result will be e.g.

```
[-Inf, 1), [1, 3), [3, 5], (5, Inf]
```

NA values in x, and values which are outside the (extended) endpoints, return NA.

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.

```
[x1, x2) ...
```

• If left is FALSE and close_end is FALSE, all breaks will look like '...(x1, x2]: R:x1,%20x2)%20...%60%0A*%20If [lbl_seq()]: R:lbl_seq() [3, 5]: R:3,%205 [3, 5]: R:3,%205

kiru is a synonym for chop. If you load tidyr, you can use it to avoid confusion with tidyr::chop().

Value

A factor of the same length as x, representing the intervals containing the value of x.

See Also

cut

Other chopping functions: chop_mean_sd(), chop_n(), chop_quantiles(), chop_width(), fillet()

```
chop(1:3, 2)
chop(1:10, c(2, 5, 8))
chop(1:10, c(2, 5, 8), extend = FALSE)
chop(1:10, c(2, 5, 5, 8))
chop(1:10, c(2, 5, 8), left = FALSE)
```

chop_mean_sd 9

```
chop(1:10, c(2, 5, 8), close_end = TRUE)
chop(1:10, brk_quantiles(c(0.25, 0.75)))
chop(1:10, c(2, 5, 8), labels = lbl_dash())
# floating point inaccuracy:
chop(0.3/3, c(0, 0.1, 0.1, 1))
```

chop_mean_sd

Chop by standard deviations

Description

Intervals of width 1 standard deviation are included on either side of the mean. The outermost pair of intervals will be shorter if sd is not a whole number.

Usage

```
chop_mean_sd(x, sd = 3, ...)
brk_mean_sd(sd = 3)
```

Arguments

x A numeric vector.

sd Positive number: include up to sd standard deviations.

... Passed to chop().

Value

For chop_* functions, a factor of the same length as x.

See Also

```
Other chopping functions: chop_n(), chop_quantiles(), chop_width(), chop(), fillet()
```

```
chop_mean_sd(1:10)
chop(1:10, brk_mean_sd())
```

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chop_n

Chop into fixed-sized groups

Description

chop_n() creates intervals containing a fixed number of elements. One interval may have fewer elements.

Usage

```
chop_n(x, n, ..., close_end = TRUE)
brk_n(n)
```

Arguments

x A numeric vector.
 n Integer: number of elements in each interval.
 ... Passed to chop().
 close_end Passed to chop().

Details

Note that $chop_n()$ sets $close_end = TRUE$ by default.

Groups may be larger than n, if there are too many duplicated elements in x. If so, a warning is given.

Value

For chop $_{-}^{*}$ functions, a factor of the same length as x.

See Also

```
Other chopping functions: chop_mean_sd(), chop_quantiles(), chop_width(), chop(), fillet()
```

```
table(chop_n(1:10, 5))
table(chop_n(1:10, 4))
# too many duplicates
x <- rep(1:2, each = 3)
chop_n(x, 2)</pre>
```

chop_quantiles 11

iles	

Description

chop_quantiles chops data by quantiles. chop_equally chops data into equal-sized groups. chop_deciles is a convenience shortcut and chops into deciles.

Usage

```
chop_quantiles(x, probs, ..., left = is.numeric(x), close_end = TRUE)
chop_deciles(x, ...)
chop_equally(x, groups, ..., left = is.numeric(x), close_end = TRUE)
brk_quantiles(probs, ...)
brk_equally(groups)
```

Arguments

```
x A numeric vector.

probs A vector of probabilities for the quantiles.

... Passed to chop(), or for brk_quantiles to stats::quantile().

left Passed to chop().

close_end Passed to chop().

groups Number of groups.
```

Details

Note that these functions set close_end = TRUE by default. This helps ensure that e.g. $chop_quantiles(x, c(0, 1/3, 2/3, 1))$ will split the data into three equal-sized groups.

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().

Value

For chop_* functions, a factor of the same length as x.

See Also

```
Other chopping functions: chop_mean_sd(), chop_n(), chop_width(), chop(), fillet()
```

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Examples

```
chop_quantiles(1:10, 1:3/4)
chop(1:10, brk_quantiles(1:3/4))
chop_deciles(1:10)
chop_equally(1:10, 5)
# to label by the quantiles themselves:
chop_quantiles(1:10, 1:3/4, lbl_intervals(raw = TRUE))
```

chop_width

Chop into equal-width intervals

Description

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

Usage

```
chop_width(x, width, start, ...)
chop_evenly(x, intervals, ..., groups, close_end = TRUE)
brk_width(width, start)
## Default S3 method:
brk_width(width, start)
brk_evenly(intervals)
```

Arguments

x A numeric vector. width Width of intervals.

start Leftpoint of first interval. By default the lowest finite x.

... Passed to chop().

intervals Integer: number of intervals to create.

groups Do not use. **Deprecated**

close_end Passed to chop().

Details

Note that chop_evenly sets close_end = TRUE by default.

Value

For chop_* functions, a factor of the same length as x.

exactly 13

See Also

```
brk_width-for-datetime
```

```
Other chopping functions: chop_mean_sd(), chop_n(), chop_quantiles(), chop(), fillet()
```

Examples

```
chop_width(1:10, 2)
chop_width(1:10, 2, start = 0)
chop(1:10, brk_width(2, 0))
chop_evenly(0:10, 5)
```

exactly

Syntactic sugar

Description

exactly lets you write chop(x,c(1,exactly(2),3)). This is the same as chop(x,c(1,2,2,3)) but conveys your intent more clearly.

Usage

```
exactly(x)
```

Arguments

Χ

A numeric vector.

Value

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

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

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fillet

Chop data precisely (for programmers)

Description

Chop data precisely (for programmers)

Usage

```
fillet(x, breaks, labels, left = TRUE, close_end = FALSE)
```

Arguments

```
x A numeric vector.
breaks Passed to chop().
labels Passed to chop().
left Passed to chop().
close_end Passed to chop().
```

Details

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().

Value

For chop_* functions, a factor of the same length as x.

See Also

```
Other chopping functions: chop_mean_sd(), chop_n(), chop_quantiles(), chop_width(), chop()
```

Examples

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

knife

Deprecated

Description

Soft-deprecated knife() is deprecated in favour of purrr::partial().

Usage

```
knife(...)
```

Arguments

... Parameters for chop().

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Value

A function.

lbl_dash

Label chopped intervals like 1 - 3, 4 - 5, ...

Description

This label style is user-friendly, but doesn't distinguish between left- and right-closed intervals.

Usage

```
lbl_dash(symbol = " - ", raw = FALSE, fmt = NULL)
```

Arguments

symbol String: symbol to use for the dash.

raw Logical. Always use raw breaks in labels, rather than e.g. quantiles or standard deviations?

fmt A format. Can be a string, passed into base::sprintf() or format() methods;

Value

A vector of labels for chop, or a function that creates labels.

See Also

```
Other labelling functions: lbl_discrete(), lbl_format(), lbl_intervals(), lbl_manual(), lbl_seq()
```

Examples

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

or a one-argument formatting function.

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lbl_discrete

Label discrete data

Description

Experimental

Usage

```
lbl_discrete(symbol = " - ", fmt = NULL)
```

Arguments

symbol String: symbol to use for the dash.

fmt A format. Can be a string, passed into base::sprintf() or format() methods;

or a one-argument formatting function.

Details

```
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".
```

No check is done that the data is discrete-valued. If it isn't, then these labels may be misleading. Here, discrete-valued means that if x < y, then x <= y -1.

Be aware that Date objects may have non-integer values. See Date.

Value

A vector of labels for chop, or a function that creates labels.

See Also

```
Other labelling functions: lbl_dash(), lbl_format(), lbl_intervals(), lbl_manual(), lbl_seq()
```

```
tab(1:7, c(1, 3, 5), lbl_discrete())
# Misleading labels for non-integer data
chop(2.5, c(1, 3, 5), lbl_discrete())
```

lbl_endpoint 17

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Label chopped intervals by their left or right endpoints

Description

This is useful when the left endpoint unambiguously indicates the interval.

Usage

```
lbl_endpoint(fmt = NULL, raw = FALSE, left = TRUE)
```

Arguments

fmt	A format. Can be a string, passed into base::sprintf() or format() methods; or a one-argument formatting function.
raw	Logical. Always use raw breaks in labels, rather than e.g. quantiles or standard deviations?
left	Flag. Use left endpoint or right endpoint?

Value

A vector of labels for chop, or a function that creates labels.

Examples

lbl_format

Label chopped intervals with arbitrary formatting

Description

Questioning

Usage

```
lbl_format(fmt, fmt1 = "%.3g", raw = FALSE)
```

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Arguments

fmt	A format. Can be a string, passed into base::sprintf() or format() methods; or a one-argument formatting function.
fmt1	Format for breaks consisting of a single value.
raw	Logical. Always use raw breaks in labels, rather than e.g. quantiles or standard deviations?

Details

These labels let you format breaks arbitrarily, using either a string (passed to sprintf()) or a function.

If fmt is a function, it must accept two arguments, representing the left and right endpoints of each interval.

If breaks are non-numeric, you can only use "%s" in a string fmt. breaks will be converted to character in this case.

lbl_format() is in the "questioning" stage. As an alternative, consider using lbl_dash() or lbl_intervals() with the fmt argument.

Value

A vector of labels for chop, or a function that creates labels.

See Also

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

Examples

lbl_intervals

Label chopped intervals using set notation

Description

Label chopped intervals using set notation

Usage

```
lbl_intervals(raw = FALSE, fmt = NULL)
```

Ibl_manual

Arguments

raw Logical. Always use raw breaks in labels, rather than e.g. quantiles or standard

deviations?

fmt A format. Can be a string, passed into base::sprintf() or format() methods;

or a one-argument formatting function.

Details

Mathematical set notation is as follows:

```
• [a,b]: all numbers x where a \le x \le b;
```

• (a,b): all numbers where a < x < b;

• [a,b): all numbers where $a \le x < b$;

• (a,b]: all numbers where $a < x \le b$;

• {a}: just the number a.

Value

A vector of labels for chop, or a function that creates labels.

See Also

```
Other labelling functions: lbl_dash(), lbl_discrete(), lbl_format(), lbl_manual(), lbl_seq()
```

Examples

lbl_manual

Label chopped intervals in a user-defined sequence

Description

lbl_manual() uses an arbitrary sequence to label intervals. If the sequence is too short, it will be pasted with itself and repeated.

Usage

```
lbl_manual(sequence, fmt = "%s")
```

Arguments

sequence A character vector of labels.

fmt A format. Can be a string, passed into base::sprintf() or format() methods;

or a one-argument formatting function.

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Value

A vector of labels for chop, or a function that creates labels.

See Also

```
Other labelling functions: lbl_dash(), lbl_discrete(), lbl_format(), lbl_intervals(), lbl_seq()
```

Examples

```
chop(1:10, c(2, 5, 8), lbl_manual(c("w", "x", "y", "z")))
# if labels need repeating:
chop(1:10, 1:10, lbl_manual(c("x", "y", "z")))
```

lbl_seq

Label chopped intervals in sequence

Description

1b1_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.

See Also

```
Other labelling functions: lbl_dash(), lbl_discrete(), lbl_format(), lbl_intervals(), lbl_manual()
```

```
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)"))
```

percent 21

percent

Simple formatter

Description

For a wider range of formatters, consider the "scales" package.

Usage

```
percent(x)
```

Arguments

Χ

Numeric values.

Value

x formatted as a percent.

Examples

```
percent(0.5)
```

tab

Tabulate data by intervals

Description

These functions call their related chop_xxx function, and call table() on the result.

Usage

```
tab(...)
tab_width(...)
tab_evenly(...)
tab_n(...)
tab_mean_sd(...)
```

Arguments

... Passed to chop

Value

```
A table().
```

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```
tab(1:10, c(2, 5, 8))
tab_mean_sd(1:10)
```

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