Package: sift (via r-universe)

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break_join

Join tables based on overlapping intervals.

Description

User-friendly interface that synthesizes power of dplyr::left_join and findInterval.

Usage

```
break_join(x, y, brk = character(), by = NULL, ...)
```

Arguments

x A data frame.

y Data frame containing desired reference information.

brk Name of column in x and y to join by via interval overlapping. Must be coercible to numeric.

by Joining variables, if needed. See mutate-joins.

additional arguments automatically directed to findInterval and dplyr::left_join. No partial matching.

Value

An object of the same type as x.

- All x rows will be returned.
- All columns between x and y are returned.
- Rows in y are matched with x based on overlapping values of brk (e.g. findInterval(x\$brk, y\$brk, ...)).

Examples

```
# joining USA + UK leaders with population time-series
break_join(us_uk_pop, us_uk_leaders, brk = c("date" = "start"))

# simple dataset
set.seed(1)
a <- data.frame(p = c(rep("A", 10), rep("B", 10)), q = runif(20, 0, 10))
b <- data.frame(p = c("A", "A", "B", "B"), q = c(3, 5, 6, 9), r = c("a1", "a2", "b1", "b2"))
break_join(a, b, brk = "q") # p identified as common variable automatically</pre>
```

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```
break_join(a, b, brk = "q", by = "p") # same result
break_join(a, b, brk = "q", all.inside = TRUE) # note missing values have been filled

# joining toll prices with vehicle time-series

library(mopac)
library(dplyr, warn.conflicts = FALSE)
library(hms)

express %>%
  mutate(time_hms = as_hms(time)) %>%
  break_join(rates, brk = c("time_hms" = "time"))
```

comms

Simulated records of radio station communications.

Description

Dataset intended to demonstrate usage of sift::conjecture.

Usage

comms

Format

An object of class tbl_df (inherits from tbl, data.frame) with 50000 rows and 4 columns.

conjecture

Specialized "long to wide" reshaping

Description

On the surface, conjecture() appears similar to tidyr::pivot_wider(), but uses different logic tailored to a specific type of dataset:

- column corresponding to names_from contains only 2 levels
- there is no determinate combination of elements to fill 2 columns per row.

See vignette("conjecture") for more details.

Usage

```
conjecture(data, sort_by, names_from, names_first)
```

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Arguments

data A data frame to reshape.

Sort_by Column name, as symbol. Plays a similar role as values_from in pivot_wider(), but also serves as sorting dimension for underlying conjecture algorithm.

Column name, as symbol. Used to differentiate anterior/posterior observations. Column must only contain 2 levels (missing values not allowed).

names_first level in variable specified by names_from indicating anterior observation.

Details

conjecture() uses the following routine to match elements:

- 1. Values in sort_by are separated into two vectors: anterior and posterior.
- 2. Each anterior element is matched with the closest posterior element measured by sort_by.

Value

An object of the same type as data.

Examples

```
# See vignette("conjecture") for more examples
conjecture(comms, timestamp, type, "send")
```

kluster

Automatically cluster 1-dimensional continuous data.

Description

Automatically cluster 1-dimensional continuous data.

Usage

```
kluster(x, bw = "SJ", fixed = FALSE)
```

Arguments

X	Vector to be clustered. Must contain at least 1 non-missing value.
bw	kernel bandwidth. Default "SJ" should suffice more application, however you can supply a custom numeric value. See ?stats::density for more information.
fixed	logical; if TRUE, performs simple 1-dimensional clustering without kernel density estimation. default FALSE.

Value

An integer vector identifying the cluster corresponding to each element in x.

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Examples

```
# Below vector clearly has 2 groups.
# kluster will identify these groups using kernel density estimation.
kluster(c(0.1, 0.2, 1))
# kluster shines in cases where manually assigning groups via "eyeballing" is impractical.
# Suppose we obtained vector 'x' without knowing how it was generated.
set.seed(1)
nodes \leftarrow runif(10, min = 0, max = 100)
x \leftarrow lapply(nodes, function(x) rnorm(10, mean = x, sd = 0.1))
kluster(x) # kluster reveals the natural grouping
kluster(x, bw = 10) # adjust bandwidth depending on application
# Example with faithful dataset
faithful$k <- kluster(faithful$eruptions)</pre>
library(ggplot2)
ggplot(faithful, aes(eruptions)) +
  geom_density() +
  geom_rug(aes(color = factor(k))) +
  theme_minimal() +
  scale_color_discrete(name = "k")
```

nyt2020

2020 New York Times Headlines

Description

Includes selected headlines and additional metadata for NYT articles throughout 2020. This dataset is not a comprehensive account of all major events from 2020.

Usage

nyt2020

Format

A data frame with 1,830 rows and 6 variables:

```
headline Article Headline
abstract Brief summary of article
byline Contributing Writers
pub_date Date of Publication
section_name NYT section in which article was published
web_url Article URL ...
```

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Source

Obtained using NYT Developer Portal (Archive API)

sift

Augmented data frame filtering.

Description

Imagine dplyr::filter that includes neighboring observations. Choose how many observations to include by adjusting inputs sift.col and scope.

Usage

```
sift(.data, sift.col, scope, ...)
```

Arguments

.data	A data frame.
sift.col	Column name, as symbol, to serve as "sifting/augmenting" dimension. Must be non-missing and coercible to numeric.
scope	Specifies augmentation bandwidth relative to "key" observations. Parameter should share the same scale as sift.col.
	If length 1, bandwidth used is +/- scope.
	If length 2, bandwidth used is (-scope[1], +scope[2]).
• • •	Expressions passed to dplyr::filter, of which the results serve as the "key" observations. The same data-masking rules used in dplyr::filter apply here.

Details

sift() can be understood as a 2-step process:

- 1. .data is passed to dplyr::filter, using subsetting expression(s) provided in We'll refer to these intermediate results as "key" observations.
- 2. For each key observation, sift expands the row selection bidirectionally along dimension specified by sift.col. Any row from the original dataset within scope units of a key observation is captured in the final result.

Essentially, this allows us to "peek" at neighboring rows surrounding the key observations.

Value

A sifted data frame, with 2 additional columns:

- .cluster <int>: Identifies resulting group formed by each key observation and its neighboring rows. When the key observations are close enough together, the clusters will overlap.
- .key <lgl>: TRUE indicates key observation.

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Examples

```
# See current events from same timeframe as 2020 Utah Monolith discovery.
sift(nyt2020, pub_date, scope = 2, grepl("Monolith", headline))

# or Biden's presidential victory.
sift(nyt2020, pub_date, scope = 2, grepl("Biden is elected", headline))

# We can specify lower & upper scope to see what happened AFTER Trump tested positive.
sift(nyt2020, pub_date, scope = c(0, 2), grepl("Trump Tests Positive", headline))

# sift recognizes dplyr group specification.
library(dplyr)
library(mopac)
express %>%
  group_by(direction) %>%
  sift(time, 30, plate == "EAS-1671") # row augmentation performed within groups.
```

us_uk_pop

Fragments of US & UK population & leaders

Description

These datasets are intended to demonstrate usage of sift::break_join.

Usage

```
us_uk_pop
us_uk_leaders
```

Source

See tidyr::who and ggplot2::presidential.

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