

# Package ‘cubble’

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**Title** A Vector Spatio-Temporal Data Structure for Data Analysis

**Version** 0.2.0

**Description** A spatiotemporal data object in a relational data structure to separate the recording of time variant/ invariant variables.

**License** MIT + file LICENSE

**Language** au

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**add\_missing\_prct**      *Compute missing summary*

### Description

Compute missing summary

### Usage

```
add_missing_prct(data, vars)
```

### Arguments

data	a cubble object
vars	variables to compute percentage missing (support tidyselect)

### Details

- `add_missing_prct()` computes the percentage of missing for the selected variables

**Value**

a cubble object with additional columns VAR\_missing

**Examples**

```
climate_aus %>% add_missing_prct(vars = prcp)
```

---

as\_cubble

*The constructor for the cubble class*

---

**Description**

The constructor for the cubble class

**Usage**

```
as_cubble(data, key, index, coords, ...)

## S3 method for class 'list'
as_cubble(data, key, index, coords, by = NULL, output = "auto-match", ...)

## S3 method for class 'tbl_df'
as_cubble(data, key, index, coords, ...)

## S3 method for class 'rowwise_df'
as_cubble(data, key, index, coords, ...)

## S3 method for class 'cubble_df'
print(
  x,
  width = NULL,
  ...,
  n_extra = NULL,
  n = NULL,
  max_extra_cols = NULL,
  max_footer_lines = NULL
)

## S3 method for class 'cubble_df'
tbl_sum(data)

is_cubble(data)

cubble(..., key, index, coords)
```

## Arguments

data	the object to be created or tested as cubble
key	the spatial identifier
index	the time identifier
coords	the coordinates that characterise the spatial dimension
...	a list object to create new cubble
by	only used in <code>as_cubble.list()</code> to specify the linking key between spatial and temporal data
output	either "all" or "unmatch", whether to output all or a list of unmatched summary
x, width, n_extra, n, max_extra_cols, max_footer_lines	see pillar <code>tbl-format.R</code>

## Value

a cubble object
a cubble object
a cubble object
a TRUE/FALSE predicate
a cubble object

## Examples

```
# Disclaimer: to make the examples easier, here we first `climate_flat` into
# different classes and show how they can be casted into a cubble. This is to
# demonstrate if your data come in one of the classes, it can be directly cast
# into a cubble. By no mean you need to first transform your data into any of
# the following class and then cast it to cubble.

# If the data is in a tibble:
climate_flat %>% as_cubble(key = id, index = date, coords = c(long, lat))

# If the spatial and temporal information are in two separate tables:
library(dplyr)
spatial <- climate_flat %>% select(id:wmo_id) %>% distinct()
temporal <- climate_flat %>% select(id, date: tmin) %>% filter(id != "ASN00009021")
as_cubble(data = list(spatial = spatial, temporal = temporal),
          key = id, index = date, coords = c(long, lat))

# If the data is already in a rowwise_df:
dt <- climate_flat %>%
  tidyr::nest(ts = date:tmin) %>%
  dplyr::rowwise()
dt %>% as_cubble(key = id, index = date, coords = c(long, lat))

# If the data is already in a tsibble, only need to supply `coords`
dt <- climate_flat %>% tsibble::as_tsibble(key = id, index = date)
dt %>% as_cubble(coords = c(long, lat))
```

```
# If the data is in netcdf:  
path <- system.file("ncdf/era5-pressure.nc", package = "cubicle")  
raw <- ncdf4::nc_open(path)  
dt <- as_cubicle(raw, vars = c("q", "z"))
```

---

**climate\_aus***Australia climate data - 639 stations*

---

**Description**

Daily measure on precipitation (prcp) maximum temperature (tmax), and minimum temperature (tmin) in 2020 for 639 stations. `stations` and `climate` are the separate spatial and temporal objects while `climate_aus` is the combined cubicle object.

**Usage**

```
climate_aus
```

**Format**

An object of class `cubicle_df` (inherits from `rowwise_df`, `tbl_df`, `tbl`, `data.frame`) with 639 rows and 7 columns.

**Details**

**id** station id  
**lat** latitude of the station  
**long** longitude of the station  
**elev** elevation of the station  
**name** station name  
**wmo\_id** the world meteorological organisation (WMO) station number  
**ts** a list-column that nests all the time-wise measures: date, prcp, tmax, and tmin

**See Also**

```
climate_subset climate_flat
```

**Examples**

```
climate_aus %>% face_temporal() %>% face_spatial()
```

---

climate_flat	<i>Australia climate data - 5 stations</i>
--------------	--

---

## Description

Daily measure on precipitation (prcp) maximum temperature (tmax), and minimum temperature (tmin) in 2020 for 5 stations.

## Usage

```
climate_flat  
stations  
climate
```

## Format

A tibble object with 155 rows and 10 columns

**id** station id  
**lat** latitude of the station  
**long** longitude of the station  
**elev** elevation of the station  
**name** station name  
**wmo\_id** the world meteorological organisation (WMO) station number  
**date** the date that prcp, tmax, and tmin recorded  
**prcp** precipitation  
**tmax** maximum temperature  
**tmin** minimum temperature

An object of class `tbl_df` (inherits from `tbl`, `data.frame`) with 5 rows and 6 columns.

An object of class `tbl_df` (inherits from `tbl`, `data.frame`) with 1830 rows and 5 columns.

## See Also

`climate_aus` `climate_subset`

## Examples

```
climate_flat %>% as_cubicle(key = id, index = date, coords = c(long, lat))
```

---

climate_subset	<i>Australia climate data - 30 stations</i>
----------------	---

---

## Description

Daily measure on precipitation (prcp) maximum temperature (tmax), and minimum temperature (tmin) in 2020 for 30 stations.

## Usage

```
climate_subset
```

## Format

A cubble object

**id** station id

**lat** latitude of the station

**long** longitude of the station

**elev** elevation of the station

**name** station name

**wmo\_id** the world meteorological organisation (WMO) station number

**ts** a list-column that nests all the time-wise measures: date, prcp, tmax, and tmin

## See Also

```
climate_aus climate_flat
```

## Examples

```
climate_subset %>% face_temporal()
```

---

extract_var	<i>Functions to extract NetCDF dimension and variables</i>
-------------	--

---

## Description

Functions to extract NetCDF dimension and variables

## Usage

```
extract_var(data, vars)
```

```
extract_longlat(data)
```

```
extract_time(data)
```

**Arguments**

- data** a NetCDF file read in from `ncdf4::nc_open()`  
**vars** variables to read, see the variables in your data with `names(data$var)`

**Value**

extracted netcdf4 components

<code>face_spatial</code>	<i>Switch a cubble object into the nested form</i>
---------------------------	--

**Description**

`face_spatial()` turns a long cubble back into a nest cubble and can be seen as the inverse operation of `face_temporal()`. The nested cubble identifies each row by key and is suitable for operations whose output doesn't involve a time index.

**Usage**

```
face_spatial(data)
```

**Arguments**

- data** a long cubble object

**Value**

a cubble object in the nested form

**Examples**

```
cb_long <- climate_flat %>%
  as_cubble(key = id, index = date, coords = c(long, lat)) %>%
  face_temporal()

cb_long %>% face_spatial()
```

---

face_temporal	<i>Switch a cubble object into the long form</i>
---------------	--

---

## Description

`face_temporal()` switches a cubble object into a long cubble, suitable for temporal operations. The long cubble uses the combination of key and index to identify each row and arranges each key as a separate group.

## Usage

```
face_temporal(data, col)
```

## Arguments

<code>data</code>	a nested cubble object
<code>col</code>	the list column to be expanded, <code>col</code> is required to be specified if there are more than one list column and the list column name is not <code>ts</code>

## Value

a cubble object in the nested form

## Examples

```
climate_flat %>%
  as_cibble(key = id, index = date, coords = c(long, lat)) %>%
  face_temporal()
```

---

fill_gaps.cubble_df	<i>tsibble methods implemented in cubble</i>
---------------------	--

---

## Description

See [fill\\_gaps](#)

## Usage

```
## S3 method for class 'cubble_df'
fill_gaps(.data, ..., .full = FALSE, .start = NULL, .end = NULL)
```

## Arguments

<code>.data, ..., .full, .start, .end</code>	see tsibble documentation
--	---------------------------

**Value**

a cubble object

---

form

*Functions to extract cubble attributes*

---

**Description**

Functions to extract cubble attributes

**Usage**

```
form(data)  
  
is_long(data)  
  
is_nested(data)  
  
spatial(data)  
  
key_vars(data)  
  
key_data(data)  
  
coords(data)  
  
coord_x(data)  
  
coord_y(data)  
  
index(data)
```

**Arguments**

data            an cubble object

**Details**

Apart from inheriting attributes names, row.names, and class from the underlying tibble, a cubble has its site identifier: key, temporal identifier, index, and spatial coordinate reference: coords.

If a cubble object is also a tsibble, then tsibble attributes (key, index, index2, interval) are also preserved and can be accessed via the relevant functions in the tsibble package. (NOT FULLY IMPLEMENTED)

**Value**

the name of cubble attributes

## Examples

```
# extract attributes of a cubble object
form(climate_aus)
spatial(climate_aus) %>% head(5)
key_data(climate_aus) %>% head(5)
key_vars(climate_aus)
index(climate_aus)
coords(climate_aus)
coord_x(climate_aus)
coord_y(climate_aus)
```

---

geom\_glyph

*Create glyph map with ggplot2*

---

## Description

Create glyph map with ggplot2

## Usage

```
geom_glyph(
  mapping = NULL,
  data = NULL,
  stat = "identity",
  position = "identity",
  ...,
  x_major = NULL,
  x_minor = NULL,
  y_major = NULL,
  y_minor = NULL,
  x_scale = identity,
  y_scale = identity,
  polar = FALSE,
  width = ggplot2::rel(2.1),
  height = ggplot2::rel(1.8),
  global_rescale = TRUE,
  show.legend = NA,
  inherit.aes = TRUE
)

geom_glyph_line(
  mapping = NULL,
  data = NULL,
  stat = "identity",
  position = "identity",
  ...,
```

```

x_major = NULL,
x_minor = NULL,
y_major = NULL,
y_minor = NULL,
polar = FALSE,
width = ggplot2::rel(2.1),
height = ggplot2::rel(2.1),
show.legend = NA,
inherit.aes = TRUE
)

geom_glyph_box(
  mapping = NULL,
  data = NULL,
  stat = "identity",
  position = "identity",
  ...,
  x_major = NULL,
  x_minor = NULL,
  y_major = NULL,
  y_minor = NULL,
  polar = FALSE,
  width = ggplot2::rel(2.1),
  height = ggplot2::rel(2.1),
  show.legend = NA,
  inherit.aes = TRUE
)

```

## Arguments

<code>mapping</code>	Set of aesthetic mappings created by <a href="#">aes()</a> . If specified and <code>inherit.aes = TRUE</code> (the default), it is combined with the default mapping at the top level of the plot. You must supply <code>mapping</code> if there is no plot mapping.
<code>data</code>	The data to be displayed in this layer. There are three options: If <code>NULL</code> , the default, the data is inherited from the plot data as specified in the call to <a href="#">ggplot()</a> . A <code>data.frame</code> , or other object, will override the plot data. All objects will be fortified to produce a data frame. See <a href="#">fortify()</a> for which variables will be created. A function will be called with a single argument, the plot data. The return value must be a <code>data.frame</code> , and will be used as the layer data. A function can be created from a <code>formula</code> (e.g. <code>~ head(.x, 10)</code> ).
<code>stat</code>	The statistical transformation to use on the data for this layer, either as a <code>ggproto</code> <code>Geom</code> subclass or as a string naming the stat stripped of the <code>stat_</code> prefix (e.g. "count" rather than "stat_count")
<code>position</code>	Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use <code>position_jitter</code> ), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.

...	Other arguments passed on to <code>layer()</code> . These are often aesthetics, used to set an aesthetic to a fixed value, like <code>colour = "red"</code> or <code>size = 3</code> . They may also be parameters to the paired geom/stat.
<code>x_major</code> , <code>x_minor</code> , <code>y_major</code> , <code>y_minor</code>	The name of the variable (as a string) for the major and minor x and y axes. Together, each unique combination of <code>x_major</code> and <code>y_major</code> specifies a grid cell.
<code>y_scale</code> , <code>x_scale</code>	The scaling function to be applied to each set of minor values within a grid cell. Defaults to <code>identity</code> so that no scaling is performed.
<code>polar</code>	A logical of length 1, specifying whether the glyphs should be drawn in polar coordinates. Defaults to FALSE.
<code>height</code> , <code>width</code>	The height and width of each glyph. Defaults to 95% of the <code>resolution</code> of the data. Specify the width absolutely by supplying a numeric vector of length 1, or relative to the resolution of the data by using <code>rel</code> .
<code>global_rescale</code>	Whether rescale is performed globally or on each individual glyph.
<code>show.legend</code>	logical. Should this layer be included in the legends? NA, the default, includes if any aesthetics are mapped. FALSE never includes, and TRUE always includes. It can also be a named logical vector to finely select the aesthetics to display.
<code>inherit.aes</code>	If FALSE, overrides the default aesthetics, rather than combining with them. This is most useful for helper functions that define both data and aesthetics and shouldn't inherit behaviour from the default plot specification, e.g. <code>borders()</code> .

## Value

a ggplot object

## Examples

```
print_p <- GGally::print_if_interactive

library(ggplot2)
# basic glyph map with reference line and box-----
p <- ggplot(data = GGally::nasa,
             aes(x_major = long, x_minor = day,
                  y_major = lat, y_minor = surftemp)) +
  geom_glyph_box() +
  geom_glyph_line() +
  geom_glyph() +
  theme_bw()
print_p(p)

# rescale on each individual glyph -----
p <- ggplot(data = GGally::nasa,
             aes(x_major = long, x_minor = day,
                  y_major = lat, y_minor = surftemp)) +
  geom_glyph(global_rescale = FALSE)
print_p(p)
```

```

# adjust width and height with relative & absolute value -----
p <- ggplot() +
  geom_glyph(data = GGally::nasa,
    aes(x_major = long, x_minor = day,
        y_major = lat, y_minor = surftemp),
    width = rel(0.8), height = 1) +
  theme_bw()
print_p(p)

# apply a re-scaling on Y and use polar coordinate
p <-
  GGally::nasa %>%
  ggplot(aes(x_major = long, x_minor = day,
    y_major = lat, y_minor = ozone)) +
  geom_glyph_box(fill=NA) +
  geom_glyph_line() +
  geom_glyph(y_scale = GGally::range01, polar = TRUE)
print_p(p)

```

**get\_centroid***Find the centroid of cubble***Description**

find the convex hull that wraps around the cluster, make it a polygon, find the centroid of the polygon and finally, extract the x and y coordinate of each centroid:

**Usage**

```
get_centroid(data)
```

**Arguments**

data	a cubble data object
------	----------------------

**Value**

a cubble object
-----------------

---

match_sites	<i>Matching sites from two data sources</i>
-------------	---

---

## Description

The function includes both spatial and temporal matching. The spatial matching is based on the distance and the distance is calculated using the Vincenty formula assuming earth is sphere with a radius of 6371 km. The temporal matching first filters out the n largest increases, determined by `temporal_n_highest`, in both datasets, constructs an interval of length `temporal_window` from one dataset and count the number that large increase from the other dataset falls into the interval constructed.

## Usage

```
match_sites(  
    major,  
    minor,  
    spatial_single_match = TRUE,  
    spatial_n_keep = 1,  
    spatial_dist_max = 10,  
    temporal_matching = TRUE,  
    temporal_by,  
    temporal_n_highest = 20,  
    temporal_independent,  
    temporal_window = 5,  
    temporal_min_match = 10  
)  
  
match_spatial(  
    major,  
    minor,  
    spatial_single_match = TRUE,  
    spatial_n_keep = 1,  
    spatial_dist_max = 10  
)  
  
match_postprocessing(major, minor, match_table)  
  
match_temporal(  
    major,  
    minor,  
    temporal_by,  
    temporal_n_highest = 20,  
    temporal_independent,  
    temporal_window = 5,  
    temporal_min_match = 10  
)
```

### Arguments

<code>major</code>	The major dataset to match, every key in the major dataset will have a match, unless filtered by <code>dist_max</code>
<code>minor</code>	The dataset to match from
<code>spatial_single_match</code>	Whether each observation in the minor dataset is only allowed to be matched once, default to TRUE
<code>spatial_n_keep</code>	The number of matching to keep
<code>spatial_dist_max</code>	The maximum distance allowed between matched pair
<code>temporal_matching</code>	Whether to perform temporal matching
<code>temporal_by</code>	The variable used for temporal matching
<code>temporal_n_highest</code>	The number of highest peak used for temporal matching
<code>temporal_independent</code>	The dataset used to construct the temporal window, need to be the name of either major or minor.
<code>temporal_window</code>	The temporal window allowed to fall in
<code>temporal_min_match</code>	The minimum number of peak matching for temporal matching
<code>match_table</code>	The spatial matching table

### Value

A cubble with matched pairs

`prcp_aus`

*Daily precipitation data from 2016 to 2020*

### Description

Daily precipitation data from 2016 to 2020

### Usage

`prcp_aus`

### Format

An object of class `cubicle_df` (inherits from `rowwise_df`, `tbl_df`, `tbl`, `data.frame`) with 663 rows and 7 columns.

---

prep_edges	<i>A function to prepare edges data for tour display</i>
------------	--

---

### Description

A function to prepare edges data for tour display

### Usage

```
prep_edges(data, edges_col, color_col)

## S3 method for class 'cubble_df'
prep_edges(data, edges_col, color_col = NULL)

prep_data(data, cols)

## S3 method for class 'cubble_df'
prep_data(data, cols = NULL)
```

### Arguments

data	a cubble object
edges_col	the variable maps to edges colour
color_col	the variable maps to point colour
cols	the numerical column selected for a tour

### Value

a list of edge linkage, edge color, and point color

---

rename_key	<i>Rename the key variable</i>
------------	--------------------------------

---

### Description

Rename the key variable

### Usage

```
rename_key(data, ...)
```

### Arguments

data	a cubble
...	argument passed to rename: NEW = OLD

**Value**

a cubble object

`river`

*Australia river data*

**Description**

Australia river data

**Usage**

```
river
```

**Format**

An object of class `cubble_df` (inherits from `rowwise_df`, `tbl_df`, `tbl`, `data.frame`) with 71 rows and 5 columns.

`simplify_sf`

*Find (multi)polygons with small area*

**Description**

Find (multi)polygons with small area

**Usage**

```
simplify_sf(data, geom, area, point_threshold = 0.9, area_threshold = 0.9)
```

**Arguments**

<code>data</code>	An sf object
<code>geom</code>	The geometry column
<code>area</code>	The area column if any
<code>point_threshold</code>	The number of point threshold used to define small crumb
<code>area_threshold</code>	The area size threshold used to define small crumb

**Value**

An sf object

the data object with additional column `crumb` indicating whether the area is a "small crumb"

---

slice\_head.cubble\_df    *Slicing a cubble*

---

## Description

Slicing can be useful when the number of site is too large to be all visualised in a single plot. The slicing family in cubble wraps around the `dplyr::slice()` family to allow slicing from top and bottom, based on a variable, or in random.

## Usage

```
## S3 method for class 'cubble_df'
slice_head(data, ...)

## S3 method for class 'cubble_df'
slice_tail(data, ...)

## S3 method for class 'cubble_df'
slice_min(data, ...)

## S3 method for class 'cubble_df'
slice_max(data, ...)

## S3 method for class 'cubble_df'
slice_sample(data, ...)
```

## Arguments

data	a cubble object to slice
...	other arguments passed to the <code>dplyr::slice()</code>

## Value

a cubble object

## Examples

```
# slice the first 50 stations from the top/ bottom
library(dplyr)
climate_aus |> slice_head(n = 50)
climate_aus |> slice_tail(n = 50)

# slice based on the max/ min of a variable

climate_aus |> slice_max(elev, n = 10)
climate_aus |> slice_min(lat, n = 10)

# random sample
climate_aus |> slice_sample(n = 10)
```

slice_nearby	<i>Location-based slicing</i>
--------------	-------------------------------

### Description

Location-based slicing

### Usage

```
slice_nearby(data, coord, buffer, n)
```

### Arguments

data	the data to slice
coord	the coordinate of used to slice nearby locations
buffer	the buffer added to the coordinate for slicing
n	the number of nearby points to slice, based on distance

### Value

a cubble object

### Examples

```
# slice locations within 1 degree of (130E, 25S)
slice_nearby(climate_aus, coord = c(130, -25), buffer = 3)

# slice the 5 closest location to (130E, 25S)
slice_nearby(climate_aus, coord = c(130, -25), n = 5)
```

strip_rowwise	<i>Remove the rowwise grouping of a cubble</i>
---------------	--

### Description

Remove the rowwise grouping of a cubble

### Usage

```
strip_rowwise(data)
```

### Arguments

data	a cubble object
------	-----------------

**Value**

a cubble object

**Examples**

```
library(dplyr)
# row number is not properly added since each row is a separate group
climate_aus |> mutate(.id = row_number())

# proper id after removing the grouping structure
climate_aus |> strip_rowwise() |> mutate(.id = row_number())
```

`switch_key`

*Switch to a different key of a cubble*

**Description**

`switch_key()` allows you select a new variable in the data to become the key. This can be used to create hierarchical data where one variable is nested in another.

**Usage**

```
switch_key(data, key)
```

**Arguments**

data	a cubble object, can be either long or nested cubble
key	the new key

**Value**

a cubble object

**Examples**

```
library(ggplot2)
library(dplyr)
# create an artificial cluster for stations
set.seed(1234)
cb <- climate_flat %>%
  as_cubble(key = id, index = date, coords = c(long, lat)) %>%
  mutate(cluster = sample(1:3, 1))

# switch the key to cluster
cb_hier <- cb %>% switch_key(cluster)
```

---

`tmax_hist`

*Victoria and Tasmania daily maximum temperature for 1970 - 1975  
and 2016 - 2020*

---

### Description

Victoria and Tasmania daily maximum temperature for 1970 - 1975 and 2016 - 2020

### Usage

```
tmax_hist
```

### Format

An object of class `cubble_df` (inherits from `rowwise_df`, `tbl_df`, `tbl`, `data.frame`) with 39 rows and 7 columns.

---

`unfold`

*Move spatial variables into the long form*

---

### Description

Some spatio-temporal transformation, i.e. glyph maps, uses both spatial and temporal variables. `unfold()` allows you to temporarily moves spatial variables into the long form for these transformations.

### Usage

```
unfold(data, ...)
```

### Arguments

<code>data</code>	a long cubble object
<code>...</code>	spatial variables to move into the long form

### Value

a cubble object in the long form

**Examples**

```
cb <- climate_flat |>
  as_cubble(key = id, index = date, coords = c(long, lat)) |>
  face_temporal()

# unfold long and lat
cb_mig <- cb |> unfold(long, lat)

# unfold is not memorised by cubble:
# if you switch to the nested cubble and then switch back,
# long and lat will not be preserved
cb_mig |> face_spatial() |> face_temporal()
```

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