

# Package ‘stars’

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**Title** Spatiotemporal Arrays, Raster and Vector Data Cubes

**Version** 0.5-1

**Description** Reading, manipulating, writing and plotting spatiotemporal arrays (raster and vector data cubes) in 'R', using 'GDAL' bindings provided by 'sf', and 'NetCDF' bindings by 'ncmeta' and 'RNetCDF'.

**License** Apache License

**URL** <https://r-spatial.github.io/stars/>,  
<https://github.com/r-spatial/stars/>

**BugReports** <https://github.com/r-spatial/stars/issues/>

**Additional\_repositories** <http://gis-bigdata.uni-muenster.de/pebesma/>

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---

aggregate.stars	<i>spatially or temporally aggregate stars object</i>
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## Description

spatially or temporally aggregate stars object, returning a data cube with lower spatial or temporal resolution

## Usage

```
## S3 method for class 'stars'
aggregate(
  x,
  by,
  FUN,
  ...,
  drop = FALSE,
  join = st_intersects,
  as_points = any(st_dimension(by) == 2, na.rm = TRUE),
  rightmost.closed = FALSE,
  left.open = FALSE,
  exact = FALSE
)
```

## Arguments

x	object of class stars with information to be aggregated
by	object of class sf or sfc for spatial aggregation, for temporal aggregation a vector with time values (Date, POSIXct, or PCICT) that is interpreted as a sequence of left-closed, right-open time intervals or a string like "months", "5 days" or the like (see <a href="#">cut.POSIXt</a> ); if by is an object of class stars, it is converted to sfc by <code>st_as_sfc(by, as_points = FALSE)</code> thus ignoring its time component.
FUN	aggregation function, such as mean
...	arguments passed on to FUN, such as <code>na.rm=TRUE</code>
drop	logical; ignored
join	function; function used to find matches of x to by
as_points	see <a href="#">st_as_sf</a> : shall raster pixels be taken as points, or small square polygons?
rightmost.closed	see <a href="#">findInterval</a>
left.open	logical; used for time intervals, see <a href="#">findInterval</a> and <a href="#">cut.POSIXt</a>
exact	logical; if TRUE, use <a href="#">coverage_fraction</a> to compute exact overlap fractions of polygons with raster cells

## Examples

```
# aggregate time dimension in format Date
tif = system.file("tif/L7_ETMs.tif", package = "stars")
t1 = as.Date("2018-07-31")
x = read_stars(c(tif, tif, tif, tif), along = list(time = c(t1, t1+1, t1+2, t1+3)))[,1:30,1:30]
st_get_dimension_values(x, "time")
x_agg_time = aggregate(x, by = t1 + c(0, 2, 4), FUN = max)

# aggregate time dimension in format Date - interval
by_t = "2 days"
x_agg_time2 = aggregate(x, by = by_t, FUN = max)
st_get_dimension_values(x_agg_time2, "time")
x_agg_time - x_agg_time2

# aggregate time dimension in format POSIXct
x = st_set_dimensions(x, 4, values = as.POSIXct(c("2018-07-31",
                                                "2018-08-01",
                                                "2018-08-02",
                                                "2018-08-03")),
                    names = "time")
by_t = as.POSIXct(c("2018-07-31", "2018-08-02"))
x_agg_posix = aggregate(x, by = by_t, FUN = max)
st_get_dimension_values(x_agg_posix, "time")
x_agg_time - x_agg_posix
aggregate(x, "2 days", mean)
# Spatial aggregation, see https://github.com/r-spatial/stars/issues/299
prec_file = system.file("nc/test_stageiv_xy.t.nc", package = "stars")
prec = read_ncdf(prec_file, curvilinear = c("lon", "lat"))
prec_slice = dplyr::slice(prec, index = 17, along = "time")
nc = sf::read_sf(system.file("gpkg/nc.gpkg", package = "sf"), "nc.gpkg")
nc = st_transform(nc, st_crs(prec_slice))
agg = aggregate(prec_slice, st_geometry(nc), mean)
plot(agg)
```

---

as

*Coerce stars object into a Raster raster or brick*


---

## Description

Coerce stars object into a Raster raster or brick

## Arguments

from                    object to coerce

## Details

If the stars object has more than three dimensions, all dimensions higher than the third will be collapsed into the third dimensions. If the stars object has only an x/y raster but multiple attributes, these are merged first, then put in a raster brick.

**Value**

RasterLayer or RasterBrick

---

c.stars	<i>combine multiple stars objects, or combine multiple attributes in a single stars object into a single array</i>
---------	--

---

**Description**

combine multiple stars objects, or combine multiple attributes in a single stars object into a single array

**Usage**

```
## S3 method for class 'stars'
c(..., along = NA_integer_, try_hard = FALSE, nms = names(list(...)))

## S3 method for class 'stars_proxy'
c(
  ...,
  along = NA_integer_,
  along_crs = FALSE,
  try_hard = FALSE,
  nms = names(list(...))
)
```

**Arguments**

...	object(s) of class star: in case of multiple arguments, these are combined into a single stars object, in case of a single argument, its attributes are combined into a single attribute. In case of multiple objects, all objects should have the same dimensionality.
along	integer; see <a href="#">read_stars</a>
try_hard	logical; if TRUE and some arrays have different dimensions, combine those that dimensions matching to the first array
nms	character; vector with array names
along_crs	logical; if TRUE, combine arrays along a CRS dimension

**Examples**

```
tif = system.file("tif/L7_ETMs.tif", package = "stars")
x = read_stars(tif)
(new = c(x, x))
c(new) # collapses two arrays into one with an additional dimension
c(x, x, along = 3)
```

---

contour.stars	<i>plot contours of a stars object</i>
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---

**Description**

plot contours of a stars object

**Usage**

```
## S3 method for class 'stars'
contour(x, ...)
```

**Arguments**

x	object of class stars
...	other parameters passed on to <a href="#">contour</a>

**Details**

this uses the R internal contour algorithm, which (by default) plots contours; [st\\_contour](#) uses the GDAL contour algorithm that returns contours as simple features.

**Examples**

```
d = st_dimensions(x = 1:ncol(volcano), y = 1:nrow(volcano))
r = st_as_stars(t(volcano))
r = st_set_dimensions(r, 1, offset = 0, delta = 1)
r = st_set_dimensions(r, 2, offset = 0, delta = -1)
plot(r, reset = FALSE)
contour(r, add = TRUE)
```

---

cut_stars	<i>cut methods for stars objects</i>
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---

**Description**

cut methods for stars objects

**Usage**

```
## S3 method for class 'array'
cut(x, breaks, ...)

## S3 method for class 'matrix'
cut(x, breaks, ...)

## S3 method for class 'stars'
cut(x, breaks, ...)
```

**Arguments**

x                    see [cut](#)  
 breaks                see [cut](#)  
 ...                    see [cut](#)

**Details**

R's factor only works for vectors, not for arrays or matrices. This is a work-around (or hack?) to keep the factor levels generated by `cut` and use them in plots.

**Value**

an array or matrix with a `levels` attribute; see details

**Examples**

```
tif = system.file("tif/L7_ETMs.tif", package = "stars")
x = read_stars(tif)
cut(x, c(0, 50, 100, 255))
cut(x[,,,1], c(0, 50, 100, 255))
plot(cut(x[,,,1], c(0, 50, 100, 255)))
tif = system.file("tif/L7_ETMs.tif", package = "stars")
x1 = read_stars(tif)
(x1_cut = cut(x1, breaks = c(0, 50, 100, Inf))) # shows factor in summary
plot(x1_cut[,,,c(3,6)]) # propagates through [ and plot
```

---

dplyr

*dplyr verbs for stars objects*


---

**Description**

dplyr verbs for stars objects

**Usage**

```
filter.stars(.data, ...)
filter.stars_proxy(.data, ...)
mutate.stars(.data, ...)
mutate.stars_proxy(.data, ...)
transmute.stars(.data, ...)
transmute.stars_proxy(.data, ...)
```

```

select.stars(.data, ...)
select.stars_proxy(.data, ...)
pull.stars(.data, var = -1)
pull.stars_proxy(.data, ...)
as.tbl_cube.stars(x, ...)
slice.stars(.data, along, index, ..., drop = length(index) == 1)
slice.stars_proxy(.data, ...)

```

### Arguments

.data	object of class stars
...	see <a href="#">filter</a>
var	see <a href="#">pull</a>
x	object of class stars
along	name or index of dimension to which the slice should be applied
index	integer value(s) for this index
drop	logical; drop dimensions that only have a single index?

### Examples

```

tif = system.file("tif/L7_ETMs.tif", package = "stars")
x1 = read_stars(tif)
library(dplyr)
x1 %>% slice("band", 2:3)
x1 %>% slice("x", 50:100)

```

---

geom\_stars

*ggplot geom for stars objects*

---

### Description

ggplot geom for stars objects

### Usage

```

geom_stars(mapping = NULL, data = NULL, ..., downsample = 0, sf = FALSE)

theme_stars(...)

```



**Arguments**

mapping	see <a href="#">geom_raster</a>
data	see <a href="#">geom_raster</a>
...	see <a href="#">geom_raster</a>
downsample	downsampling rate: e.g. 3 keeps rows and cols 1, 4, 7, 10 etc.; a value of 0 does not downsample
sf	logical; if TRUE rasters will be converted to polygons and plotted using <a href="#">geom_sf</a> .

**Details**

`geom_stars` returns (a call to) either [geom\\_raster](#), [geom\\_tile](#), or [geom\\_sf](#), depending on the raster or vector geometry; for the first to, an `aes` call is constructed with the raster dimension names and the first array as fill variable. Further calls to [coord\\_equal](#) and [facet\\_wrap](#) are needed to control aspect ratio and the layers to be plotted; see examples.

**Examples**

```
system.file("tif/L7_ETMs.tif", package = "stars") %>% read_stars() -> x
library(ggplot2)
ggplot() + geom_stars(data = x) +
  coord_equal() +
  facet_wrap(~band) +
  theme_void() +
  scale_x_discrete(expand=c(0,0))+
  scale_y_discrete(expand=c(0,0))
```

---

make_intervals	<i>create an intervals object</i>
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---

**Description**

create an intervals object, assuming left-closed and right-open intervals

**Usage**

```
make_intervals(start, end)
```

**Arguments**

start	vector with start values, or 2-column matrix with start and end values in column 1 and 2, respectively
end	vector with end values

---

merge	<i>merge or split stars object</i>
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---

### Description

merge attributes into a dimension, or split a dimension over attributes

### Usage

```
## S3 method for class 'stars'
split(x, f = length(dim(x)), drop = TRUE, ...)

## S3 method for class 'stars'
merge(x, y, ..., name = "attributes")
```

### Arguments

x	object of class stars
f	the name or index of the dimension to split; by default the last dimension
drop	ignored
...	if defined, the first unnamed argument is used for dimension values, if not defined, attribute names are used for dimension values
y	needs to be missing
name	name for the new dimension

### Details

split.stars works on the first attribute, and will give an error when more than one attribute is present

### Value

merge merges attributes of a stars object into a new dimension; split splits a dimension over attributes

---

ops_stars	<i>S3 Ops Group Generic Functions for stars objects</i>
-----------	---

---

### Description

Ops functions for stars objects, including comparison, product and divide, add, subtract

**Usage**

```
## S3 method for class 'stars'
Ops(e1, e2)

## S3 method for class 'stars'
Math(x, ...)

## S3 method for class 'stars_proxy'
Ops(e1, e2)

## S3 method for class 'stars_proxy'
Math(x, ...)
```

**Arguments**

e1	object of class stars
e2	object of class stars
x	object of class stars
...	parameters passed on to the Math functions

**Value**

object of class stars

**Examples**

```
tif = system.file("tif/L7_ETMs.tif", package = "stars")
x = read_stars(tif)
x * x
x / x
x + x
x + 10
all.equal(x * 10, 10 * x)
tif = system.file("tif/L7_ETMs.tif", package = "stars")
x = read_stars(tif)
a = sqrt(x)
b = log(x, base = 10)
```

---

plot	<i>plot stars object, with subplots for each level of first non-spatial dimension</i>
------	---

---

**Description**

plot stars object, with subplots for each level of first non-spatial dimension, and customization of legend key

**Usage**

```

## S3 method for class 'stars'
plot(
  x,
  y,
  ...,
  join_zlim = TRUE,
  main = make_label(x, 1),
  axes = FALSE,
  downsample = TRUE,
  nbreaks = 11,
  breaks = "quantile",
  col = grey(1:(nbreaks - 1)/nbreaks),
  key.pos = get_key_pos(x, ...),
  key.width = lcm(1.8),
  key.length = 0.618,
  reset = TRUE,
  box_col = grey(0.8),
  center_time = FALSE,
  hook = NULL,
  mfrow = NULL
)

## S3 method for class 'stars'
image(
  x,
  ...,
  band = 1,
  attr = 1,
  asp = NULL,
  rgb = NULL,
  maxColorValue = ifelse(inherits(rgb, "data.frame"), 255, max(x[[attr]], na.rm =
    TRUE)),
  xlab = if (!axes) "" else names(d)[1],
  ylab = if (!axes) "" else names(d)[2],
  xlim = st_bbox(extent)$xlim,
  ylim = st_bbox(extent)$ylim,
  text_values = FALSE,
  text_color = "black",
  axes = FALSE,
  interpolate = FALSE,
  as_points = FALSE,
  key.pos = NULL,
  logz = FALSE,
  key.width = lcm(1.8),
  key.length = 0.618,
  add.geom = NULL,
  border = NA,

```

```

    useRaster = isTRUE(dev.capabilities("rasterImage")$rasterImage == "yes"),
    extent = x
)

## S3 method for class 'stars_proxy'
plot(x, y, ..., downsample = get_downsample(dim(x)))

```

### Arguments

<code>x</code>	object of class <code>stars</code>
<code>y</code>	ignored
<code>...</code>	further arguments: for <code>plot</code> , passed on to <code>image.stars</code> ; for <code>image</code> , passed on to <code>image.default</code> or <code>rasterImage</code> .
<code>join_zlim</code>	logical; if <code>TRUE</code> , compute a single, joint <code>zlim</code> (color scale) for all subplots from <code>x</code>
<code>main</code>	character; subplot title prefix; use <code>""</code> to get only time, use <code>NULL</code> to suppress subplot titles
<code>axes</code>	logical; should axes and box be added to the plot?
<code>downsample</code>	logical or numeric; if <code>TRUE</code> will try to plot not many more pixels than actually are visible, if <code>FALSE</code> , no downsampling takes place, if numeric, the downsampling rate; see <code>Details</code> .
<code>nbreaks</code>	number of color breaks; should be one more than number of colors. If missing and <code>col</code> is specified, it is derived from that.
<code>breaks</code>	actual color breaks, or a method name used for <a href="#">classIntervals</a> .
<code>col</code>	colors to use for grid cells
<code>key.pos</code>	integer; side to plot a color key: 1 bottom, 2 left, 3 top, 4 right; set to <code>NULL</code> to omit key. Ignored if multiple columns are plotted in a single function call. Default depends on plot size, map aspect, and, if set, parameter <code>asp</code> .
<code>key.width</code>	amount of space reserved for width of the key (labels); relative or absolute (using <code>lcm</code> )
<code>key.length</code>	amount of space reserved for length of the key (labels); relative or absolute (using <code>lcm</code> )
<code>reset</code>	logical; if <code>FALSE</code> , keep the plot in a mode that allows adding further map elements; if <code>TRUE</code> restore original mode after plotting; see <code>details</code> .
<code>box_col</code>	color for box around sub-plots; use <code>0</code> to suppress plotting of boxes around sub-plots.
<code>center_time</code>	logical; if <code>TRUE</code> , sub-plot titles will show the center of time intervals, otherwise their start
<code>hook</code>	<code>NULL</code> or function; hook function that will be called on every sub-plot.
<code>mfrow</code>	length-2 integer vector with <code>nrows</code> , <code>ncolumns</code> of a composite plot, to override the default layout
<code>band</code>	integer; which band (dimension) to plot
<code>attr</code>	integer; which attribute to plot

asp	numeric; aspect ratio of image
rgb	integer; specify three bands to form an rgb composite. Experimental: rgb color table; see Details.
maxColorValue	numeric; passed on to <a href="#">rgb</a>
xlab	character; x axis label
ylab	character; y axis label
xlim	x axis limits
ylim	y axis limits
text_values	logical; print values as text on image?
text_color	character; color for printed text values
interpolate	logical; when using <a href="#">rasterImage</a> (rgb), should pixels be interpolated?
as_points	logical; for curvilinear or sheared grids: parameter passed on to <a href="#">st_as_sf</a> , determining whether raster cells will be plotted as symbols (fast, approximate) or small polygons (slow, exact)
logz	logical; if TRUE, use log10-scale for the attribute variable. In that case, breaks and at need to be given as log10-values; see examples.
add.geom	object of class <code>sfc</code> , or list with arguments to <code>plot</code> , that will be added to an image or sub-image
border	color used for cell borders (only in case x is a curvilinear or rotated/sheared grid)
useRaster	logical; use the <code>rasterImage</code> capabilities of the graphics device?
extent	object which has a <code>st_bbox</code> method; sets the plotting extent

## Details

Downsampling: a value for `downsample` of 0 or 1 causes no downsampling, 2 that every second dimension value is sampled, 3 that every third dimension value is sampled, and so on.

use of an rgb color table is experimental; see <https://github.com/r-spatial/mapview/issues/208>

when plotting a subsetting `stars_proxy` object, the default value for argument `downsample` will not be computed correctly, and it has to be set manually.

## Examples

```
tif = system.file("tif/L7_ETMs.tif", package = "stars")
x = read_stars(tif)
image(x, col = grey((3:9)/10))
image(x, rgb = c(1,3,5)) # rgb composite
```

---

predict.stars	<i>Predict values, given a model object, for a stars or stars_proxy object</i>
---------------	--

---

**Description**

Predict values, given a model object, for a stars or stars\_proxy object

**Usage**

```
## S3 method for class 'stars'
predict(object, model, ...)

## S3 method for class 'stars_proxy'
predict(object, model, ...)
```

**Arguments**

object	object of class 'stars'
model	model object of a class that has a predict method; check with 'methods(class = class(object))'
...	arguments passed on to this predict method

**Details**

separate predictors in object need to be separate attributes in object; in case they are e.g. in a band dimension, use 'split(object)'

---

read_ncdf	<i>Read NetCDF into stars object</i>
-----------	--------------------------------------

---

**Description**

Read data from a file (or source) using the NetCDF library directly.

**Usage**

```
read_ncdf(
  .x,
  ...,
  var = NULL,
  ncsb = NULL,
  curvilinear = character(0),
  eps = 1e-12,
  ignore_bounds = FALSE,
  make_time = TRUE,
  make_units = TRUE
)
```

**Arguments**

<code>.x</code>	NetCDF file or source
<code>...</code>	ignored
<code>var</code>	variable name or names (they must be on matching grids)
<code>ncsub</code>	matrix of start, count columns (see Details)
<code>curvilinear</code>	length two character named vector with names of variables holding longitude and latitude values for all raster cells. 'stars' attempts to figure out appropriate curvilinear coordinates if they are not supplied.
<code>eps</code>	numeric; dimension value increases are considered identical when they differ less than eps
<code>ignore_bounds</code>	logical; should bounds values for dimensions, if present, be ignored?
<code>make_time</code>	if TRUE (the default), an attempt is made to provide a date-time class from the "time" variable
<code>make_units</code>	if TRUE (the default), an attempt is made to set the units property of each variable

**Details**

The following logic is applied to coordinates. If any coordinate axes have regularly spaced coordinate variables they are reduced to the offset/delta form with 'affine = c(0, 0)', otherwise the values of the coordinates are stored and used to define a rectilinear grid.

If the data has two or more dimensions and the first two are regular they are nominated as the 'raster' for plotting.

If the `curvilinear` argument is used it specifies the 2D arrays containing coordinate values for the first two dimensions of the data read. It is currently assumed that the coordinates are 2D and that they relate to the first two dimensions in that order.

If `var` is not set the first set of variables on a shared grid is used.

`start` and `count` columns of `ncsub` must correspond to the variable dimension (`nrows`) and be valid index using `var.get.nc` convention (`start` is 1-based). If the count value is NA then all steps are included. Axis order must match that of the variable/s being read.

**Examples**

```
f <- system.file("nc/reduced.nc", package = "stars")
read_ncdf(f)
read_ncdf(f, var = c("anom"))
read_ncdf(f, ncsub = cbind(start = c(1, 1, 1, 1), count = c(10, 12, 1, 1)))

#' precipitation data in a curvilinear NetCDF
prec_file = system.file("nc/test_stageiv_xyt.nc", package = "stars")
prec = read_ncdf(prec_file, curvilinear = c("lon", "lat"), ignore_bounds = TRUE)

##plot(prec) ## gives error about unique breaks
## remove NAs, zeros, and give a large number
## of breaks (used for validating in detail)
qu_0_omit = function(x, ..., n = 22) {
```



```

x = units::drop_units(na.omit(x))
c(0, quantile(x[x > 0], seq(0, 1, length.out = n)))
}
library(dplyr)
prec_slice = slice(prec, index = 17, along = "time")
plot(prec_slice, border = NA, breaks = qu_0_omit(prec_slice[[1]]), reset = FALSE)
nc = sf::read_sf(system.file("gpkg/nc.gpkg", package = "sf"), "nc.gpkg")
plot(st_geometry(nc), add = TRUE, reset = FALSE, col = NA)

```

---

read_stars	<i>read raster/array dataset from file or connection</i>
------------	--

---

## Description

read raster/array dataset from file or connection

## Usage

```

read_stars(
  .x,
  ...,
  options = character(0),
  driver = character(0),
  sub = TRUE,
  quiet = FALSE,
  NA_value = NA_real_,
  along = NA_integer_,
  RasterIO = list(),
  proxy = !length(curvilinear) && is_big(.x, sub = sub, driver = driver, ...),
  curvilinear = character(0),
  normalize_path = TRUE,
  RAT = character(0)
)

is_big(x, ..., sub = sub, n_proxy = options("stars.n_proxy")[[1]] %||% 1e+08)

```

## Arguments

.x	character vector with name(s) of file(s) or data source(s) to be read
...	passed on to <a href="#">st_as_stars</a> if curvilinear was set
options	character; opening options
driver	character; driver to use for opening file. To override fixing for subdatasets and autodetect them as well, use NULL.
sub	character, integer or logical; name, index or indicator of sub-dataset(s) to be read
quiet	logical; print progress output?

NA_value	numeric value to be used for conversion into NA values; by default this is read from the input file
along	length-one character or integer, or list; determines how several arrays are combined, see Details.
RasterIO	list with named parameters for GDAL's RasterIO, to further control the extent, resolution and bands to be read from the data source; see details.
proxy	logical; if TRUE, an object of class stars_proxy is read which contains array metadata only; if FALSE the full array data is read in memory. Always FALSE for curvilinear grids. If not set, defaults to TRUE when the number of cells to be read is larger than options(stars.n_proxy, or to 1e8 if that option was not set.
curvilinear	length two character vector with names of subdatasets holding longitude and latitude values for all raster cells, or named length 2 list holding longitude and latitude matrices; the names of this list should correspond to raster dimensions referred to
normalize_path	logical; if FALSE, suppress a call to <a href="#">normalizePath</a> on .x
RAT	character; raster attribute table column name to use as factor levels
x	object to be read with <a href="#">read_stars</a>
n_proxy	integer; number of cells above which .x will be read as stars proxy object, i.e. not as in-memory arrays but left on disk

## Details

In case .x contains multiple files, they will all be read and combined with [c.stars](#). Along which dimension, or how should objects be merged? If along is set to NA it will merge arrays as new attributes if all objects have identical dimensions, or else try to merge along time if a dimension called time indicates different time stamps. A single name (or positive value) for along will merge along that dimension, or create a new one if it does not already exist. If the arrays should be arranged along one of more dimensions with values (e.g. time stamps), a named list can be passed to along to specify them; see example.

RasterIO is a list with zero or more of the following named arguments: nXOff, nYOff (both 1-based: the first row/col has offset value 1), nXSize, nYSize, nBufXSize, nBufYSize, bands, coderesample. see <https://www.gdal.org/classGDALDataset.html#a80d005ed10aefafa8a55dc539c2f69da> for their meaning; bands is an integer vector containing the band numbers to be read (1-based: first band is 1) Note that if nBufXSize or nBufYSize are specified for downsampling an image, resulting in an adjusted geotransform. resample reflects the resampling method and has to be one of: "nearest\_neighbour" (the default), "bilinear", "cubic", "cubic\_spline", "lanczos", "average", "mode", or "Gauss".

## Value

object of class stars

## Examples

```
tif = system.file("tif/L7_ETMs.tif", package = "stars")
(x1 = read_stars(tif))
(x2 = read_stars(c(tif, tif)))
```

```

(x3 = read_stars(c(tif, tif), along = "band"))
(x4 = read_stars(c(tif, tif), along = "new_dimensions")) # create 4-dimensional array
x1o = read_stars(tif, options = "OVERVIEW_LEVEL=1")
t1 = as.Date("2018-07-31")
# along is a named list indicating two dimensions:
read_stars(c(tif, tif, tif, tif), along = list(foo = c("bar1", "bar2"), time = c(t1, t1+2)))

m = matrix(1:120, nrow = 12, ncol = 10)
dim(m) = c(x = 10, y = 12) # named dim
st = st_as_stars(m)
attr(st, "dimensions")$y$delta = -1
attr(st, "dimensions")$y$offset = 12
st
tmp = tempfile(fileext = ".tif")
write_stars(st, tmp)
(red <- read_stars(tmp))
read_stars(tmp, RasterIO = list(nXOff = 1, nYOff = 1, nXSize = 10, nYSize = 12,
  nBufXSize = 2, nBufYSize = 2))[[1]]
(red <- read_stars(tmp, RasterIO = list(nXOff = 1, nYOff = 1, nXSize = 10, nYSize = 12,
  nBufXSize = 2, nBufYSize = 2)))
red[[1]] # cell values of subsample grid:
## Not run:
plot(st, reset = FALSE, axes = TRUE, ylim = c(-.1,12.1), xlim = c(-.1,10.1),
  main = "nBufXSize & nBufYSize demo", text_values = TRUE)
plot(st_as_sfc(red, as_points = TRUE), add = TRUE, col = 'red', pch = 16)
plot(st_as_sfc(st_as_stars(st), as_points = FALSE), add = TRUE, border = 'grey')
plot(st_as_sfc(red, as_points = FALSE), add = TRUE, border = 'green', lwd = 2)

## End(Not run)
file.remove(tmp)

```

---

redimension

*redimension array, or collapse attributes into a new dimension*


---

## Description

redimension array, or collapse attributes into a new dimension

## Usage

```
st_redimension(x, new_dims, along, ...)
```

```

## S3 method for class 'stars'
st_redimension(
  x,
  new_dims = st_dimensions(x),
  along = list(new_dim = names(x)),
  ...
)

```

```
## S3 method for class 'stars_proxy'
st_redimension(
  x,
  new_dims = st_dimensions(x),
  along = list(new_dim = names(x)),
  ...
)
```

### Arguments

x	object of class stars
new_dims	target dimensions: either a 'dimensions' object or an integer vector with the dimensions' sizes
along	named list with new dimension name and values
...	ignored

---

stars_subset	<i>subset stars objects</i>
--------------	-----------------------------

---

### Description

subset stars objects

### Usage

```
## S3 method for class 'stars'
x[i = TRUE, ..., drop = FALSE, crop = !is_curvilinear(x)]

## S3 replacement method for class 'stars'
x[i] <- value

st_flip(x, which = 1)
```

### Arguments

x	object of class stars
i	first selector: integer, logical or character vector indicating attributes to select, or object of class sf or sfc used as spatial selector; see details
...	further (logical or integer vector) selectors, matched by order, to select on individual dimensions
drop	logical; if TRUE, degenerate dimensions (with only one value) are dropped
crop	logical; if TRUE and parameter i is a spatial geometry (sf or sfc) object, the extent (bounding box) of the result is cropped to match the extent of i using <a href="#">st_crop</a> . Cropping curvilinear grids is not supported.

value	array of dimensions equal to those in x, or a vector or value that will be recycled to such an array
which	character or integer; dimension(s) to be flipped

### Details

if *i* is an object of class *sf*, *sfc* or *bbox*, the spatial subset covering this geometry is selected, possibly followed by cropping the extent. Array values for which the cell centre is not inside the geometry are assigned NA.

in an assignment (or replacement form, [*<-*]), argument *i* needs to be a *stars* object with dimensions identical to *x*, and *value* will be recycled to the dimensions of the arrays in *x*.

### Value

*st\_flip* flips (reverts) the array values along the chosen dimension without(s) changing the dimension properties

### Examples

```
tif = system.file("tif/L7_ETMs.tif", package = "stars")
x = read_stars(tif)
x[,,,1:3] # select bands
x[,1:100,100:200,] # select x and y by range
x["L7_ETMs.tif"] # select attribute
xy = structure(list(x = c(293253.999046018, 296400.196497684), y = c(9113801.64775462,
9111328.49619133)), .Names = c("x", "y"))
pts = st_as_sf(data.frame(do.call(cbind, xy)), coords = c("x", "y"), crs = st_crs(x))
image(x, axes = TRUE)
plot(st_as_sfc(st_bbox(pts)), col = NA, add = TRUE)
bb = st_bbox(pts)
(xx = x[bb])
image(xx)
plot(st_as_sfc(bb), add = TRUE, col = NA)
image(x)
pt = st_point(c(x = 290462.103109179, y = 9114202.32594085))
buf = st_buffer(st_sfc(pt, crs = st_crs(x)), 1500)
plot(buf, add = TRUE)

buf = st_sfc(st_polygon(list(st_buffer(pt, 1500)[[1]], st_buffer(pt, 1000)[[1]])),
  crs = st_crs(x))
image(x[buf])
plot(buf, add = TRUE, col = NA)
image(x[buf, crop=FALSE])
plot(buf, add = TRUE, col = NA)
lc = read_stars(system.file("tif/lc.tif", package = "stars"))
x = c(orig = lc,
      flip_x = st_flip(lc, "x"),
      flip_y = st_flip(lc, "y"),
      flip_xy = st_flip(lc, c("x", "y")),
      along = 3)
plot(x)
```

---

st\_apply

*st\_apply apply a function to one or more array dimensions*


---

## Description

st\_apply apply a function to array dimensions: aggregate over space, time, or something else

## Usage

```
## S3 method for class 'stars'
st_apply(
  X,
  MARGIN,
  FUN,
  ...,
  CLUSTER = NULL,
  PROGRESS = FALSE,
  FUTURE = FALSE,
  rename = TRUE,
  .fname
)
```

## Arguments

X	object of class stars
MARGIN	see <a href="#">apply</a> ; index number(s) or name(s) of the dimensions over which FUN will be applied
FUN	see <a href="#">apply</a>
...	arguments passed on to FUN
CLUSTER	cluster to use for parallel apply; see <a href="#">makeCluster</a>
PROGRESS	logical; if TRUE, use pbapply::pbapply to show progress bar
FUTURE	logical; if TRUE, use future.apply::future_apply
rename	logical; if TRUE and X has only one attribute and FUN is a simple function name, rename the attribute of the returned object to the function name
.fname	function name for the new attribute name (if one or more dimensions are reduced) or the new dimension (if a new dimension is created); if missing, the name of FUN is used

## Value

object of class stars with accordingly reduced number of dimensions; in case FUN returns more than one value, a new dimension is created carrying the name of the function used; see the examples.

**Examples**

```
tif = system.file("tif/L7_ETMs.tif", package = "stars")
x = read_stars(tif)
st_apply(x, 1:2, mean) # mean band value for each pixel
st_apply(x, c("x", "y"), mean) # equivalent to the above
st_apply(x, 3, mean) # mean of all pixels for each band
st_apply(x, "band", mean) # equivalent to the above
st_apply(x, 1:2, range) # min and max band value for each pixel
# to get a progress bar also in non-interactive mode, specify:
if (require(pbapply)) { # install it, if FALSE
  pboptions(type = "timer")
}
```

st\_as\_sf

*Convert stars object into an sf object***Description**

Convert stars object into an sf object

**Usage**

```
## S3 method for class 'stars'
st_as_sf(x, ..., as_points, which = seq_len(prod(dim(x)[1:2])))

## S3 method for class 'stars'
st_as_sf(
  x,
  ...,
  as_points = FALSE,
  merge = FALSE,
  na.rm = TRUE,
  use_integer = is.logical(x[[1]]) || is.integer(x[[1]]),
  long = FALSE,
  connect8 = FALSE
)
```

**Arguments**

x	object of class stars
...	ignored
as_points	logical; should cells be converted to points or to polygons? See details.
which	linear index of cells to keep (this argument is not recommended to be used)
merge	logical; if TRUE, cells with identical values are merged (using GDAL_Polygonize or GDAL_FPolygonize); if FALSE, a polygon for each raster cell is returned; see details

na.rm	logical; should missing valued cells be removed, or also be converted to features?
use_integer	(relevant only if merge is TRUE): if TRUE, before polygonizing values are rounded to 32-bits signed integer values (GDALPolygonize), otherwise they are converted to 32-bit floating point values (GDALFPolygonize).
long	logical; if TRUE, return a long table form sf, with geometries and other dimensions recycled
connect8	logical; if TRUE, use 8 connectedness. Otherwise the 4 connectedness algorithm will be applied.

### Details

If merge is TRUE, only the first attribute is converted into an sf object. If na.rm is FALSE, areas with NA values are also written out as polygons. Note that the resulting polygons are typically invalid, and use [st\\_make\\_valid](#) to create valid polygons out of them.

### Examples

```
tif = system.file("tif/L7_ETMs.tif", package = "stars")
x = read_stars(tif)
x = x[,1:100,1:100,6] # subset of a band with lower values in it
x[[1]][x[[1]] < 30] = NA # set lower values to NA
x[[1]] = x[[1]] < 100 # make the rest binary
x
(p = st_as_sf(x)) # removes NA areas
(p = st_as_sf(x[,,,1], merge = TRUE)) # glues polygons together
all(st_is_valid(p)) # not all valid, see details
plot(p, axes = TRUE)
(p = st_as_sf(x, na.rm = FALSE, merge = TRUE)) # includes polygons with NA values
plot(p, axes = TRUE)
```

---

st_as_stars	<i>convert objects into a stars object</i>
-------------	--

---

### Description

convert objects into a stars object

### Usage

```
st_as_stars(.x, ...)

## S3 method for class 'list'
st_as_stars(.x, ..., dimensions = NULL)

## Default S3 method:
st_as_stars(.x = NULL, ..., raster = NULL)
```



```
## S3 method for class 'stars'  
st_as_stars(.x, ..., curvilinear = NULL, crs = st_crs(4326))  
  
## S3 method for class 'bbox'  
st_as_stars(  
  .x,  
  ...,  
  nx,  
  ny,  
  dx = dy,  
  dy = dx,  
  xlim = .x[c("xmin", "xmax")],  
  ylim = .x[c("ymin", "ymax")],  
  values = 0,  
  n = 64800,  
  pretty = FALSE,  
  inside = FALSE,  
  nz  
)  
  
## S3 method for class 'sf'  
st_as_stars(.x, ..., name = attr(.x, "sf_column"))  
  
## S3 method for class 'Raster'  
st_as_stars(.x, ..., att = 1, ignore_file = FALSE)  
  
## S3 method for class 'ncdfgeom'  
st_as_stars(.x, ..., sf_geometry = NA)  
  
## S3 method for class 'stars_proxy'  
st_as_stars(  
  .x,  
  ...,  
  downsample = 0,  
  url = attr(.x, "url"),  
  envir = parent.frame()  
)  
  
## S3 method for class 'data.frame'  
st_as_stars(.x, ..., dims = 1:2, xy = dims[1:2], y_decreasing = TRUE)  
  
## S3 method for class 'xts'  
st_as_stars(.x, ..., dimensions)  
  
## S3 method for class 'OpenStreetMap'  
st_as_stars(.x, ..., as_col = FALSE)
```

**Arguments**

<code>.x</code>	object to convert
<code>...</code>	in case <code>.x</code> is of class <code>bbox</code> , arguments passed on to <a href="#">pretty</a>
<code>dimensions</code>	object of class <code>dimensions</code>
<code>raster</code>	character; the names of the dimensions that denote raster dimensions
<code>curvilinear</code>	only for creating curvilinear grids: named length 2 list holding longitude and latitude matrices; the names of this list should correspond to raster dimensions referred to
<code>crs</code>	object of class <code>crs</code> with the coordinate reference system of the values in <code>curvilinear</code> ; see details
<code>nx</code>	integer; number of cells in x direction; see details
<code>ny</code>	integer; number of cells in y direction; see details
<code>dx</code>	numeric; cell size in x direction; see details
<code>dy</code>	numeric; cell size in y direction; see details
<code>xlim</code>	length 2 numeric vector with extent (min, max) in x direction
<code>ylim</code>	length 2 numeric vector with extent (min, max) in y direction
<code>values</code>	value(s) to populate the raster values with
<code>n</code>	the (approximate) target number of grid cells
<code>pretty</code>	logical; should cell coordinates have <a href="#">pretty</a> values?
<code>inside</code>	logical; should all cells entirely fall inside the <code>bbox</code> , potentially not covering it completely?
<code>nz</code>	integer; number of cells in z direction; if missing no z-dimension is created.
<code>name</code>	character; name for the geometry dimensions
<code>att</code>	see <a href="#">factorValues</a> ; column in the <code>RasterLayer</code> 's attribute table
<code>ignore_file</code>	logical; if TRUE, ignore the Raster object file name
<code>sf_geometry</code>	<code>sf</code> data.frame with geometry and attributes to be added to stars object. Must have same number of rows as timeseries instances.
<code>downsample</code>	integer: if larger than 0, downsample with this rate (number of pixels to skip in every row/column); if length 2, specifies downsampling rate in x and y.
<code>url</code>	character; URL of the stars endpoint where the data reside
<code>envir</code>	environment to resolve objects in
<code>dims</code>	the column names or indices that form the cube dimensions
<code>xy</code>	the x and y raster dimension names or indices; only takes effect after <code>dims</code> has been specified
<code>y_decreasing</code>	logical; if TRUE, (numeric) y values get a negative delta (decrease with increasing index)
<code>as_col</code>	logical; return rgb numbers (FALSE) or (character) color values (TRUE)?

## Details

if `curvilinear` is a stars object with longitude and latitude values, its coordinate reference system is typically not that of the latitude and longitude values.

For the `bbox` method: if `pretty` is `TRUE`, raster cells may extend the coordinate range of `.x` on all sides. If in addition to `nx` and `ny`, `dx` and `dy` are also missing, these are set to a single value computed as  $\sqrt{\text{diff}(x\text{lim}) * \text{diff}(y\text{lim}) / n}$ . If `nx` and `ny` are missing, they are computed as the ceiling of the ratio of the (x or y) range divided by (dx or dy), unless `inside` is `TRUE`, in which case ceiling is replaced by floor. Postive `dy` will be made negative. Further named arguments (...) are passed on to `pretty`.

For the `ncdfgeom` method: objects are point-timeseries with optional line or polygon geometry for each timeseries specified with the `sf_geometry` parameter. See **ncdfgeom** for more about this NetCDF-based format for geometry and timeseries.

for the `xts` methods, if dimensions are provided, time has to be the first dimension.

## Examples

```
data(Produc, package = "plm")
st_as_stars(Produc, y_decreasing = FALSE)
```

---

st\_contour

*Compute or plot contour lines or sets*

---

## Description

Compute contour lines or sets

## Usage

```
st_contour(
  x,
  na.rm = TRUE,
  contour_lines = FALSE,
  breaks = classInt::classIntervals(na.omit(as.vector(x[[1]])))$brks
)
```

## Arguments

<code>x</code>	object of class stars
<code>na.rm</code>	logical; should missing valued cells be removed, or also be converted to features?
<code>contour_lines</code>	logical; if <code>FALSE</code> , polygons are returned (contour sets), otherwise contour lines
<code>breaks</code>	numerical; values at which to "draw" contour levels

## Details

this function requires GDAL >= 2.4.0

**See Also**

for polygonizing rasters following grid boundaries, see [st\\_as\\_sf](#) with arguments `as_points=FALSE` and `merge=TRUE`; [contour](#) plots contour lines using R's native algorithm (which also plots contour levels)

---

st_coordinates	<i>retrieve coordinates for raster or vector cube cells</i>
----------------	---

---

**Description**

retrieve coordinates for raster or vector cube cells

**Usage**

```
## S3 method for class 'stars'
st_coordinates(x, ..., add_max = FALSE, center = TRUE)

## S3 method for class 'stars'
as.data.frame(x, ..., add_max = FALSE, center = NA)

as_tibble.stars(.x, ..., add_max = FALSE, center = NA)
```

**Arguments**

x	object of class stars
...	ignored
add_max	logical; if TRUE, dimensions are given with a min (x) and max (x_max) value
center	logical; (only if add_max is FALSE): should grid cell center coordinates be returned (TRUE) or offset values (FALSE)? center can be a named logical vector or list to specify values for each dimension.
.x	object to be converted to a tibble

---

st_crop	<i>crop a stars object</i>
---------	----------------------------

---

**Description**

crop a stars object

**Usage**

```
## S3 method for class 'stars_proxy'
st_crop(
  x,
  y,
  ...,
  crop = TRUE,
  epsilon = sqrt(.Machine$double.eps),
  collect = TRUE
)

## S3 method for class 'stars'
st_crop(
  x,
  y,
  ...,
  crop = TRUE,
  epsilon = sqrt(.Machine$double.eps),
  as_points = all(st_dimension(y) == 2, na.rm = TRUE)
)
```

**Arguments**

x	object of class stars
y	object of class sf, sfc or bbox; see Details below.
...	ignored
crop	logical; if TRUE, the spatial extent of the returned object is cropped to still cover obj, if FALSE, the extent remains the same but cells outside y are given NA values.
epsilon	numeric; factor to shrink the bounding box of y towards its center before cropping.
collect	logical; if TRUE, repeat cropping on stars object, i.e. after data has been read
as_points	logical; only relevant if y is of class sf or sfc: if FALSE, treat x as a set of points, else as a set of small polygons. Default: TRUE if y is two-dimensional, else FALSE; see Details

**Details**

for raster x, st\_crop selects cells that intersect with y. For intersection, are raster cells interpreted as points or as small polygons? If y is of class stars, x raster cells are interpreted as points; if y is of class bbox, x cells are interpreted as cells (small polygons). Otherwise, if as\_points is not given, cells are interpreted as points if y has a two-dimensional geometry.

**Examples**

```
l7 = read_stars(system.file("tif/L7_ETMs.tif", package = "stars"))
d = st_dimensions(l7)
```

```

# area around cells 3:10 (x) and 4:11 (y):
offset = c(d[["x"]]$offset, d[["y"]]$offset)
res = c(d[["x"]]$delta, d[["y"]]$delta)
bb = st_bbox(c(xmin = offset[1] + 2 * res[1],
ymin = offset[2] + 11 * res[2],
xmax = offset[1] + 10 * res[1],
ymax = offset[2] + 3 * res[2]), crs = st_crs(17))
l7[bb]

plot(l7[,1:13,1:13,1], reset = FALSE)
image(l7[bb,,1], add = TRUE, col = sf.colors())
plot(st_as_sf(bb), add = TRUE, border = 'green', lwd = 2)

# slightly smaller bbox:
bb = st_bbox(c(xmin = offset[1] + 2.1 * res[1],
ymin = offset[2] + 10.9 * res[2],
xmax = offset[1] + 9.9 * res[1],
ymax = offset[2] + 3.1 * res[2]), crs = st_crs(17))
l7[bb]

plot(l7[,1:13,1:13,1], reset = FALSE)
image(l7[bb,,1], add = TRUE, col = sf.colors())
plot(st_as_sf(bb), add = TRUE, border = 'green', lwd = 2)

# slightly larger bbox:
bb = st_bbox(c(xmin = offset[1] + 1.9 * res[1],
ymin = offset[2] + 11.1 * res[2],
xmax = offset[1] + 10.1 * res[1],
ymax = offset[2] + 2.9 * res[2]), crs = st_crs(17))
l7[bb]

plot(l7[,1:13,1:13,1], reset = FALSE)
image(l7[bb,,1], add = TRUE, col = sf.colors())
plot(st_as_sf(bb), add = TRUE, border = 'green', lwd = 2)

# half a cell size larger bbox:
bb = st_bbox(c(xmin = offset[1] + 1.49 * res[1],
ymin = offset[2] + 11.51 * res[2],
xmax = offset[1] + 10.51 * res[1],
ymax = offset[2] + 2.49 * res[2]), crs = st_crs(17))
l7[bb]

plot(l7[,1:13,1:13,1], reset = FALSE)
image(l7[bb,,1], add = TRUE, col = sf.colors())
plot(st_as_sf(bb), add = TRUE, border = 'green', lwd = 2)

```

---

st\_dimensions

*get dimensions from stars object*


---

## Description

get dimensions from stars object

**Usage**

```

st_dimensions(.x, ...)

## S3 method for class 'stars'
st_dimensions(.x, ...)

st_dimensions(x) <- value

## S3 replacement method for class 'stars'
st_dimensions(x) <- value

## S3 replacement method for class 'list'
st_dimensions(x) <- value

## S3 method for class 'array'
st_dimensions(.x, ...)

## Default S3 method:
st_dimensions(
  .x,
  ...,
  .raster,
  affine = c(0, 0),
  cell_midpoints = FALSE,
  point = FALSE
)

st_set_dimensions(
  .x,
  which,
  values = NULL,
  point = NULL,
  names = NULL,
  xy,
  ...
)

st_get_dimension_values(.x, which, ..., max = FALSE, center = NA)

```

**Arguments**

.x	object to retrieve dimensions information from
...	further arguments
x	object of class dimensions
value	new object of class dimensions, with matching dimensions
.raster	length 2 character array with names (if any) of the raster dimensions
affine	numeric; specify parameters of the affine transformation

cell_midpoints	logical; if TRUE AND the dimension values are strictly regular, the values are interpreted as the cell midpoint values rather than the cell offset values when calculating offset (i.e., the half-cell-size correction is applied); can have a value for each dimension, or else is recycled
point	logical; does the pixel value (measure) refer to a point (location) value or to a pixel (area) summary value?
which	integer or character; index or name of the dimension to be changed
values	values for this dimension (e.g. sfc list-column), or length-1 dimensions object
names	character; vector with new names for all dimensions, or with the single new name for the dimension indicated by which
xy	length-2 character vector; (new) names for the x and y raster dimensions
max	logical; if TRUE return the end, rather than the beginning of an interval
center	logical; if TRUE return the center of an interval; if NA return the center for raster dimensions, and the start of intervals in other cases

### Details

dimensions can be specified in two ways. The simplest is to pass a vector with numeric values for a numeric dimension, or character values for a categorical dimension. Parameter `cell_midpoints` is used to specify whether numeric values refer to the offset (start) of a dimension interval (default), or to the center; the center case is only available for regular dimensions. For rectilinear numeric dimensions, one can specify either a vector with cell borders (start values), or a data.frame with two columns named "start" and "end", with the respective interval start and end values. In the first case, the end values are computed from the start values by assuming the last two intervals have equal width.

### Value

the dimensions attribute of `x`, of class `dimensions`

### Examples

```
x = read_stars(system.file("tif/L7_ETMs.tif", package = "stars"))
# Landsat 7 ETM+ band semantics: https://landsat.gsfc.nasa.gov/the-enhanced-thematic-mapper-plus/
# set bands to values 1,2,3,4,5,7:
(x1 = st_set_dimensions(x, "band", values = c(1,2,3,4,5,7), names = "band_number", point = TRUE))
# set band values as bandwidth
rbind(c(0.45,0.515), c(0.525,0.605), c(0.63,0.69), c(0.775,0.90), c(1.55,1.75), c(2.08,2.35)) %>%
  units::set_units("um") -> bw # or: units::set_units(um) -> bw
# set bandwidth midpoint:
(x2 = st_set_dimensions(x, "band", values = 0.5 * (bw[,1]+bw[,2]),
  names = "bandwidth_midpoint", point = TRUE))
# set bandwidth intervals:
(x3 = st_set_dimensions(x, "band", values = make_intervals(bw), names = "bandwidth"))
```



---

st_dim_to_attr	<i>create an array with dimension values</i>
----------------	--

---

**Description**

create an array with dimension values

**Usage**

```
st_dim_to_attr(x, which = seq_along(dim(x)))
```

**Arguments**

x	object of class stars
which	integer; indices of the dimensions to address (default: all)

**Value**

stars object with dimension values as attributes

**Examples**

```
tif = system.file("tif/L7_ETMs.tif", package = "stars")
x1 = read_stars(tif)
(x = st_dim_to_attr(x1))
plot(x)
(x = st_dim_to_attr(x1, 2:3))
plot(x)
(x = st_dim_to_attr(x1, 3))
plot(x)
```

---

st_extract	<i>Extract cell values at point locations</i>
------------	---

---

**Description**

Extract cell values at point locations

**Usage**

```
st_extract(x, ...)

## S3 method for class 'stars'
st_extract(
  x,
  pts,
```

```

    ...,
    bilinear = FALSE,
    time_column = attr(pts, "time_column") %||% attr(pts, "time_col"),
    interpolate_time = bilinear
  )

```

### Arguments

x	object of class stars or stars_proxy
...	ignored
pts	object of class sf or sfc with POINT geometries
bilinear	logical; use bilinear interpolation rather than nearest neighbour?
time_column	character or integer; name or index of a column with time or date values that will be matched to values of the dimension "time" in x, after which this dimension is reduced. This is useful to extract data cube values along a trajectory; see <a href="https://github.com/r-spatial/stars/issues/352">https://github.com/r-spatial/stars/issues/352</a> .
interpolate_time	logical; should time be interpolated? if FALSE, time instances are matched using the coinciding or the last preceding time in the data cube.

### Details

points outside the raster are returned as NA values.

### Value

if x has more dimensions than only x and y (raster), an object of class stars with POINT geometries replacing x and y raster dimensions; otherwise an object of sf with extracted values.

### Examples

```

tif = system.file("tif/L7_ETMs.tif", package = "stars")
r = read_stars(tif)
pnt = st_sample(st_as_sfc(st_bbox(r)), 10)
st_extract(r, pnt)
st_extract(r, pnt) %>% st_as_sf()
st_extract(r[, , 1], pnt)

```

---

st\_intersects.stars    *spatial intersect predicate for stars and sfc object*

---

### Description

spatial intersect predicate for stars and sfc object

**Usage**

```
## S3 method for class 'stars'
st_intersects(x, y, sparse = TRUE, ..., as_points = NA, transpose = FALSE)
```

**Arguments**

x	object of class stars
y	object that has an 'st_geometry' method: of class 'sf' or 'sfc', or 'stars' object with an 'sfc' dimension
sparse	logical; if TRUE, return the a sparse logical matrix (object of class 'sgbp'), if FALSE, return a logical matrix
...	ignored, or passed on to 'st_intersects.sf' for curvilinear grids
as_points	logical, should grid cells be considered as points (TRUE) or polygons (FALSE)? Default: FALSE and warning emitted
transpose	logical; should the transpose of the 'sgbp' object be returned?

**Details**

curvilinear grids are always converted to polygons, so points on grid boundaries may intersect with two cells touched; for other grids each cell boundary or corner belongs only to one cell.

**Value**

'sgbp' object if sparse = TRUE, logical matrix otherwise

---

st_join.stars	<i>Spatially join a stars and an 'sf' object</i>
---------------	--

---

**Description**

Spatially join a stars and an 'sf' object

**Usage**

```
## S3 method for class 'stars'
st_join(
  x,
  y,
  join = st_intersects,
  ...,
  what = "left1",
  as_points = NA,
  warn = TRUE
)
```

**Arguments**

x	object of class stars
y	object of class sf, or one that can be coerced into that by <a href="#">st_as_sf</a>
join	the join function, which should return an sgbp object; see details
...	arguments that will be passed on to the join function
what	"left1", "right" or "inner"; see details
as_points	logical; controls whether grid cells in x will be treated as points, or as cell areas; the <a href="#">st_intersects.stars</a> method by default will derive this from x's metadata, or else assume areas.
warn	logical; if TRUE, warn on 1-to-many matches when what is "left1"

**Details**

When there is more than one match to a single x value, the first matching record from y is taken (and if warn is TRUE a warning is raised). If what is "inner", an object of class sf with all matching records of x and y.

**Value**

If what is "left1", an object of class stars with the (first) value of y at spatial instances of x

---

st_mosaic	<i>build mosaic (composite) of several spatially disjoint stars objects</i>
-----------	---

---

**Description**

build mosaic (composite) of several spatially disjoint stars objects

**Usage**

```
st_mosaic(.x, ...)

## S3 method for class 'stars'
st_mosaic(
  .x,
  ...,
  dst = tempfile(fileext = file_ext),
  options = c("-vrtndata", "-9999", "-srcndata", "nan"),
  file_ext = ".tif"
)

## S3 method for class 'character'
st_mosaic(
  .x,
  ...,
```

```

    dst = tempfile(fileext = file_ext),
    options = c("-vrt nodata", "-9999"),
    file_ext = ".tif"
)

## S3 method for class 'stars_proxy'
st_mosaic(
  .x,
  ...,
  dst = tempfile(fileext = file_ext),
  options = c("-vrt nodata", "-9999"),
  file_ext = ".tif"
)

```

### Arguments

.x	object of class stars, or character vector with input dataset names
...	further input stars objects
dst	character; destination file name
options	character; options to the gdalbuildvrt command
file_ext	character; file extension, determining the format used to write to (".tif" implies GeoTIFF)

### Details

the gdal function buildvrt builds a mosaic of input images; these input images can be multi-band, but not higher-dimensional data cubes or stars objects with multiple attributes

uses [gdal\\_utils](#) to internally call buildvrt; no executables external to R are called.

### Value

the stars method returns a stars object with the composite of the input; the character method returns the file name of the file with the mosaic; see also the GDAL documentation of gdalbuildvrt

### Examples

```

x = read_stars(system.file("tif/L7_ETMs.tif", package = "stars"))
x1 = x[,100:200,100:200,]
x2 = x[,150:300,150:300,]
plot(st_mosaic(x1, x2))

```

---

st_rasterize	<i>rasterize simple feature geometries</i>
--------------	--

---

## Description

rasterize simple feature geometries

## Usage

```
st_rasterize(
  sf,
  template = guess_raster(sf, ...) %||% st_as_stars(st_bbox(sf), values = NA_real_,
  ...),
  file = tempfile(),
  driver = "GTiff",
  options = character(0),
  ...
)
```

## Arguments

sf	object of class sf
template	stars object with desired target geometry
file	temporary file name
driver	driver for temporary file
options	character; options vector for GDALRasterize
...	arguments passed on to <a href="#">st_as_stars</a>

## Examples

```
demo(nc, echo = FALSE, ask = FALSE)
(x = st_rasterize(nc)) # default grid:
plot(x, axes = TRUE)
# a bit more customized grid:
(x = st_rasterize(nc, st_as_stars(st_bbox(nc), nx = 100, ny = 50, values = NA_real_)))
plot(x, axes = TRUE)
(ls = st_sf(a = 1:2, st_sfc(st_linestring(rbind(c(0.1, 0), c(1.1, 1))),
  st_linestring(rbind(c(0, 0.05), c(1, 0.05))))))
(grd = st_as_stars(st_bbox(ls), nx = 10, ny = 10, xlim = c(0, 1.0), ylim = c(0, 1),
  values = NA_real_))
# Only the left-top corner is part of the grid cell:
sf_extSoftVersion()["GDAL"]
plot(st_rasterize(ls, grd), axes = TRUE, reset = FALSE) # ALL_TOUCHED=FALSE;
plot(ls, add = TRUE, col = "red")
plot(st_rasterize(ls, grd, options = "ALL_TOUCHED=TRUE"), axes = TRUE, reset = FALSE)
plot(ls, add = TRUE, col = "red")
# add lines to existing 0 values, summing values in case of multiple lines:
```

```
(grd = st_as_stars(st_bbox(ls), nx = 10, ny = 10, xlim = c(0, 1.0), ylim = c(0, 1), values = 0))
r = st_rasterize(ls, grd, options = c("MERGE_ALG=ADD", "ALL_TOUCHED=TRUE"))
plot(r, axes = TRUE, reset = FALSE)
plot(ls, add = TRUE, col = "red")
```

---

st\_raster\_type            *get the raster type (if any) of a stars object*

---

### Description

get the raster type (if any) of a stars object

### Usage

```
st_raster_type(x)
```

### Arguments

x                        object of class stars

### Value

one of NA (if the object does not have raster dimensions), "curvilinear", "rectilinear", "affine", or "regular"

### Examples

```
tif = system.file("tif/L7_ETMs.tif", package = "stars")
x = read_stars(tif)
st_raster_type(x)
```

---

st\_rgb                    *reduce dimension to rgb (alpha) hex values*

---

### Description

reduce dimension to rgb (alpha) hex values

### Usage

```
st_rgb(
  x,
  dimension = 3,
  use_alpha = FALSE,
  maxColorValue = 255L,
  probs = c(0, 1),
  stretch = FALSE
)
```

**Arguments**

x	object of class stars
dimension	dimension name or number to reduce
use_alpha	logical; if TRUE, the fourth band will be used as alpha values
maxColorValue	integer; maximum value for colors
probs	probability values for quantiles used for stretching
stretch	logical; if TRUE, each band is stretched to 0 ... maxColorValue

**Details**

the dimension's bands are mapped to red, green, blue, alpha; if a different ordering is wanted, use [\[.stars\]](#) to reorder a dimension, see examples

**See Also**

[st\\_apply](#), [rgb](#)

**Examples**

```
tif = system.file("tif/L7_ETMs.tif", package = "stars")
x = read_stars(tif)
st_rgb(x, 3)
r = st_rgb(x[, , c(6,5,4,3)], 3, use_alpha=TRUE) # now R=6,G=5,B=4,alpha=3
if (require(ggplot2)) {
  ggplot() + geom_stars(data = r) + scale_fill_identity()
}
```

---

st\_set\_bbox

*set bounding box parameters of regular grid*

---

**Description**

set bounding box parameters of regular grid

**Usage**

```
st_set_bbox(x, value, ...)
```

**Arguments**

x	object of class dimensions, stars or stars_proxy
value	object of class bbox
...	ignored



---

st_sfc2xy	<i>replace POINT simple feature geometry list with an x y raster</i>
-----------	--

---

**Description**

replace POINT simple feature geometry list with an x y raster

**Usage**

```
st_sfc2xy(x, ...)
```

**Arguments**

x	object of class stars, or of class sf
...	passed on to <a href="#">as.data.frame.stars</a>

**Value**

object of class stars with a POINT list replaced by x and y raster dimensions. This only works when the points are distributed over a regular or rectilinear grid.

---

st_transform	<i>transform geometries in stars objects to a new coordinate reference system, without warping</i>
--------------	--

---

**Description**

transform geometries in stars objects to a new coordinate reference system, without warping

**Usage**

```
## S3 method for class 'stars'
st_transform(x, crs, ...)
```

```
## S3 method for class 'stars'
st_transform_proj(x, crs, ...)
```

**Arguments**

x	object of class stars, with either raster or simple feature geometries
crs	object of class crs with target crs
...	ignored

## Details

For simple feature dimensions, `st_transform` is called, leading to lossless transformation. For gridded spatial data, a curvilinear grid with transformed grid cell (centers) is returned, which is also lossless. To convert this to a regular grid in the new CRS, use `st_warp` (which is in general lossy).

## See Also

[st\\_warp](#)

## Examples

```
geomatrix = system.file("tif/geomatrix.tif", package = "stars")
(x = read_stars(geomatrix))
new = st_crs(4326)
y = st_transform(x, new)
plot(st_transform(st_as_sfc(st_bbox(x)), new), col = NA, border = 'red')
plot(st_as_sfc(y, as_points=FALSE), col = NA, border = 'green', axes = TRUE, add = TRUE)
image(y, col = heat.colors(12), add = TRUE)
plot(st_as_sfc(y, as_points=TRUE), pch=3, cex=.5, col = 'blue', add = TRUE)
plot(st_transform(st_as_sfc(x, as_points=FALSE), new), add = TRUE)
```

---

st\_warp

*Warp (resample) grids in stars objects to a new grid, possibly in an new coordinate reference system*

---

## Description

Warp (resample) grids in stars objects to a new grid, possibly in an new coordinate reference system

## Usage

```
st_warp(
  src,
  dest,
  ...,
  crs = NA_crs_,
  cellsize = NA_real_,
  segments = 100,
  use_gdal = FALSE,
  options = character(0),
  no_data_value = NA_real_,
  debug = FALSE,
  method = "near"
)
```

**Arguments**

src	object of class stars with source raster
dest	object of class stars with target raster geometry
...	ignored
crs	coordinate reference system for destination grid, only used when dest is missing
cellsize	length 1 or 2 numeric; cellsize in target coordinate reference system units
segments	(total) number of segments for segmentizing the bounding box before transforming to the new crs
use_gdal	logical; if TRUE, use gdalwarp, through <a href="#">gdal_utils</a>
options	character vector with options, passed on to gdalwarp
no_data_value	value used by gdalwarp for no_data (NA) when writing to temporary file
debug	logical; if TRUE, do not remove the temporary gdalwarp destination file, and print its name
method	character; see details for options; methods other than near only work when use_gdal=TRUE

**Details**

method should be one of near, bilinear, cubic, cubicspline, lanczos, average, mode, max, min, med, q1 or q3; see <https://github.com/r-spatial/stars/issues/109>

For gridded spatial data (dimensions  $x$  and  $y$ ), see figure; the existing grid is transformed into a regular grid defined by dest, possibly in a new coordinate reference system. If dest is not specified, but crs is, the procedure used to choose a target grid is similar to that of [projectRaster](#) (currently only with method='ngb'). This entails: (i) the envelope (bounding box polygon) is transformed into the new crs, possibly after segmentation (red box); (ii) a grid is formed in this new crs, touching the transformed envelope on its East and North side, with (if cellsize is not given) a cellsize similar to the cell size of src, with an extent that at least covers  $x$ ; (iii) for each cell center of this new grid, the matching grid cell of  $x$  is used; if there is no match, an NA value is used.

**Examples**

```
geomatrix = system.file("tif/geomatrix.tif", package = "stars")
(x = read_stars(geomatrix))
new_crs = st_crs(4326)
y = st_warp(x, crs = new_crs)
plot(st_transform(st_as_sfc(st_bbox(x)), new_crs), col = NA, border = 'red')
plot(st_as_sfc(y, as_points=FALSE), col = NA, border = 'green', axes = TRUE, add = TRUE)
image(y, add = TRUE, nbreaks = 6)
plot(st_as_sfc(y, as_points=TRUE), pch=3, cex=.5, col = 'blue', add = TRUE)
plot(st_transform(st_as_sfc(x, as_points=FALSE), new_crs), add = TRUE)
# warp 0-360 raster to -180-180 raster:
r = read_stars(system.file("nc/reduced.nc", package = "stars"))
r %>% st_set_crs(4326) %>% st_warp(st_as_stars(st_bbox(), dx = 2)) -> s
plot(r, axes = TRUE) # no CRS set, so no degree symbols in labels
plot(s, axes = TRUE)
```

---

<code>st_xy2sfc</code>	<i>replace x y raster dimensions with simple feature geometry list (points, or polygons = rasterize)</i>
------------------------	--

---

**Description**

replace x y raster dimensions with simple feature geometry list (points, or polygons = rasterize)

**Usage**

```
st_xy2sfc(x, as_points, ..., na.rm = TRUE)
```

**Arguments**

<code>x</code>	object of class <code>stars</code>
<code>as_points</code>	logical; if TRUE, generate points at cell centers, else generate polygons
<code>...</code>	arguments passed on to <code>st_as_sfc</code>
<code>na.rm</code>	logical; omit (remove) cells which are entirely missing valued (across other dimensions)?

**Value**

object of class `stars` with x and y raster dimensions replaced by a single `sfc` geometry list column containing either points, or polygons. Adjacent cells with identical values are not merged; see `st_rasterize` for this.

---

<code>write_stars</code>	<i>write stars object to gdal dataset (typically: to file)</i>
--------------------------	--

---

**Description**

write stars object to gdal dataset (typically: to file)

**Usage**

```
write_stars(obj, dsn, layer, ...)
```

```
## S3 method for class 'stars'
write_stars(
  obj,
  dsn,
  layer = 1,
  ...,
  driver = detect.driver(dsn),
  options = character(0),
```

```

    type = "Float32",
    NA_value = NA_real_,
    update = FALSE,
    normalize_path = TRUE
  )

## S3 method for class 'stars_proxy'
write_stars(
  obj,
  dsn,
  layer = 1,
  ...,
  driver = detect.driver(dsn),
  options = character(0),
  type = "Float32",
  NA_value = NA_real_,
  chunk_size = c(dim(obj)[1], floor(2.5e+07/dim(obj)[1])),
  progress = TRUE
)

detect.driver(filename)

```

### Arguments

obj	object of class stars
dsn	gdal dataset (file) name
layer	attribute name; if missing, the first attribute is written
...	passed on to <a href="#">gdal_write</a>
driver	driver driver name; see <a href="#">st_drivers</a>
options	character vector with options
type	character; output binary type, one of: Byte for eight bit unsigned integer, UInt16 for sixteen bit unsigned integer, Int16 for sixteen bit signed integer, UInt32 for thirty two bit unsigned integer, Int32 for thirty two bit signed integer, Float32 for thirty two bit floating point, Float64 for sixty four bit floating point.
NA_value	non-NA value that should represent R's NA value in the target raster file; if set to NA, it will be ignored.
update	logical; if TRUE, an existing file is being updated
normalize_path	logical; see <a href="#">read_stars</a>
chunk_size	length two integer vector with the number of pixels (x, y) used in the read/write loop; see details.
progress	logical; if TRUE, a progress bar is shown
filename	character; used for guessing driver short name based on file extension; see examples

**Details**

`write_stars` first creates the target file, then updates it sequentially by writing blocks of `chunk_size`.

**Examples**

```
detect.driver("L7_ETMs.tif")
```

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