

# Package: stars (via r-universe)

January 19, 2025

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

**Version** 0.6-8

**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** <https://cran.uni-muenster.de/pebesma/>

**LazyData** true

**Depends** R (>= 3.3.0), abind, sf (>= 1.0-19)

**Imports** methods, parallel, classInt (>= 0.4-1), rlang, units

**Suggests** Cairo, OpenStreetMap, PCICt, RNetCDF (>= 1.8-2), clue, covr, cubble (>= 0.3.0), cubelyr, digest, dplyr (>= 0.7-0), exactextractr, FNN, future.apply, ggforce, ggplot2, ggthemes, gstat, httr, jsonlite, knitr, lwgeom, maps, mapdata, ncdfgeom, ncmeta (>= 0.0.3), pbapply, plm, randomForest, raster, rmarkdown, sp, spacetime, spatstat (>= 2.0-1), spatstat.geom, starsdata, terra (>= 1.4-22), testthat, tibble, tidyr, tsibble, viridis, xts, zoo

**VignetteBuilder** knitr

**Encoding** UTF-8

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**Collate** 'aggregate.R' 'cubble.R' 'datasets.R' 'dimensions.R'  
'extract.R' 'factors.R' 'geom.R' 'gt.R' 'init.R' 'intervals.R'  
'mdim.R' 'mosaic.R' 'ncdf.R' 'ncproxy.R' 'OpenStreetMap.R'  
'ops.R' 'plot.R' 'prcomp.R' 'proxy.R' 'rasterize.R' 'raster.R'  
'read.R' 'rotate.R' 'sample.R' 'sf.R' 'spacetime.R'  
'spatstat.R' 'sp.R' 'stars.R' 'subset.R' 'tidyverse.R' 'tile.R'  
'transform.R' 'values.R' 'warp.R' 'write.R' 'xts.R'

**Config/pak/sysreqs** libgdal-dev gdal-bin libgeos-dev libproj-dev  
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**Repository** <https://fastverse.r-universe.dev>

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

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aggregate.stars	<i>spatially or temporally aggregate stars object</i>
-----------------	---

---

**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> ), or a function that cuts time into intervals; if by is

an object of class `stars`, it is converted to `sfc` by `st_as_sfc`(`by`, `as_points = FALSE`) thus ignoring its time component. Note: each pixel is assigned to only a single group (in the order the groups occur) so non-overlapping spatial features and temporal windows are recommended.

`FUN` aggregation function, such as `mean`  
`...` arguments passed on to `FUN`, such as `na.rm=TRUE`  
`drop` logical; ignored  
`join` function; function used to find matches of `x` to `by`  
`as_points` see `st_as_sf`: shall raster pixels be taken as points, or small square polygons?  
`rightmost.closed`  
 see `findInterval`  
`left.open` logical; used for time intervals, see `findInterval` and `cut.POSIXt`  
`exact` logical; if `TRUE`, use `coverage_fraction` to compute exact overlap fractions of polygons with raster cells

### See Also

[aggregate](#), [st\\_interpolate\\_aw](#), [st\\_extract](#), <https://github.com/r-spatial/stars/issues/317>

### 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")
#TBD:
#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")
#TBD:
# x_agg_time - x_agg_posix
aggregate(x, "2 days", mean)
if (require(ncmeta, quietly = TRUE)) {
  # Spatial aggregation, see https://github.com/r-spatial/stars/issues/299
```

```

prec_file = system.file("nc/test_stageiv_xyt.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)
}

# example of using a function for "by": aggregate by month-of-year
d = c(10, 10, 150)
a = array(rnorm(prod(d)), d) # pure noise
times = Sys.Date() + seq(1, 2000, length.out = d[3])
m = as.numeric(format(times, "%m"))
signal = rep(sin(m / 12 * pi), each = prod(d[1:2])) # yearly period
s = (st_as_stars(a) + signal) %>%
  st_set_dimensions(3, values = times)
f = function(x, format = "%B") {
  months = format(as.Date(paste0("01-", 1:12, "-1970")), format)
  factor(format(x, format), levels = months)
}
agg = aggregate(s, f, mean)
plot(agg)

```

as

*Coerce stars object into a Raster raster or brick***Description**

Coerce stars object into a Raster raster or brick

Coerce stars object into a terra SpatRaster

**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.

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 SpatRaster.

**Value**

RasterLayer or RasterBrick

SpatRaster

---

bcsd_obs	<i>Monthly Gridded Meteorological Observations</i>
----------	--

---

### Description

These are the monthly observational data used for BCSD downscaling. See: <[https://gdo-dcp.ucllnl.org/downscaled\\_cmip\\_pr](https://gdo-dcp.ucllnl.org/downscaled_cmip_pr) for more information." ; "Atmospheric Temperature, Air Temperature Atmosphere, Precipitation, Rain, Maximum Daily Temperature, Minimum Daily Temperature" ;

### Usage

```
bcsd_obs
```

### Format

An object of class stars\_proxy (inherits from stars) of dimension 81 x 33 x 12.

---

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_proxy'
c(
  ...,
  along = NA_integer_,
  along_crs = FALSE,
  try_hard = FALSE,
  nms = names(list(...)),
  tolerance = sqrt(.Machine$double.eps)
)

## S3 method for class 'stars'
c(
  ...,
  along = NA_integer_,
  try_hard = FALSE,
  nms = names(list(...)),
  tolerance = sqrt(.Machine$double.eps)
)
```

**Arguments**

...	object(s) of class <code>star</code> : in case of multiple arguments, these are combined into a single <code>stars</code> 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.
<code>along</code>	integer; see <a href="#">read_stars</a>
<code>along_crs</code>	logical; if TRUE, combine arrays along a CRS dimension
<code>try_hard</code>	logical; if TRUE and some arrays have different dimensions, combine those that dimensions matching to the first array
<code>nms</code>	character; vector with array names
<code>tolerance</code>	numeric; values used in <a href="#">all.equal</a> to compare dimension values combine those that dimensions matching to the first array

**Details**

An error is raised when attempting to combine arrays with different measurement units into a single array. If this was intended, `drop_units` can be used to remove units of a `stars` object before merging.

**Value**

a single `stars` object with merged (binded) arrays.

**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)
```

---

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

---

**Description**

plot contours of a `stars` object

**Usage**

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

**Arguments**

<code>x</code>	object of class <code>stars</code>
...	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

*cut methods for stars objects*


---

**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 <a href="#">cut</a>
breaks	see <a href="#">cut</a>
...	see <a href="#">cut</a>

**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; package dplyr needs to be loaded before these methods can be used 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, ...)
rename.stars(.data, ...)
rename.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, along, index, ...)
```

```
replace_na.stars(data, replace, ...)
```

```
replace_na.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?
data	data set to work on
replace	see <a href="#">replace_na</a> : list with variable=value pairs, where value is the replacement value for NA's

### Examples

```
tif = system.file("tif/L7_ETMs.tif", package = "stars")
x1 = read_stars(tif)
if (require(dplyr, quietly = TRUE)) {
  x1 %>% slice("band", 2:3)
  x1 %>% slice("x", 50:100)
}
```

---

expand\_dimensions      *expand the dimension values into a list*

---

### Description

expand the dimension values into a list

### Usage

```
expand_dimensions(x, ...)
```

```
## S3 method for class 'dimensions'
```

```
expand_dimensions(x, ..., max = FALSE, center = NA)
```

**Arguments**

x	object of class 'stars' or 'dimensions'
...	ignored
max	logical; if 'TRUE' return the max (end) values of the dimensions intervals
center	logical; if 'TRUE' return the center values of intervals, otherwise return offset (start) of intervals; if 'NA' (default) return centers for x/y dimensions, offsets for all others

---

geom_stars	<i>ggplot geom for stars objects</i>
------------	--------------------------------------

---

**Description**

ggplot geom for stars objects

**Usage**

```
geom_stars(
  mapping = NULL,
  data = NULL,
  ...,
  downsample = 0,
  sf = FALSE,
  na.action = na.pass
)

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; can be specified for each dimension, e.g. <code>c(5,5,0)</code> to downsample the first two dimensions but not the third.
sf	logical; if TRUE rasters will be converted to polygons and plotted using <a href="#">geom_sf</a> .
na.action	function; if NA values need to be removed before plotting use the value <code>na.omit</code> here (only applies to objects with raster dimensions)

## 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. If a stars array contains hex color values, and no `fill` parameter is given, the color values are used as fill color; see the example below.

If visual artefacts occur (Moiré-Effekt), then see the details section of [plot.stars](#)

## Examples

```
system.file("tif/L7_ETMs.tif", package = "stars") %>% read_stars() -> x
if (require(ggplot2, quietly = TRUE)) {
  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))
# plot rgb composite:
st_as_stars(L7_ETMs)[,,1:3] |> st_rgb() -> x # x contains colors as pixel values
ggplot() + geom_stars(data = x)
}
```

---

L7\_ETMs

*Landsat-7 bands for a selected region around Olinda, BR*

---

## Description

Probably containing the six 30 m bands:

- Band 1 Visible (0.45 - 0.52  $\mu\text{m}$ ) 30 m
- Band 2 Visible (0.52 - 0.60  $\mu\text{m}$ ) 30 m
- Band 3 Visible (0.63 - 0.69  $\mu\text{m}$ ) 30 m
- Band 4 Near-Infrared (0.77 - 0.90  $\mu\text{m}$ ) 30 m
- Band 5 Short-wave Infrared (1.55 - 1.75  $\mu\text{m}$ ) 30 m
- Band 7 Mid-Infrared (2.08 - 2.35  $\mu\text{m}$ ) 30 m

## Usage

L7\_ETMs

## Format

An object of class `stars_proxy` (inherits from `stars`) of dimension 349 x 352 x 6.

---

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

---

**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

---

mdim	<i>Read or write data using GDAL's multidimensional array API</i>
------	---

---

**Description**

Read or write data using GDAL's multidimensional array API

**Usage**

```
read_mdim(
  filename,
  variable = character(0),
  ...,
  options = character(0),
  raster = NULL,
  offset = integer(0),
  count = integer(0),
  step = integer(0),
  proxy = FALSE,
  debug = FALSE,
  bounds = TRUE,
  curvilinear = NA
)

write_mdim(
  x,
  filename,
  driver = detect.driver(filename),
```

```

...,
root_group_options = character(0),
options = character(0),
as_float = TRUE
)

```

### Arguments

filename	name of the source or destination file or data source
variable	name of the array to be read; if "", a list of array names is returned, with group name as list element names.
...	ignored
options	character; driver specific options regarding the opening (read_mdime) or creation (write_mdime) of the dataset
raster	names of the raster variables (default: first two dimensions)
offset	integer; zero-based offset for each dimension (pixels) of sub-array to read, defaults to 0 for each dimension (requires sf >= 1.0-9)
count	integer; size for each dimension (pixels) of sub-array to read (default: read all); a value of NA will read the corresponding dimension entirely; counts are relative to the step size (requires sf >= 1.0-9)
step	integer; step size for each dimension (pixels) of sub-array to read; defaults to 1 for each dimension (requires sf >= 1.0-9)
proxy	logical; return proxy object?
debug	logical; print debug info?
bounds	logical or character: if TRUE tries to infer from "bounds" attribute; if character, named vector of the form c(longitude="lon_bnds", latitude="lat_bnds") with names dimension names
curvilinear	control reading curvilinear (geolocation) coordinate arrays; if NA try reading the x/y dimension names; if character, defines the arrays to read; if FALSE do not try; see also <a href="#">read_stars</a>
x	stars object
driver	character; driver name
root_group_options	character; driver specific options regarding the creation of the root group
as_float	logical; if TRUE write 4-byte floating point numbers, if FALSE write 8-byte doubles

### Details

it is assumed that the first two dimensions are easting and northing

### See Also

[gdal\\_utils](#), in particular util mdiminfo to query properties of a file or data source containing arrays

**Examples**

```

set.seed(135)
m = matrix(runif(10), 2, 5)
names(dim(m)) = c("stations", "time")
times = as.Date("2022-05-01") + 1:5
pts = st_as_sfc(c("POINT(0 1)", "POINT(3 5)"))
s = st_as_stars(list(Precipitation = m)) |>
  st_set_dimensions(1, values = pts) |>
  st_set_dimensions(2, values = times)
nc = tempfile(fileext=".nc")
if (compareVersion(sf_extSoftVersion()["GDAL"], "3.4.0") > -1) {
  write_mdims(s, nc)
  # try ncdump on the generated file
  print(read_mdims(nc))
}

```

---

merge

*merge or split stars object*


---

**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

*S3 Ops Group Generic Functions for stars objects*


---

**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

**Details**

if e1 or e2 is a numeric vector, or e2 has less or smaller dimensions than e1, then e2 is recycled such that it fits e1, using usual R array recycling rules. The user needs to make sure this is sensible; it may be needed to use aperm to permutate dimensions first.

**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 'nc_proxy'
plot(x, y, ..., downsample = get_downsample(dim(x)), max_times = 16)

## 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 = kw_dflt(x, key.pos),
  key.length = 0.618,
  key.lab = ifelse(length(main) == 1, main, ""),
  reset = TRUE,
  box_col = NA,
  center_time = FALSE,
  hook = NULL,
```

```

    mfrow = NULL,
    compact = TRUE
)

## 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 = kw_dflt(x, key.pos),
  key.length = 0.618,
  add.geom = NULL,
  border = NA,
  useRaster = isTRUE(dev.capabilities()$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>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 number of pixels/lines/bands etc that will be skipped; see <code>Details</code> .
<code>max_times</code>	integer; maximum number of time steps to attempt to plot.
<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>

main	character; subplot title prefix; use "" to get only time, use NULL to suppress subplot titles
axes	logical; should axes and box be added to the plot?
nbreaks	number of color breaks; should be one more than number of colors. If missing and col is specified, it is derived from that.
breaks	numeric vector with actual color breaks, or a style name used in <a href="#">classIntervals</a> .
col	colors to use for grid cells, or color palette function
key.pos	numeric; side to plot a color key: 1 bottom, 2 left, 3 top, 4 right; set to NULL 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 asp. If it has length 2, the second value, ranging from 0 to 1, determines where the key is placed in the available space (default: 0.5, center).
key.width	amount of space reserved for width of the key (labels); relative or absolute (using lcm)
key.length	amount of space reserved for length of the key (labels); relative or absolute (using lcm)
key.lab	character; label for color key in case of multiple subplots, use "" to suppress
reset	logical; if FALSE, keep the plot in a mode that allows adding further map elements; if TRUE restore original mode after plotting
box_col	color for box around sub-plots; use NA to suppress plotting of boxes around sub-plots.
center_time	logical; if TRUE, sub-plot titles will show the center of time intervals, otherwise their start
hook	NULL or function; hook function that will be called on every sub-plot; see examples.
mfrow	length-2 integer vector with nrows, ncolumns of a composite plot, to override the default layout
compact	logical; place facets compactly (TRUE), or spread over the plotting device area?
band	integer; which band (dimension) to plot
attr	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 <a href="#">Details</a> .
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 <code>rasterImage</code> (rgb), should pixels be interpolated?
as_points	logical; for curvilinear or sheared grids: parameter passed on to <code>st_as_sf</code> , 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 <code>x</code> 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

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

Downsampling: a value for `downsample` of 0: no downsampling, 1: after every dimension value (pixel/line/band), one value is skipped (half of the original resolution), 2: after every dimension value, 2 values are skipped (one third of the original resolution), etc. If `downsample` is TRUE or a length 1 numeric vector, downsampling is only applied to the raster [x] and [y] dimensions.

To remove unused classes in a categorical raster, use the `droplevels` function.

When bitmaps show visual artefacts (Moiré effects), make sure that device `png` is used rather than `ragg::agg_png` as the latter uses antialiasing for filled polygons which causes this; see also <https://github.com/r-spatial/stars/issues/573>.

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 has to be set manually.

### Examples

```
st_bbox(L7_ETMs) |> st_as_sfc() |> st_centroid() |> st_coordinates() -> pt
hook1 = function() {
  text(pt[,"X"], pt[,"Y"], "foo", col = 'orange', cex = 2)
}
plot(L7_ETMs, hook = hook1)
x = st_set_dimensions(L7_ETMs, 3, paste0("B_", 1:6))
hook2 = function(..., row, col, nr, nrow, ncol, value, bbox) {
  str = paste0("row ", row, "/", nrow, ", col ", col, "/", ncol, "\nnr: ", nr, " value: ", value)
  bbox |> st_as_sfc() |> st_centroid() |> st_coordinates() -> pt
  text(pt[,"X"], pt[,"Y"], str, col = 'red', cex = 2)
}
plot(x, hook = hook2, col = grey(c(.2, .25, .3, .35)))
if (isTRUE(dev.capabilities()$rasterImage == "yes")) {
  lc = read_stars(system.file("tif/lc.tif", package = "stars"))
  levels(lc[[1]]) = abbreviate(levels(lc[[1]]), 6) # so it's not only legend
  plot(lc, key.pos=4)
}
```

```
tif = system.file("tif/L7_ETMs.tif", package = "stars")
x = read_stars(tif)
image(x, col = grey((3:9)/10))
if (isTRUE(dev.capabilities()$rasterImage == "yes")) {
  image(x, rgb = c(1,3,5)) # false color composite
}
```

prcomp

*Principle components of stars object***Description**

Compute principle components of stars object

**Usage**

```
## S3 method for class 'stars_proxy'
prcomp(x, ..., downsample = 0)

## S3 method for class 'stars'
prcomp(x, ..., quiet = FALSE)
```

**Arguments**

x	object of class 'stars' or 'stars_proxy'
...	see <a href="#">prcomp</a>
downsample	see <a href="#">st_as_stars</a>
quiet	logical; if 'TRUE', suppress message that PCs will be computed on last dimension; see details

**Details**

if 'x' has only one attribute, principle components will be computed in the space of the last dimension of 'x' to predict PC scores into a 'stars' object, use [predict.stars](#); see example below

**Value**

object of class 'prcomp', see [prcomp](#)

**Examples**

```
l7 = split(st_as_stars(L7_ETMs), 3) # use bands as features
l7 |> prcomp() |> plot()
l7 |> prcomp() |> predict(l7, model = _) |> merge() |> plot()
```

---

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_proxy'
predict(object, model, ...)

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

**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
drop_dimensions	logical; if 'TRUE', remove dimensions (coordinates etc) from 'data.frame' with predictors

**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)'

---

print_stars	<i>print stars or dimensions object</i>
-------------	---

---

**Description**

print stars or dimensions object

**Usage**

```
## S3 method for class 'dimensions'
as.data.frame(
  x,
  ...,
  digits = max(3, getOption("digits") - 3),
  usetz = TRUE,
```

```

    stars_crs = getOption("stars.crs") %||% 28,
    all = FALSE
  )

  ## S3 method for class 'dimensions'
  print(x, ...)

  ## S3 method for class 'stars'
  print(x, ..., n = 1e+05, abbrev = 30)

```

### Arguments

x	object of class stars or of class dimensions
...	passed on to <code>as.data.frame.dimensions</code>
digits	number of digits to print numbers
usetz	logical; used to format PCICt or POSIXct values
stars_crs	maximum width of string for CRS objects
all	logical; if TRUE print also fields entirely filled with NA or NULL
n	when $\text{prod}(\text{dim}(x)) > 10 * n$ , the first n cells are used for attribute summary statistics
abbrev	number of characters to abbreviate attribute names to

---

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,
  ncsub = NULL,
  curvilinear = character(0),
  eps = sqrt(.Machine$double.eps),
  ignore_bounds = FALSE,
  make_time = TRUE,
  make_units = TRUE,
  proxy = NULL,
  downsample = 0
)

```

**Arguments**

<code>.x</code>	NetCDF file or source as a character vector or an <code>nc_proxy</code> object.
<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 <code>eps</code>
<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
<code>proxy</code>	logical; if TRUE, an object of class <code>stars_proxy</code> is read which contains array metadata only; if FALSE the full array data is read in memory. If not set, defaults to TRUE when the number of cells to be read is larger than <code>options(stars.n_proxy)</code> , or to 1e8 if that option was not set.
<code>downsample</code>	integer; number of cells to omit between samples along each dimension. e.g. <code>c(1,1,2)</code> would return every other cell in x and y and every third cell in the third dimension (z or t). If 0, no downsampling is applied. Note that this transformation is applied AFTER NetCDF data are read using <code>st_downsample</code> . As such, if <code>proxy=TRUE</code> , this option is ignored.

**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")
if (require(ncmeta, quietly = TRUE)) {
  read_ncdf(f)
}
```



```

read_ncdf(f, var = c("anom"))
read_ncdf(f, ncs = cbind(start = c(1, 1, 1, 1), count = c(10, 12, 1, 1)))
}

if (require(ncmeta, quietly = TRUE)) {
  #' 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)))
}
if (require(dplyr, quietly = TRUE)) {
  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

*read raster/array dataset from file or connection*


---

## Description

read raster/array dataset from file or connection

## Usage

```

read_stars(
  .x,
  sub = TRUE,
  ...,
  options = character(0),
  driver = character(0),
  quiet = FALSE,
  NA_value = NA_real_,
  along = NA_integer_,
  RasterIO = list(),
  proxy = getOption("stars.n_proxy") %||% 1e+08,
  curvilinear = character(0),
  normalize_path = TRUE,
  RAT = character(0),
  tolerance = 1e-10,
  exclude = "",

```

```

    shorten = TRUE
  )

```

### Arguments

<code>.x</code>	character vector with name(s) of file(s) or data source(s) to be read, or a function that returns such a vector
<code>sub</code>	character, integer or logical; name, index or indicator of sub-dataset(s) to be read
<code>...</code>	passed on to <code>st_as_stars</code> if <code>curvilinear</code> was set
<code>options</code>	character; opening options
<code>driver</code>	character; driver to use for opening file. To override fixing for subdatasets and autodetect them as well, use <code>NULL</code> .
<code>quiet</code>	logical; print progress output?
<code>NA_value</code>	numeric value to be used for conversion into NA values; by default this is read from the input file
<code>along</code>	length-one character or integer, or list; determines how several arrays are combined, see <code>Details</code> .
<code>RasterIO</code>	list with named parameters for GDAL's <code>RasterIO</code> , to further control the extent, resolution and bands to be read from the data source; see <code>details</code> .
<code>proxy</code>	logical; if <code>TRUE</code> , an object of class <code>stars_proxy</code> is read which contains array metadata only; if <code>FALSE</code> the full array data is read in memory. Always <code>FALSE</code> for <code>curvilinear</code> girds. If set to a number, defaults to <code>TRUE</code> when the number of cells to be read is larger than that number.
<code>curvilinear</code>	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
<code>normalize_path</code>	logical; if <code>FALSE</code> , suppress a call to <code>normalizePath</code> on <code>.x</code>
<code>RAT</code>	character; raster attribute table column name to use as factor levels
<code>tolerance</code>	numeric; passed on to <code>all.equal</code> for comparing dimension parameters.
<code>exclude</code>	character; vector with category value(s) to exclude
<code>shorten</code>	logical or character; if <code>TRUE</code> and <code>length(.x) &gt; 1</code> , remove common start and end parts of array names; if character a new prefix

### 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, resample. See <https://gdal.org/en/latest/doxygen/classGDALDataset.html> 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".

Data that are read into memory (proxy=FALSE) are read into a numeric (double) array, except for categorical variables which are read into an numeric (integer) array of class factor.

## 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	<i>redimension array, or collapse attributes into a new dimension</i>
-------------	---

---

**Description**

redimension array, or collapse attributes into a new dimension

**Usage**

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

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

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

**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
name	character name of the new dimension

---

stars_sentinel2	<i>Sentinel-2 sample tile</i>
-----------------	-------------------------------

---

**Description**

Sentinel-2 sample tile, downloaded from <<https://scihub.copernicus.eu/>> reads the four 10-m bands: B2 (490 nm), B3 (560 nm), B4 (665 nm) and B8 (842 nm)

**Usage**

```
stars_sentinel2
```

**Format**

An object of class `stars_proxy` (inherits from `stars`) of dimension 10980 x 10980 x 4.

---

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

---

**Description**

subset stars objects

**Usage**

```
## S3 replacement method for class 'stars_proxy'
x[i, downsample = 0] <- value

## 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**

<code>x</code>	object of class <code>stars</code>
<code>i</code>	first selector: integer, logical or character vector indicating attributes to select, or object of class <code>sf</code> , <code>sfc</code> , <code>bbox</code> , or <code>stars</code> used as spatial selector; see details
<code>downsample</code>	downsampling rate used in case <code>i</code> is a <code>stars_proxy</code> object
<code>value</code>	array of dimensions equal to those in <code>x</code> , or a vector or value that will be recycled to such an array
<code>...</code>	further (logical or integer vector) selectors, matched by order, to select on individual dimensions
<code>drop</code>	logical; if <code>TRUE</code> , degenerate dimensions (with only one value) are dropped
<code>crop</code>	logical; if <code>TRUE</code> and parameter <code>i</code> is a spatial geometry ( <code>sf</code> or <code>sfc</code> ) object, the extent (bounding box) of the result is cropped to match the extent of <code>i</code> using <a href="#">st_crop</a> . Cropping curvilinear grids is not supported.
<code>which</code>	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. If *i* is of class *stars*, and attributes of *i* are logical, cells in *x* corresponding to NA or FALSE cells in *i* are assigned an NA. Dimension ranges containing negative values or NA may be partially supported.

in an assignment (or replacement form, [*<-*], argument *i* needs to be either (i) a *stars* object with logical attribute(s) that has dimensions matching (possibly after recycling) those of *x*, in which case the TRUE cells will be replaced and *i* and/or value will be recycled to the dimensions of the arrays in *x*, or (ii) a length-one integer or character vector indicating which array to replace, in which case value may be *stars* object or a vector or array (that will be recycled).

## 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)
# with i of class stars:
x[x > 75] # generates lots of NA's; pattern for each band
x[x[,,,1] > 75] # recycles a single band template for all bands
x = read_stars(tif)
# replace, using a logical stars selector: cuts all values above 90 to 90
x[x > 90] = 90
# replace a single attribute when there are more than one:
```

```

s = split(x)
names(s) = paste0("band", 1:6)
# rescale only band 1:
s[1] = s[1] * 0.75
# rescale only attribute named "band2":
s["band2"] = s["band2"] * 0.85
# create a new attribute from a numeric vector:
s["rnorm"] = rnorm(prod(dim(s)))
s
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,
  single_arg = has_single_arg(FUN, list(...)) || can_single_arg(FUN),
  keep = FALSE
)

```

## 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> and see Details.
...	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
single_arg	logical; if TRUE, FUN takes a single argument (like fn_ndvi1 below), if FALSE FUN takes multiple arguments (like fn_ndvi2 below).
keep	logical; if TRUE, preserve dimension metadata (e.g. time stamps)

### Details

FUN is a function which either operates on a single object, which will be the data of each iteration step over dimensions MARGIN, or a function that has as many arguments as there are elements in such an object. See the NDVI examples below. The second form can be VERY much faster e.g. when a trivial function is not being called for every pixel, but only once (example).

The heuristics for the default of single\_arg work often, but not always; try setting this to the right value when st\_apply gives an error.

### 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. Following the logic of [apply](#), This new dimension is put before the other dimensions; use [aperm](#) to rearrange this, see last example.

### 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
## Not run:
st_apply(x, "band", mean) # equivalent to the above
st_apply(x, 1:2, range) # min and max band value for each pixel
fn_ndvi1 = function(x) (x[4]-x[3])/(x[4]+x[3]) # ONE argument: will be called for each pixel
fn_ndvi2 = function(red,nir) (nir-red)/(nir+red) # n arguments: will be called only once
ndvi1 = st_apply(x, 1:2, fn_ndvi1)
# note that we can select bands 3 and 4 in the first argument:
ndvi2 = st_apply(x[, , 3:4], 1:2, fn_ndvi2)
all.equal(ndvi1, ndvi2)
# compute the (spatial) variance of each band; https://github.com/r-spatial/stars/issues/430
st_apply(x, 3, function(x) var(as.vector(x))) # as.vector is required!
# to get a progress bar also in non-interactive mode, specify:
if (require(pbapply)) { # install it, if FALSE
  pboptions(type = "timer")
}
```



```

}
st_apply(x, 1:2, range) # dimension "range" is first; rearrange by:
st_apply(x, 1:2, range) %>% aperm(c(2,3,1))

## End(Not run)

```

---

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
)

## S3 method for class 'stars_proxy'
st_as_sf(x, ..., downsample = 0)

```

### 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.
downsample	see <a href="#">st_as_stars</a>

### 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

```
## S3 method for class 'cubple_df'
st_as_stars(.x, ..., check_times = FALSE)

## S3 method for class 'ncdfgeom'
st_as_stars(.x, ..., sf_geometry = NA)

## S3 method for class 'OpenStreetMap'
st_as_stars(.x, ..., as_col = FALSE)
```

```
## 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 = coords, xy, y_decreasing = TRUE, coords = 1:2)

## S3 method for class 'Raster'
st_as_stars(.x, ..., att = 1, ignore_file = FALSE)

## S3 method for class 'SpatRaster'
st_as_stars(
  .x,
  ...,
  ignore_file = FALSE,
  as_attributes = all(terra::is.factor(.x))
)

## S3 method for class 'sf'
st_as_stars(.x, ..., dims = attr(.x, "sf_column"))

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("OGC:CRS84"))

## 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,
    proxy = FALSE
)

## S3 method for class 'xts'
st_as_stars(.x, ..., dimensions, name = "attr")

```

### 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> . In case <code>.x</code> is of class <code>nc_proxy</code> , arguments passed on to <a href="#">read_ncdf</a> .
<code>check_times</code>	logical; should we check that the time stamps of all time series are identical?
<code>sf_geometry</code>	sf data.frame with geometry and attributes to be added to stars object. Must have same number of rows as timeseries instances.
<code>as_col</code>	logical; return rgb numbers (FALSE) or (character) color values (TRUE)?
<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, see details
<code>y_decreasing</code>	logical; if TRUE, (numeric) y values get a negative delta (decrease with increasing index)
<code>coords</code>	same as <code>dims</code> , for symmetry with <a href="#">st_as_sf</a>
<code>att</code>	see <a href="#">factorValues</a> ; column in the RasterLayer's attribute table
<code>ignore_file</code>	logical; if TRUE, ignore the SpatRaster object file name
<code>as_attributes</code>	logical; if TRUE and <code>.x</code> has more than one layer, load these as separate attributes rather than as a band or time dimension (only implemented for the case where <code>ignore_file</code> is TRUE)
<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 or stars arrays, or the names of the corresponding attributes in <code>.x</code> ; the names of this vector should correspond to raster dimensions the matrices are associated with; see <a href="#">Details</a> .
<code>crs</code>	object of class <code>crs</code> with the coordinate reference system of the values in <code>curvilinear</code> ; see details

nx	integer; number of cells in x direction; see details
ny	integer; number of cells in y direction; see details
dx	numeric or object of class units; cell size in x direction; see details
dy	numeric or object of class units; cell size in y direction; see details
xlim	length 2 numeric vector with extent (min, max) in x direction
ylim	length 2 numeric vector with extent (min, max) in y direction
values	value(s) to populate the raster values with
n	the (approximate) target number of grid cells
pretty	logical; should cell coordinates have <a href="#">pretty</a> values?
inside	logical; should all cells entirely fall inside the bbox, potentially not covering it completely (TRUE), or always cover the bbox (FALSE), or find a good approximation (NA, default)?
nz	integer; number of cells in z direction; if missing no z-dimension is created.
proxy	logical; should a stars_proxy object be created? (requires gdal_create binary when sf < 1.0-6)
name	character; attribute name for array from an xts object

## Details

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.

If `xy` is not specified and the first two dimensions in `dims` are both numeric, then it is set to these two dimensions.

The `st_as_stars` method for `sf` objects without any additional arguments returns a one-dimensional data cube with a dimension for the simple features geometries, and all remaining attributes as data cube attributes. When used with further arguments, the method for `data.frames` is called.

if `curvilinear` is a list with `stars` objects with longitude and latitude values, its coordinate reference system is typically not that of the latitude and longitude values. If `curvilinear` contains the names of two arrays in `.x`, then these are removed from the returned object.

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  $\sqrt{\text{diff}(xlim) * \text{diff}(ylim) / n}$ .

If `nx` and `ny` are missing and `values` is a matrix, the number of columns and rows of the matrix are taken.

Otherwise, if `nx` and `ny` are missing, they are computed as the (ceiling, floor, or rounded to integer value) of the ratio of the (x or y) range divided by (dx or dy), depending on the value of `inside`. Positive `dy` will be made negative. Further named arguments (...) are passed on to `pretty`. If `dx` or `dy` are units objects, their value is converted to the units of `st_crs(.x)` (only when `sf` >= 1.0-7).

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

**Examples**

```

if (require(plm, quietly = TRUE)) {
  data(Produc, package = "plm")
  st_as_stars(Produc)
}
if (require(dplyr, quietly = TRUE)) {
  # https://stackoverflow.com/questions/77368957/
  spatial_dim <- st_sf(
    ID = 1:3,
    geometry = list(
      st_polygon(list(
        cbind(c(0, 1, 1, 0, 0), c(0, 0, 1, 1, 0))
      )),
      st_polygon(list(
        cbind(c(1, 2, 2, 1, 1), c(0, 0, 1, 1, 0))
      )),
      st_polygon(list(
        cbind(c(2, 3, 3, 2, 2), c(0, 0, 1, 1, 0))
      ))
    )
  )
  weekdays_dim <- data.frame(weekdays = c("Monday", "Tuesday", "Wednesday",
    "Thursday", "Friday", "Saturday", "Sunday"))
  hours_dim <- data.frame(hours = c("8am", "11am", "4pm", "11pm"))
  sf_dta <- spatial_dim |>
    cross_join(weekdays_dim)|>
    cross_join(hours_dim) |>
    mutate(population = rnorm(n(), mean = 1000, sd = 200)) |>
    select(everything(), geometry)

  st_as_stars(sf_dta, dims = c("weekdays", "hours", "geometry"))
}
demo(nc, echo=FALSE,ask=FALSE)
st_as_stars(nc)
st_as_stars(st_drop_geometry(nc), dims = "NAME")
data.frame(expand.grid(x=1:5, y = 1:5), z = rnorm(25)) |> st_as_stars()
nc = st_read(system.file("gpkg/nc.gpkg", package="sf"))
st_as_stars(nc)

```

---

st\_cells

*return the cell index corresponding to the location of a set of points*


---

**Description**

If the object has been cropped without normalization, then the indices return are relative to the original uncropped extent. See [st\\_crop](#)

**Usage**

```
st_cells(x, sf)
```

**Arguments**

x                    object of class stars  
 sf                    object of class sf or sfc

**Examples**

```
set.seed(1345)
st_bbox(L7_ETMs) |>
  st_as_sfc() |>
  st_sample(10) -> pts
(x <- st_cells(L7_ETMs, pts))
# get the pixel values (first band only):
st_as_stars(L7_ETMs)[[1]][x]
# get pixel values for all bands:
st_as_stars(L7_ETMs) |> split() |> sapply(`[, x)
# compare with st_extract():
st_as_stars(L7_ETMs) |> split() |> st_extract(pts)
```

---

 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**

x                    object of class stars  
 na.rm                logical; should missing valued cells be removed, or also be converted to features?  
 contour\_lines       logical; if FALSE, polygons are returned (contour sets), otherwise contour lines  
 breaks               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, add_coordinates = TRUE)

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.
add_coordinates	logical; if 'TRUE', columns with dimension values precede the array values, otherwise they are omitted
.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 'mdim'
st_crop(x, y, ...)

## 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),
  normalize = FALSE
)
```

## 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

`normalize` logical; if TRUE then pass the cropped object to `st_normalize` before returning. This typically changes the 'offset' field and resets the 'from' field to 1, and changes the bounding box of the returned object accordingly.

## 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(l7))
l7[bb]
# equivalent:
st_crop(l7, bb)

plot(l7[,1:13,1:13,1], reset = FALSE)
image(l7[bb,,1], add = TRUE, col = sf.colors())
plot(st_as_sfc(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(l7))
l7[bb]

plot(l7[,1:13,1:13,1], reset = FALSE)
image(l7[bb,,1], add = TRUE, col = sf.colors())
plot(st_as_sfc(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(l7))
l7[bb]

plot(l7[,1:13,1:13,1], reset = FALSE)
image(l7[bb,,1], add = TRUE, col = sf.colors())
plot(st_as_sfc(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	<i>get dimensions from stars object</i>
---------------	---

---

## 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 'stars_proxy'
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, ..., where = NA, 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 an 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; setting special value NULL removes dimension values, for instance to remove curvilinear raster coordinates
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
where	character, one of 'start', 'center' or 'end'. Set to NA (default) to ignore and use max and center explicitly. This argument provides a convenient alternative to setting max and center.
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(µm) -> 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"))
m = matrix(1:20, nrow = 5, ncol = 4)
dim(m) = c(x = 5, y = 4) # named dim
(s = st_as_stars(m))
st_get_dimension_values(s, 'x', where = "start")
st_get_dimension_values(s, 'x', center = FALSE)
st_get_dimension_values(s, 'x', where = "center")
st_get_dimension_values(s, 'x', center = TRUE)
st_get_dimension_values(s, 'x', where = "end")
st_get_dimension_values(s, 'x', max = TRUE)
```

---

<code>st_dim_to_attr</code>	<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\_downsample                    *downsample stars or stars\_proxy objects*

---

**Description**

downsample a stars or stars\_proxy object either by skipping rows, columns and bands, or by computing a single value (e.g. the mean) from the sub-tiles involved

**Usage**

```
st_downsample(x, n, ...)

## S3 method for class 'stars'
st_downsample(x, n, ..., offset = 0, FUN)

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

**Arguments**

x                    object of class stars or stars\_proxy  
 n                    integer; for each dimension the number of pixels/lines/bands etc that will be skipped; see Details.  
 ...                  arguments passed on to FUN (e.g., na.rm = TRUE to ignore missing values if FUN is mean)  
 offset              integer; offset(s) for downsampling, in pixels, starting at the offset of each dimension; should be smaller or equal to n  
 FUN                  function; if given, downsampling will apply FUN to each of the the subtiles

## Details

If all `n == 0`, no downsampling takes place; if it is 1, every second row/column/band is skipped, if it is 2, every second+third row/column/band are skipped, etc.

Downsampling a `stars_proxy` object returns a `stars` object, is equivalent to calling `st_as_stars(x, downsample = 2)`, and only downsamples the first two (x and y) dimensions.

Downsampled regular rasters keep their dimension offsets, have a cell size (delta) that is `n[i]+1` times larger, and may result in a (slightly) different extent.

Note that terra's [aggregate](#) with `fact=2` corresponds to `st_downsample(x, n = 1, FUN = mean)`: `fact` is one larger than `n`.

## Examples

```
(m = matrix(1:121, 11, 11))
(s = st_as_stars(m))
st_downsample(s, 1)
st_downsample(s, 1)[[1]]
st_downsample(s, 1, offset = 1)
st_downsample(s, 1, offset = 1)[[1]]
st_downsample(s, 1, offset = c(0,1))
st_downsample(s, 1, offset = c(0,1))[[1]]
st_downsample(s, 1, FUN = mean)
st_downsample(s, 1, FUN = mean)[[1]]
st_downsample(s, 1, offset = 1, FUN = mean)
st_downsample(s, 1, offset = c(0,1), FUN = mean)[[1]]
```

---

st\_extract

*Extract cell values at point locations*

---

## Description

Extract cell values at point locations

## Usage

```
st_extract(x, ...)

## S3 method for class 'stars'
st_extract(
  x,
  at,
  ...,
  bilinear = FALSE,
  time_column = attr(at, "time_column") %||% attr(at, "time_col"),
  interpolate_time = bilinear,
  FUN = mean,
  resampling = c("nearest", "bilinear", "cubic", "cubicspline")
)
```

**Arguments**

x	object of class <code>stars</code> or <code>stars_proxy</code>
...	passed on to <code>aggregate.stars</code> when geometries are not exclusively POINT geometries
at	object of class <code>sf</code> or <code>sfc</code> with geometries, or two-column matrix with coordinate points in rows, indicating where to extract values of x
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 first temporal dimension (matching classes <code>POSIXct</code> , <code>POSIXt</code> , <code>Date</code> , or <code>PCICt</code> ), 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 <code>FALSE</code> , time instances are matched using the coinciding or the last preceding time in the data cube.
FUN	function used to aggregate pixel values when geometries of <code>at</code> intersect with more than one pixel
resampling	character; resampling method; for method <code>cubic</code> or <code>cubicspline</code> , ‘ <code>stars_proxy</code> ’ objects should be used and GDAL should have version <code>&gt;= 3.10.0</code>

**Details**

points outside the raster are returned as NA values. For large sets of points for which extraction is needed, passing a matrix as to `at` may be much faster than passing an `sf` or `sfc` object.

**Value**

if `at` is of class `matrix`, a matrix with extracted values is returned; otherwise: 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, if this is not the case, 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_extract(r, st_coordinates(pnt)) # "at" is a matrix: return a matrix
```



---

st_geotransform	<i>get or set the geotransform, or rotation matrix</i>
-----------------	--

---

**Description**

get or set the geotransform, or rotation matrix

**Usage**

```
st_geotransform(x, ...)

st_geotransform(x) <- value

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

**Arguments**

x	object of class stars or dimensions
...	ignored
value	length 6 numeric vector, or 2 x 2 (scaled) rotation matrix

**Examples**

```
# using the "classical" rotation matrix, see https://en.wikipedia.org/wiki/Rotation\_matrix :
rot = function(theta, dxdy = c(1., -1.)) {
  th = theta / 180 * pi
  matrix(c(cos(th), sin(th), -sin(th), cos(th)), 2, 2) %*%
    matrix(c(dxdy[2], 0, 0, dxdy[1]), 2, 2)
}
l = st_downsample(st_as_stars(L7_ETMs), 9) # save time in plotting
st_geotransform(l) = rot(20, c(28.5, 28.5)) # clockwise, 20 degrees, scale by cell size
plot(l[, , 1])
m = rot(20, c(1, 2))
g = expand.grid(x = 0:4, y = 0:4)
plot(g[1:2], asp = 1)
text(g[,1], g[,2], labels = seq_along(g[,1]), pos = 4)
g = t(m %*% t(as.matrix(g)))
points(g, col = 'red')
text(g[,1], g[,2], labels = seq_along(g[,1]), pos = 4, col = 'red')

m = matrix(1:20, 4)
s0 = st_as_stars(m)
s = s0
# dy > 0, clockwise rotation:
st_geotransform(s) = rot(10, c(1,1))
plot(s0, reset = FALSE)
plot(s, add = TRUE)
# dy < 0, counter clockwise rotation, + expansion in x-direction:
```

```

layout(1)
s0 = st_as_stars(st_bbox(s0), dx = 1)
s0$values = 1:20
s0
plot(s0, reset = FALSE)
s = s0
st_geotransform(s) = rot(10, c(2,1))
plot(s, add = TRUE)

```

---

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("-vrtnodata", "-9999", "-srcnodata", "nan"),
  file_ext = ".tif"
)

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

## S3 method for class 'stars_proxy'
st_mosaic(
  .x,
  ...,
  dst = tempfile(fileext = file_ext),
  options = c("-vrtnodata", "-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; this will be a VRT file with references to the source file(s), see details
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; note that for the 'stars' method, the 'dst' file may contain references to temporary files that are going to be removed at termination of the R session.

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

*rasterize simple feature geometries*


---

**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),
  align = FALSE,
  proxy = FALSE,
  ...
)
```

**Arguments**

sf	object of class sf
template	optional; stars object with desired target geometry, or target geometry alignment if align=TRUE; see details
file	temporary file name

driver	driver for temporary file
options	character; options vector for GDALRasterize
align	logical; if TRUE, template is only used for the geometry <code>_alignment_</code> , informing target resolution and offset
proxy	logical; should a proxy object be returned?
...	arguments passed on to <code>st_as_stars</code>

### Details

if ‘template’ is a ‘stars’ object, non-NA cells that are not covered by ‘sf’ receive the value in ‘template’; see also argument ‘align’.

### 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, dimension = character(0))
```

**Arguments**

x                    object of class stars  
 dimension           optional: numbers or names of dimension(s) to get per-dimension type

**Details**

categories "curvilinear" and "affine" only refer to the relationship between a pair of spatial (raster) dimensions.

**Value**

if dimension is not specified, return the spatial raster type: one of NA (if the object does not have raster dimensions), "curvilinear", "rectilinear", "affine", or "regular". In case dimension(s) are specified, return one of "regular", "rectilinear" (irregular but numeric), or "discrete" (anything else).

**Examples**

```
tif = system.file("tif/L7_ETMs.tif", package = "stars")
x = read_stars(tif)
st_raster_type(x)
st_raster_type(x, 1:3)
```

---

st_res	<i>obtain (spatial) resolution of a stars object</i>
--------	--

---

**Description**

obtain resolution(s) of a stars object: by default only the (absolute) x/y raster dimensions, optionally all delta dimension parameters

**Usage**

```
st_res(x, all = FALSE, absolute = !all)
```

**Arguments**

x                    an object of class stars  
 all                  logical; if FALSE return a vector with the x/y raster resolution  
 absolute            logical; only works when all = FALSE; if TRUE return absolute resolution values, if FALSE return delta values

**Value**

if all = FALSE a vector with x/y raster resolutions, otherwise a list with delta values

**Examples**

```

st_res(L7_ETMs)
st_res(L7_ETMs, absolute = FALSE)
st_res(L7_ETMs, all = TRUE)
if (require(starsdata)) {
  paste0("netcdf/", c("avhrr-only-v2.19810901.nc",
    "avhrr-only-v2.19810902.nc",
    "avhrr-only-v2.19810903.nc",
    "avhrr-only-v2.19810904.nc")) |>
  system.file(package = "starsdata") |>
  read_stars(quiet = TRUE) -> x
  st_res(x) |> print()
  st_res(x, all = TRUE) |> print()
}

```

---

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 = dim(x)[dimension] == 4,
  maxColorValue = 255L,
  probs = c(0, 1),
  stretch = NULL
)

```

**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 by "percent".
stretch	logical or character; if TRUE or "percent", each band is stretched to 0 ... maxColorValue by "percent clip" method using probs values. If "histogram", a "histogram equalization" is performed (probs values are ignored). If stretch is NULL or FALSE, no stretching is performed. Other character values are interpreted as "percent" and a message will be printed.



**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. Alternatively, you can use [plot.stars](#) with the `rgb` argument to create a three-band composition.

**See Also**

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

**Examples**

```
tif = system.file("tif/L7_ETMs.tif", package = "stars")
x = read_stars(tif)
st_rgb(x[,,,3:1])
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()
}
r = st_rgb(x[,,,3:1],
           probs = c(0.01, 0.99),
           stretch = "percent")
plot(r)
r = st_rgb(x[,,,3:1],
           probs = c(0.01, 0.99),
           stretch = "histogram")
plot(r)
```

---

st_rotate	<i>Transform rotated pole long/lat regular grid to unrotated curvilinear grid</i>
-----------	---

---

**Description**

Transform rotated long/lat regular grid to unrotated curvilinear grid

**Usage**

```
## S3 method for class 'stars'
st_rotate(.x, lon0, lat0, north = TRUE, ...)

## S3 method for class 'sfc'
st_rotate(.x, lon0, lat0, north = TRUE, ...)

## S3 method for class 'sf'
st_rotate(.x, lon0, lat0, north = TRUE, ...)
```

**Arguments**

.x	object of class stars
lon0	longitude of the rotated pole in degrees
lat0	latitude of the rotated pole in degrees
north	logical; if TRUE the pole refers to the North pole, otherwise the South pole
...	ignored

**Value**

curvilinear stars object with coordinates in regular long/lat (North pole at lat=90)

**Examples**

```
if (require("starsdata") && require("maps")) {
  # data downloaded from https://esgf-data.dkrz.de/search/cosmo-rea/
  nc = "netcdf/ts_EUR-6km_ECMWF-ERAINT_REA6_r1i1p1f1_COSMO_v1_mon_201801-201812.nc"
  f = system.file(nc, package = "starsdata")
  m = read_mdin(f, "ts")
  print(m)
  # NOTE this function is obsolete when reading m as
  # m = read_mdin(f, "ts", curvilinear = c("longitude", "latitude"))
  if (require(RNetCDF)) {
    x = open.nc(f)
    lon = att.get.nc(x, "rotated_latitude_longitude", "grid_north_pole_longitude")
    lat = att.get.nc(x, "rotated_latitude_longitude", "grid_north_pole_latitude")
    close.nc(x)
    print(c(lon = lon, lat = lat))
  } else {
    lon = -162
    lat = 39.25
  }
  m1 = st_rotate(m, lon, lat)
  print(m1)
  h = function() maps::map(add = TRUE)
  plot(m1, downsample = c(10, 10, 5), axes = TRUE, hook = h, mfrow = c(1, 2))
  # curvilinear grid: downsample for plotting speed
  m2 = st_warp(m1, crs = st_crs("OGC:CRS84"), threshold = .1)
  plot(m2, hook = h, mfrow = c(3, 4)) # regular grid: plots fast
}
```

---

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_tile	<i>Specify parameters to load raster in blocks</i>
---------	--

---

**Description**

Helper function for specifying the block parameters (nXOff, nYOff, nXsize, and nYSize) required by RasterIO argument in [read\\_stars](#)

**Usage**

```
st_tile(img_rows, img_cols, x_window, y_window, overlap = 0)
```

**Arguments**

img_rows	number of input raster rows (integer)
img_cols	number of input raster columns (integer)
x_window	number of rows in block (integer)
y_window	number of columns in block (integer)
overlap	number of overlapping pixels (integer)

**Value**

matrix with specified nXOff, nYOff, nXsize, and nYSize parameters for every block

**Examples**

```
## Not run:
tif = system.file("tif/L7_ETMs.tif", package = "stars")
r = read_stars(tif, proxy = TRUE)
tiles = st_tile(nrow(r), ncol(r), 256, 256)
for (i in seq_len(nrow(tiles))) {
  tile = read_stars(tif, proxy = FALSE, RasterIO = tiles[i, ])
  # write tiles to separate files
  write_stars(tile, dsn = paste0(i, ".tif"))
}

## End(Not run)
```

---

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, ...)

st_transform_proj.stars(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).

If array values contain geometries and an array as a whole is of class ‘sfc’ and has a non-missing CRS, array geometries are also transformed.

**See Also**

[st\\_warp](#)

**Examples**

```
geomatrix = system.file("tif/geomatrix.tif", package = "stars")
(x = read_stars(geomatrix))
new = st_crs('OGC:CRS84')
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",
  threshold = NA_real_
)
```

## 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 gdal's warp or warper, 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; not setting this when use_gdal is TRUE leads to a warning
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
threshold	numeric; distance threshold for warping curvilinear grids: new cells at distances larger than threshold are assigned NA values.

## 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](#). 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('OGC:CRS84')
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('OGC:CRS84') %>% 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)
# downsample raster (90 to 270 m)
r = read_stars(system.file("tif/olinda_dem_utm25s.tif", package = "stars"))
r270 = st_as_stars(st_bbox(r), dx = 270)
r270 = st_warp(r, r270)

```

---

st_xy2sfc	<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

x	object of class stars
as_points	logical; if TRUE, generate points at cell centers, else generate polygons
...	arguments passed on to st_as_sfc
na.rm	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.

---

write_stars	<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 = if (is.factor(obj[[1]]) && length(levels(obj[[1]])) < 256) "Byte" else "Float32",
  NA_value = NA_real_,
  update = FALSE,
  normalize_path = TRUE,
  scale_offset = c(1, 0)
)

## S3 method for class 'stars_proxy'
write_stars(
  obj,
  dsn,
  layer = 1,
  ...,
  driver = detect.driver(dsn),
  options = character(0),
  scale_offset = c(1, 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 dataset creation options, passed on to GDAL
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>
scale_offset	length 2 numeric vector with scale, offset values: raw values computed by raw = (value - offset) / scale are written to dsn; scale and offset values are written to dsn or else a warning is raised
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. in case obj is a multi-file stars\_proxy object, all files are written as layers into the output file dsn

**Examples**

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

---

%in%,stars-method      *evaluate whether cube values are in a given set*

---

**Description**

evaluate whether cube values are in a given set

**Usage**

```
## S4 method for signature 'stars'
x %in% table
```

**Arguments**

x	data cube value
table	values of the set

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