# Package 'starsExtra' 

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Description Miscellaneous functions for working with 'stars' objects, mainly single-band rasters. Currently includes functions for: (1) focal filtering, (2) detrending of Digital Elevation Models, (3) calculating flow length, (4) calculating the Convergence Index, (5) calculating topographic aspect and topographic slope.
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aspect Calculate topographic aspect from a DEM

## Description

Calculates topographic aspect given a Digital Elevation Model (DEM) raster. Input and output are rasters of class stars, single-band (i.e., only '"x"‘ and '"y"' dimensions), with one attribute.

## Usage

$\operatorname{aspect}(x$, na_flag $=-9999)$

## Arguments

$x \quad$ A raster (class stars) with two dimensions: $x$ and $y$, i.e., a single-band raster, representing a DEM.
na_flag Value used to mark NA values in C code. This should be set to a value which is guaranteed to be absent from the input raster $\times$ (default is -9999).

## Value

A stars raster with topographic slope, i.e., the azimuth where the terrain is tilted towards, in decimal degrees $(0-360)$ clockwise from north. Aspect of flat terrain, i.e., where all values in the neighborhood are equal, is set to -1 . Returned raster values are of class units (decimal degrees).

## Note

Aspect calculation results in NA when at least one of the cell neighbors is NA, including the outermost rows and columns. Given that the focal window size in aspect calculation is $3 * 3$, this means that the outermost one row and one column are given an aspect value of NA.
The raster must be in projected CRS, and units of $x / y$ resolution are assumed to be the same as units of elevation (typically meters).

## References

The topographic aspect algorithm is based on the How aspect works article in the ArcGIS documentation:
https://desktop.arcgis.com/en/arcmap/10.3/tools/spatial-analyst-toolbox/how-aspect-works. htm

## Examples

```
# Small example
data(dem)
dem_aspect = aspect(dem)
plot(
    dem, text_values = TRUE, breaks = "equal",
    col = hcl.colors(11, "Spectral"), main = "input (elevation)"
)
plot(
    dem_aspect, text_values = TRUE, breaks = "equal",
    col = hcl.colors(11, "Spectral"), main = "output (aspect)"
)
# Larger example
data(carmel)
carmel_aspect = aspect(carmel)
plot(
    carmel, breaks = "equal",
    col = hcl.colors(11, "Spectral"), main = "input (elevation)"
)
plot(
    carmel_aspect, breaks = "equal",
    col = hcl.colors(11, "Spectral"), main = "output (aspect)"
)
```

```
carmel Digital Elevation Model of Mount Carmel
```


## Description

A stars object representing a Digital Elevation Model (DEM) Digital Elevation Model of Mount Carmel, at 90 m resolution

## Usage

carmel

## Format

A stars object with 1 attribute:
elevation Elevation above sea level, in meters

## Examples

```
plot(carmel, breaks = "equal", col = terrain.colors(11))
```


## Description

Calculates the Convergence Index (CI) given a topographic slope raster. Input and output are rasters of class stars, single-band (i.e., only ‘"x"‘ and ‘"y"‘ dimensions), with one attribute.

## Usage

CI(x, k, na.rm = FALSE, na_flag = -9999)

## Arguments

x
k
na.rm Should NA values be ignored when calculating CI? Default is FALSE, i.e., when at least one aspect value in the neighborhood is NA the CI is also set to NA.
na_flag Value used to mark NA values in C code. This should be set to a value which is guaranteed to be absent from the input raster $\times$ (default is -9999).

## Value

A stars raster with CI values.

## Note

The raster is "padded" with $(k-1) / 2$ more rows and columns of NA values on all sides, so that the neighborhood of the outermost rows and columns is still a complete neighborhood. Those rows and columns are removed from the final result before returning it. Aspect values of -1 , specifying flat terrain, are assigned with a CI value of 0 regardless of their neighboring values.

## References

The Convergence Index algorithm is described in:
Thommeret, N., Bailly, J. S., \& Puech, C. (2010). Extraction of thalweg networks from DTMs: application to badlands.

## Examples

```
# Small example
data(dem)
dem_asp = aspect(dem)
dem_ci = CI(dem_asp, k = 3)
r=c(dem, round(dem_ci, 1), along = 3)
r = st_set_dimensions(r, 3, values = c("input (aspect)", "output (CI, k=3)"))
plot(r, text_values = TRUE, breaks = "equal", col = terrain.colors(10), mfrow = c(1, 2))
# Larger example
data(golan)
golan_asp = aspect(golan)
golan_ci = CI(golan_asp, k = 25)
plot(golan_asp, breaks = "equal", col = hcl.colors(11, "Spectral"), main = "input (aspect)")
plot(golan_ci, breaks = "equal", col = hcl.colors(11, "Spectral"), main = "output (CI, k=25)")
```

dem Small Digital Elevation Model

## Description

A stars object representing a small 13*11 Digital Elevation Model (DEM), at 90 m resolution

## Usage

dem

## Format

A stars object with 1 attribute:
elevation Elevation above sea level, in meters

## Examples

```
    plot(dem, text_values = TRUE, breaks = "equal", col = terrain.colors(11))
```

```
    detrend Detrend a Digital Elevation Model
```


## Description

Detrends a Digital Elevation Model (DEM) raster, by subtracting a trend surface. The trend is computed using mgcv: : gam or mgcv: : bam (when parallel>1) with formula $z \sim s(x, y)$.

## Usage

detrend(x, parallel = 1)

## Arguments

$x \quad$ A two-dimensional stars object representing the DEM
parallel Number of parallel processes. With parallel=1 uses ordinary, non-parallel processing.

## Value

A two-dimensional stars object, with two attributes:

- resid - the detrended result, i.e., "residual"
- trend - the estimated "trend" which was subtracted from the actual elevation to obtain resid


## Examples

```
# Small example
data(dem)
dem1 = detrend(dem)
dem1 = st_redimension(dem1)
dem1 = st_set_dimensions(dem1, 3, values = c("resid", "trend"))
plot(round(dem1), text_values = TRUE, col = terrain.colors(11))
# Larger example 1
data(carmel)
carmel1 = detrend(carmel, parallel = 2)
carmel1 = st_redimension(carmel1)
carmel1 = st_set_dimensions(carmel1, 3, values = c("resid", "trend"))
```

```
plot(carmel1, col = terrain.colors(11))
# Larger example 2
data(golan)
golan1 = detrend(golan, parallel = 2)
golan1 = st_redimension(golan1)
golan1 = st_set_dimensions(golan1, 3, values = c("resid", "trend"))
plot(golan1, col = terrain.colors(11))
```

dist_to_nearest Calculate raster of distances to nearest feature

## Description

Given a stars raster and an sf vector layer, returns a new raster with the distances of each cell centroid to the nearest feature in the vector layer.

## Usage

dist_to_nearest(x, v, progress = TRUE)

## Arguments

$x \quad$ A stars layer, used as a "grid" for distance calculations
$v \quad$ An sf, sfc or sfg object
progress Display progress bar? The default is TRUE

## Value

A stars raster with distances to nearest feature

## Examples

```
\# Sample 'sf' layer
\(\mathrm{x}=\) st_point(c(0,0))
y = st_point(c(1,1))
\(x=s t \_s f c(x, y)\)
x = st_sf(x)
x = st_buffer (x, 0.5)
\# Make grid
r = make_grid(x, res = 0.1, buffer = 0.5)
\(d=\) dist_to_nearest \((r, x\), progress \(=\) FALSE \()\)
\# Plot
plot(d, breaks = "equal", axes = TRUE, reset = FALSE)
plot(st_geometry (x), add = TRUE, pch = 4, cex = 3)
```


## Description

Extract raster values by lines or polygons, summarizing for each feature using a function specified by the user. This function is aimed to reproduce (some of) the functionality of raster: : extract.

## Usage

extract2(x, v, fun, progress = TRUE, ...)

## Arguments

| $x$ | A stars object |
| :--- | :--- |
| $v$ | An sf layer that determines values to extract |
| fun | A function to summarize cell values per feature/band |
| progress | Display progress bar? The default is TRUE |
| $\ldots$ | Further arguments passed to fun |

## Value

A vector (single-band raster) or matrix (multi-band raster) with the extracted and summarized values

## Examples

```
# Polygons
pol = st_bbox(landsat)
pol = st_as_sfc(pol)
set.seed(1)
pol = st_sample(pol, 5)
pol = st_buffer(pol, 100)
pol = c(pol, pol)
# Plot
plot(landsat[,,,1,drop=TRUE], reset = FALSE)
plot(pol, add = TRUE)
# Single-band raster
aggregate(landsat[,,,1,drop=TRUE], pol, mean, na.rm = TRUE)[[1]] ## Duplicated areas get 'NA'
extract2(landsat[,,,1,drop=TRUE], pol, mean, na.rm = TRUE, progress = FALSE)
# Multi-band example
extract2(landsat, pol, mean, na.rm = TRUE, progress = FALSE)
# Lines
```

```
lines = st_cast(pol, "LINESTRING")
# Single-band raster
extract2(landsat[,,,1,drop=TRUE], lines, mean, na.rm = TRUE, progress = FALSE)
# Multi-band example
extract2(landsat, lines, mean, na.rm = TRUE, progress = FALSE)
```

flowlength Calculate flow length

## Description

Calculates flow length for each pixel in a Digital Elevation Model (DEM) raster. Inputs and output are rasters of class stars, single-band (i.e., only '"x"‘ and ‘"y"‘ dimensions), with one attribute.

## Usage

flowlength(elev, veg, progress = TRUE)

## Arguments

| elev | A numeric stars raster representing a Digital Elevation Model (DEM). |
| :--- | :--- |
| veg | A matching logical stars raster representing vegetation presence. TRUE val- <br> ues represent vegetated cells where flow is absorbed (i.e. sinks), FALSE values <br> represent cells where flow is unobstructed. |
| progress | Display progress bar? The default is TRUE |

## Value

A numeric stars raster where each cell value is flow length, in resolution units.

## References

The algorithm is described in:
Mayor, A. G., Bautista, S., Small, E. E., Dixon, M., \& Bellot, J. (2008). Measurement of the connectivity of runoff source areas as determined by vegetation pattern and topography: A tool for assessing potential water and soil losses in drylands. Water Resources Research, 44(10).

## Examples

\# Example from Fig. 2 in Mayor et al. 2008
elev $=$ rbind(
$c(8,8,8,8,9,8,9)$, c(7, 7, 7, 7, 9, 7, 7), $c(6,6,6,6,6,5,7)$,

```
    c(4, 5, 5, 3, 5, 4, 7),
    c(4, 5, 4, 5, 4, 6, 5),
    c(3, 3, 3, 3, 2, 3, 3),
    c(2, 2, 2, 3, 4, 1, 3)
)
veg = rbind(
    c(TRUE, TRUE, TRUE, TRUE, FALSE, FALSE, TRUE),
    c(TRUE, TRUE, trUE, trUE, TRUE, FALSE, FALSE),
    c(FALSE, FALSE, FALSE, FALSE, FALSE, FALSE, FALSE),
    c(FALSE, TRUE, FALSE, FALSE, FALSE, FALSE, TRUE),
    c(TRUE, TRUE, FALSE, FALSE, FALSE, FALSE, FALSE),
    c(TRUE, TRUE, TRUE, FALSE, FALSE, FALSE, FALSE),
    c(FALSE, TRUE, TRUE, FALSE, FALSE, TRUE, TRUE)
)
elev = matrix_to_stars(elev)
veg = matrix_to_stars(veg)
# Calculate flow length
fl = flowlength(elev, veg, progress = FALSE)
# Plot
plot(
        round(elev, 1), text_values = TRUE, breaks = "equal",
        col = terrain.colors(6), main = "input (elevation)"
)
plot(
    veg*1, text_values = TRUE, breaks = "equal",
    col = rev(terrain.colors(2)), main = "input (vegetation)"
)
plot(
    round(fl, 1), text_values = TRUE, breaks = "equal",
    col = terrain.colors(6), main = "output (flowlength)"
)
# Larger example
data(carmel)
elev = carmel
elev[is.na(elev)] = 0
veg = elev > 100
fl = flowlength(elev, veg, progress = FALSE)
plot(fl, breaks = "equal", col = hcl.colors(11), main = "flowlength (m)")
```


## focal2 Apply a focal filter on a raster

## Description

Applies a focal filter with weighted neighborhood $w$ on a raster. The weights (w) can be added to, subtracted from, multiplied by or divided with the raster values (as specified with weight_fun).

The focal cell is then taken as the mean, sum, minimum or maximum of the weighted values (as specified with fun). Input and output are rasters of class stars, single-band (i.e., only '"x"‘ and ‘"y"‘ dimensions), with one attribute.

```
Usage
    focal2(
        x,
        w,
        fun = "mean",
        weight_fun = "*",
        na.rm = FALSE,
        mask = FALSE,
        na_flag = -9999
    )
```


## Arguments

x
w
fun A function to aggregate the resulting values for each neighborhood. Possible values are: "mean", "sum", "min", "max", and "mode". The default is "mean", i.e., the resulting values per neighborhood are averaged before being assigned to the new focal cell value.
weight_fun An operator which is applied on each pair of values comprising the cell value and the respective weight value, as in raster_value-weight. Possible values are: "+", "-", "*", "/". The default is "*", i.e., each cell value is multiplied by the respective weight.
na.rm Should NA values in the neighborhood be removed from the calculation? Default is FALSE.
mask If TRUE, pixels with NA in the input are set to NA in the output as well, i.e., the output is "masked" using the input (default is FALSE).
na_flag Value used to mark NA values in C code. This should be set to a value which is guaranteed to be absent from the input raster x (default is -9999).

## Value

The filtered stars raster.

## Note

The raster is "padded" with (nrow(w)-1)/2 more rows and columns of NA values on all sides, so that the neighborhood of the outermost rows and columns is still a complete neighborhood. Those rows and columns are removed from the final result before returning it. This means, for instance, that the outermost rows and columns in the result will be NA when using na.rm=FALSE.

## References

The function interface was inspired by function raster: : focal. The C code for this function is a modified and expanded version of the C function named applyKernel included with R package spatialfil.

## Examples

```
# Small example
data(dem)
dem_mean3 = focal2(dem, matrix(1, 3, 3), "mean")
r = c(dem, round(dem_mean3, 1), along = 3)
r = st_set_dimensions(r, 3, values = c("input", "output (mean, k=3)"))
plot(r, text_values = TRUE, breaks = "equal", col = terrain.colors(11))
# Larger example
data(carmel)
carmel_mean15 = focal2(carmel, matrix(1, 15, 15), "mean")
r = c(carmel, carmel_mean15, along = 3)
r = st_set_dimensions(r, 3, values = c("input", "output (mean, k=15)"))
plot(r, breaks = "equal", col = terrain.colors(11))
```

focal2r Apply a focal filter on a raster ( $R$ )

## Description

Applies a focal filter with neighborhood size $\mathrm{k} * \mathrm{k}$ on a raster (class stars), using R code. This function is relatively slow, provided here mainly for testing purposes or for custom using functions which are not provided by focal2.

## Usage

focal2r(x, w, fun, mask = FALSE, ...)

## Arguments

x
W
fun

A raster (class stars) with two dimensions: $x$ and $y$, i.e., a single-band raster
Weights matrix defining the neighborhood size around the focal cell, as well as the weights. For example, matrix $(1,3,3)$ implies a neighborhood of size $3 * 3$ with equal weights of 1 for all cells. Focal cell values are multiplied by the matrix values before being passed to function fun. The matrix must be square, i.e., with an odd number of rows and columns.

A function to be applied on each neighborhood, after it has been multiplied by the matrix. The function needs to accepts a vector (of length equal to length(w) and return a vector of length 1

| mask | If TRUE, pixels with NA in the input are set to NA in the output as well, i.e., the |
| :--- | :--- |
| output is "masked" with the input (default FALSE) |  |

## Value

The filtered stars raster

## Note

The raster is "padded" with one more row/column of NA values on all sides, so that the neigborhood of the outermost rows and columns is still a complete $3 \times 3$ neighborhood. Those rows and columns are removed from the filtered result before returning it.

## Examples

```
# Small example
data(dem)
dem1 = focal2r(dem, matrix(1,3,3), mean, na.rm = TRUE)
dem2 = focal2r(dem, matrix(1,3,3), min, na.rm = TRUE)
dem3 = focal2r(dem, matrix(1,3,3), max, na.rm = TRUE)
r = c(dem, round(dem1, 1), dem2, dem3, along = 3)
r = st_set_dimensions(r, 3, values = c("input", "mean", "min", "max"))
plot(r, text_values = TRUE, breaks = "equal", col = terrain.colors(10))
# Larger example
data(carmel)
carmel1 = focal2r(carmel, matrix(1,3,3), mean, na.rm = TRUE, mask = TRUE)
carmel2 = focal2r(carmel, matrix(1,9,9), mean, na.rm = TRUE, mask = TRUE)
carmel3 = focal2r(carmel, matrix (1,15,15), mean, na.rm = TRUE, mask = TRUE)
r = c(carmel, carmel1, carmel2, carmel3, along = 3)
r = st_set_dimensions(r, 3, values = c("input", "k=3", "k=9", "k=15"))
plot(r, breaks = "equal", col = terrain.colors(100))
```


## Description

Calculates a polygon layer with the footprints of raster images.

## Usage

footprints(x)

## Arguments

## Value

An sf layer with the footprints (i.e., bounding box polygons) of the rasters

## Examples

```
    # Create sample files
    file1 = tempfile(fileext = ".tif")
    file2 = tempfile(fileext = ".tif")
    file3 = tempfile(fileext = ".tif")
    r1 = landsat[,1:100, 1:100,]
    r2 = landsat[,101:200, 101:200,]
    r3 = landsat[,21:40, 51:120,]
    write_stars(r1, file1)
    write_stars(r2, file2)
    write_stars(r3, file3)
    # Calculate footprints
    files = c(file1, file2, file3)
    pol = footprints(files)
    pol
    # Plot
    plot(pol)
```

    golan Digital Elevation Model of Golan Heights
    
## Description

A stars object representing a Digital Elevation Model (DEM) Digital Elevation Model of part of the Golan Heights and Lake Kinneret, at 90 m resolution

## Usage

golan

## Format

A stars object with 1 attribute:
elevation Elevation above sea level, in meters

## Examples

```
plot(golan, breaks = "equal", col = terrain.colors(11))
```

landsat $\quad$ RGB image of Mount Carmel

## Description

A stars object representing an RGB image of part of Mount Carmel, at 30m resolution. The data source is Landsat-8 Surface Reflectance product.

## Usage

landsat

## Format

A stars object with 1 attribute:
refl Reflectance, numeric value between 0 and 1

## Examples

```
plot(landsat, breaks = "equal")
```

    layer_to_matrix Get stars layer values as matrix
    
## Description

Extracts the values of a single layer in a stars object to a matrix.

## Usage

layer_to_matrix(x, check = TRUE)

## Arguments

| $x$ | A stars raster with one attribute and two dimensions, $x$ and $y$, i.e., a single-band |
| :--- | :--- |
| raster. |  |$\quad$| Whether to check (and fix if necessary) that input has one attribute, one layer |
| :--- |
| and $x-y$ as dimensions 1-2 (default is TRUE). |

## Value

A matrix with the layer values, having the same orientation as the raster (i.e., rows represent the y -axis and columns represent the x -axis).

## Examples

```
data(dem)
m = layer_to_matrix(dem)
m
```

layer_to_vector Get stars layer values as vector

## Description

Extracts the values of a single layer in a stars object to a vector. Cell values are ordered from top-left corner to the right.

## Usage

layer_to_vector (x, check = TRUE)

## Arguments

| $x$ | A raster (class stars) with two dimensions: $x$ and $y$, i.e., a single-band raster. |
| :--- | :--- |
| check | Whether to check (and fix if necessary) that input has one attribute, one layer <br> and $x-y$ as dimensions 1-2 (default is TRUE). |

## Value

A vector with cell values, ordered by rows, starting from the top left corner (north-west) and to the right.

## Examples

```
data(dem)
v = layer_to_vector(dem)
v
```

make_grid

```
    make_grid Make 'stars' grid from 'sf' layer
```


## Description

Create 'stars' raster grid from bounding box of 'sf' vector layer, possibly buffered, with specified resolution.

## Usage

make_grid(x, res, buffer = 0)

## Arguments

| $x$ | An $s f, s f c$ or $s f g$ object |
| :--- | :--- |
| res | Required grid resolution, in CRS units of $x$ |
| buffer | Buffer size around $x$ (default is 0, i.e., no buffer) |

## Value

A stars raster with the grid, with all cell values equal to 1

## Examples

\# Sample 'sf' layer
$x=$ st_point $(c(0,0))$
$y=s t \_p o i n t(c(1,1))$
$x=s t \_s f c(x, y)$
$x=s t \_s f(x)$
\# Make grid
$r=$ make_grid(x, res = 0.1, buffer $=0.5$ )
$r[[1]][]=\operatorname{rep}(1: 3$, length.out $=$ length $(r[[1]]))$
\# Plot
plot $(r$, axes $=$ TRUE, reset $=$ FALSE $)$
plot(st_geometry(x), add = TRUE, pch = 4, cex = 3, col = "red")

```
matrix_extend Extend matrix
```


## Description

Adds n rows and columns with NA values on all sides of a matrix.

## Usage

matrix_extend(m, $\mathrm{n}=1$, fill = NA)

## Arguments

m
A matrix
$\mathrm{n} \quad$ By how many rows/columns to extend the matrix on each side?
fill Fill value (default is NA)

## Value

An extended matrix

## Examples

```
m = matrix(1:6, nrow = 2, ncol = 3)
m
matrix_extend(m, 1)
matrix_extend(m, 2)
matrix_extend(m, 3)
```

matrix_get_neighbors Get neighboring cell values for given matrix cell

## Description

Get the values of a $k^{*} k$ neighborhood, as vector and by row, given a matrix, $k$, and focal cell position (row and column).

## Usage

matrix_get_neighbors(m, pos, $k=3$ )

## Arguments

m
pos
k

A matrix.
The focal cell position, a numeric vector of length two of the form c (row, column).
Neighborhood size around the focal cell. For example, $k=3$ implies a neighborhood of size $3 * 3$. Must be an odd positive integer.

## Value

A vector with cell values, ordered by rows, starting from the top left corner of the neighborhood and to the right. When neighborhood extends beyond matrix bounds, only the "existing" values are returned.

## Examples

```
m = matrix(1:12, nrow = 3, ncol = 4)
m
matrix_get_neighbors(m, pos = c(2, 2), k = 3)
matrix_get_neighbors(m, pos = c(2, 2), k = 5)
matrix_get_neighbors(m, pos = c(2, 2), k = 7) # Same result
```

```
matrix_to_stars Convert matrix to stars
```


## Description

Converts matrix to a single-band stars raster, conserving the matrix orientation where rows become the $y$-axis and columns become the $y$-axis. The bottom-left corner of the axis is set to $(0,0)$ coordinate, so that x and y coordinates are positive across the raster extent.

## Usage

matrix_to_stars(m, res = 1)

## Arguments

m
A matrix
res
The cell size, default is 1

## Value

A stars raster

## Examples

```
data(volcano)
r = matrix_to_stars(volcano, res = 10)
plot(r)
```

```
matrix_trim Trim matrix
```


## Description

Removes n rows and columns with NA values on all sides of a matrix.

## Usage

matrix_trim(m, $\mathrm{n}=1)$

## Arguments

m
A matrix
n
By how many rows/columns to trim the matrix on each side?

## Value

A trimmed matrix, or NULL if trimming results in an "empty" matrix.

## Examples

```
m = matrix(1:80, nrow = 8, ncol = 10)
m
matrix_trim(m, 1)
matrix_trim(m, 2)
matrix_trim(m, 3)
matrix_trim(m, 4)
```

| mode_value $\quad$ Mode |
| :--- | :--- |

## Description

Find the mode (i.e., most common value) in a numeric vector. In case of ties, the first encountered value is returned.

## Usage

mode_value(x, na_flag = -9999)

## Arguments

$x \quad$ A numeric or logical vector
na_flag Value used to mark NA values in C code. This should be set to a value which is guaranteed to be absent from the input vector $x$ (default is -9999).

## Value

The mode, numeric vector of length 1

## Examples

```
x = c(3, 2, 5, 5, 3, 10, 2, 5)
mode_value(x)
```

normalize_2d Normalize a 2 D 'stars' object

## Description

Check, and possibly correct, that the input stars object:

- Has exactly one attribute.
- Has exactly two dimensions.
- The dimensions are spatial, named $x$ and $y$ (in that order).


## Usage

normalize_2d(x)

## Arguments

x
A stars object

## Value

A new stars object

## Examples

```
# Small example
data(dem)
normalize_2d(dem)
```

```
normalize_3d Normalize a 3D 'stars' object
```


## Description

Check, and possibly correct, that the input stars object:

- Has exactly one attribute.
- Has exactly three dimensions.
- The first two dimensions are spatial, named $x$ and $y$ (in that order).


## Usage

normalize_3d(x)

## Arguments

x
A stars object

## Value

A new stars object

## Examples

\# Small example
data(landsat)
normalize_3d(landsat)

```
rgb_to_greyscale Convert RGB to greyscale
```


## Description

Convert a 3-band RGB raster to 1-band greyscale raster. Based on wvtool : : rgb2gray.

## Usage

rgb_to_greyscale(x, rgb $=1: 3$, coefs $=c(0.3,0.59,0.11)$ )

## Arguments

| $x$ | A three-dimensional stars object with RGB values |
| :--- | :--- |
| rgb | Indices of RGB bands, default is $c(1,2,3)$ |
| coefs | RGB weights, default is $c(0.30,0.59,0.11)$ |

## Value

A two-dimensional stars object greyscale values

## Examples

```
data(landsat)
plot(landsat, rgb = 1:3)
landsat_grey = rgb_to_greyscale(landsat)
plot(landsat_grey, breaks = "equal")
```

slope Calculate topographic slope from a DEM

## Description

Calculates topographic slope given a Digital Elevation Model (DEM) raster. Input and output are rasters of class stars, single-band (i.e., only " $x$ "‘ and ‘"y"‘ dimensions), with one attribute.

## Usage

slope(x, na_flag = -9999)

## Arguments

$x \quad$ A raster (class stars) with two dimensions: $x$ and $y$, i.e., a single-band raster, representing a DEM.
na_flag Value used to mark NA values in C code. This should be set to a value which is guaranteed to be absent from the input raster $\times$ (default is -9999).

## Value

A stars raster with topographic slope, i.e., the azimuth where the terrain is tilted towards, in decimal degrees (0-360) clockwise from north.

Note
Slope calculation results in NA when at least one of the cell neighbors is NA, including the outermost rows and columns. Given that the focal window size in slope calculation is $3 * 3$, this means that the outermost one row and one column are given an slope value of NA.
The raster must be in projected CRS, and units of $x / y$ resolution are assumed to be the same as units of elevation (typically meters).

## References

The topographic slope algorithm is based on the How slope works article in the ArcGIS documentation:

```
https://desktop.arcgis.com/en/arcmap/10.3/tools/spatial-analyst-toolbox/how-slope-works.
```

htm

## Examples

```
# Small example
data(dem)
dem_slope = slope(dem)
plot(
    dem, text_values = TRUE, breaks = "equal",
    col = hcl.colors(11, "Spectral"), main = "input (elevation)"
)
plot(
    dem_slope, text_values = TRUE, breaks = "equal",
    col = hcl.colors(11, "Spectral"), main = "output (slope)"
)
# Larger example
data(carmel)
carmel_slope = slope(carmel)
plot(
    carmel, breaks = "equal",
    col = hcl.colors(11, "Spectral"), main = "input (elevation)"
)
plot(
    carmel_slope, breaks = "equal",
    col = hcl.colors(11, "Spectral"), main = "output (slope)"
)
```

    trim2
        Remove empty outer rows and columns
    
## Description

Removes complete outer rows and columns which have NA values.

## Usage

$\operatorname{trim} 2(x)$

## Arguments

x
A two-dimensional stars object

## Value

A new stars object with empty outer rows and columns removed

## Examples

```
# Single-band example
data(dem)
dem[[1]][1,] = NA
dem1 = trim2(dem)
# Multi-band example
data(landsat)
landsat[[1]][1:100,,] = NA
landsat1 = trim2(landsat)
```


## Description

Creates a matrix with directions (i.e., azimuth) to central cell, of specified size k. The matrix can be used as weight matrix when calculating the convergence index (see Examples).

## Usage

w_azimuth(k)

## Arguments

k Neighborhood size around focal cell. Must be an odd number. For example, k=3 implies a $3 * 3$ neighborhood.

## Value

A matrix where each cell value is the azimuth from that cell towards the matrix center.

## Examples

```
m = w_azimuth(3)
m
m = w_azimuth(5)
m
```


## Description

Creates a matrix with where a circular pattern is filled with values of 1 and the remaining cells are filled with values of 0 (see Examples).

## Usage

w_circle(k)

## Arguments

k Neighborhood size around focal cell. Must be an odd number. For example, k=3 implies a 3*3 neighborhood.

## Value

A matrix with a circular pattern.

## Examples

```
m = w_circle(3)
image(m, asp = 1, axes = FALSE)
m = w_circle(5)
image(m, asp = 1, axes = FALSE)
m = w_circle(15)
image(m, asp = 1, axes = FALSE)
m = w_circle(35)
image(m, asp = 1, axes = FALSE)
m = w_circle(91)
image(m, asp = 1, axes = FALSE)
m = w_circle(151)
image(m, asp = 1, axes = FALSE)
```


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