Package ‘habtools’

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Author Joshua Madin [aut] (<https://orcid.org/0000-0002-5005-6227>), Nina Schiettekatte [aut, cre] (<https://orcid.org/0000-0002-1925-3484>)
Maintainer Nina Schiettekatte <nina.schiettekatte@gmail.com>
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cell_count_1d

---

**Description**

A helper function for segment, box and cube counting fractal methods. The function divide the array into \( n \) pieces and counts how many are occupied.

**Usage**

\[
\text{cell_count}_1d(\text{pts}, \text{xmin}, \text{xmax}, \text{n})
\]

**Arguments**

- **pts**: Data frame with x coordinates
- **xmin**: Minimum x-value
- **xmax**: Maximum x-value
- **n**: Multiplier

**Value**

Number of filled cells

**Examples**

pts <- data.frame(x = rnorm(200, 0, 5))

\[
\text{cell_count}_1d(\text{pts}, \text{xmin} = \min(\text{pts}\$x), \text{xmax} = \max(\text{pts}\$x), \text{n} = 5)
\]

---

**cell_count_2d**

---

**Description**

A helper function for segment, box and cube counting fractal methods. The function divide the array into \( n \) pieces and counts how many are occupied.

**Usage**

\[
\text{cell_count}_2d(\text{pts}, \text{xmin}, \text{xmax}, \text{ymin}, \text{ymax}, \text{n})
\]
cell_count_3d

Arguments

- **pts**: Data frame with x and y coordinates
- **xmin**: Minimum x-value
- **xmax**: Maximum x-value
- **ymin**: Minimum y-value
- **ymax**: Maximum y-value
- **n**: Multiplier

Value

Number of filled cells

---

Description

A helper function for segment, box and cube counting fractal methods. The function divide the array into $n$ pieces and counts how many are occupied.

Usage

```r
cell_count_3d(pts, xmin, xmax, ymin, ymax, zmin, zmax, n)
```

Arguments

- **pts**: Data frame with x, y, and z coordinates
- **xmin**: Minimum x-value
- **xmax**: Maximum x-value
- **ymin**: Minimum y-value
- **ymax**: Maximum y-value
- **zmin**: Minimum z-value
- **zmax**: Maximum z-value
- **n**: Multiplier

Value

Number of filled cells
**centroid**

*Calculate the centroid of 3D points*

**Description**
Calculates the centroid for a given set of XYZ coordinates.

**Usage**

```r
centroid(data)
```

**Arguments**

- `data` A data frame with x, y, and z coordinates.

**Value**
The coordinates of the centroid.

**Examples**

```r
data <- mesh_to_points(mcap)
centroid(data)
```

---

**circularity**

*Calculate circularity of a 2D shape*

**Description**
The perimeter of the 2D shape is divided by the perimeter of a circle with the same area as the shape. The more irregular the shape is, the closer the output value is to zero. The closer the shape is to a circle, the closer the output value is to 1.

**Usage**

```r
circularity(data)
```

**Arguments**

- `data` A data frame with the first two columns x and y coordinates, respectively.

**Value**
A value between 0 (infinitely irregular) and 1 (a perfect circle).
convexity

See Also

sphericity()

Examples

mcap_2d <- mesh_to_2d(mcap)
plot(mcap_2d, asp=1)
circularity(mcap_2d)

circ <- sim_circle() # simulate xy coordinates for a circle
plot(circ, asp=1)
circularity(circ)

convexity

Calculate convexity of a 3D mesh

Description

The ratio of the volume of the object and the volume of the convex hull around the object. Objects with fewer concavities will be closer to 1.

Usage

convexity(mesh)

Arguments

mesh A triangular mesh of class mesh3d. #'

Value

The convexity value.

Examples

convexity(mcap)
**Description**

Calculates mechanical vulnerability of rigid, cantilever-type structural elements.

**Usage**

csf(mesh, z_min, res, keep_data = FALSE)

**Arguments**

- **mesh**: A triangular mesh of class mesh3d.
- **z_min**: The z plane about which csf should be calculated. Defaults to min(z).
- **res**: The resolution to be used for the calculation. Defaults to the resolution of the mesh.
- **keep_data**: Logical. Return list with supplemental info? Defaults to FALSE.

**Details**

This function calculates the mechanical vulnerability of a structural element, like a hard coral colony, to fluid flow. While developed for corals, and originally called the Colony Shape Factor (CSF), the function is applicable to any attached, rigid cantilever type structure. CSF is dimensionless and can be used to compare the vulnerability among structures. Mechanistically, if the CSF of a structure becomes greater than the dislodgement mechanical threshold, breakage occurs. This threshold is a function of material tensile strength and inversely related to fluid velocity and density (Madin & Connolly 2006).

**Value**

A value for csf or if keep_data = TRUE, a list containing the colony shape factor (csf), the parallel to flow (dy) and perpendicular (dx) diameters of the cantilever base, and the bending moment (mom).

**Note**

The orientation of the 3D mesh is important for this function. The function assumes the fluid flow is parallel with the y-axis. The function also assumes the base of the cantilever over which the bending moment acts can be approximated as an ellipse with the diameter on the y-axis parallel with flow (dy). You can set a z_min if the base of your mesh is not flat at the base (i.e., shift the plane upon which the cantilever is attached upwards). The function output includes dy and dx for monitoring anticipated values.

**References**

Examples

csf(mcap, z_min = -3.65)
csf(mcap, z_min = -3.65, keep_data = TRUE)

dem_cropped

Crop DEM around points

Description

A function for sampling a DEM by cropping squares of a given size around xy coordinates.

Usage

dem_cropped(data, x0, y0, L, plot = FALSE)

Arguments

- `data`: A DEM in RasterLayer format.
- `x0`: A value or vector of central x coordinate(s).
- `y0`: A value or vector of central y coordinate(s).
- `L`: Size of squares to cropped from the DEM.
- `plot`: Logical. Plot the DEM and the cropped sections?

Value

A cropped RasterLayer or list of RasterLayers.

Examples

# around one point
dem_cropped <- dem_cropped(horseshoe, -468, 1266, L = 2)
raster::plot(dem_cropped)
points(-468, 1266)

# around multiple points
points <- data.frame(x = c(-467, -465, -466), y = c(1270, 1265, 1268))
dem_list <- dem_cropped(horseshoe, points$x, points$y, L = 1, plot = TRUE)

# plot the first element
raster::plot(dem_list[[1]])
**dem_sample**

Sample a random DEM with specified size from a larger DEM

**Usage**

dem_sample(data, L, allow_NA = TRUE, plot = FALSE, max_iter = 100)

**Arguments**

data  
Digital elevation model of class RasterLayer.

L  
Size of square to cut out of DEM.

allow_NA  
Logical. Allow NAs in the sample? Useful when DEM is not regular.

plot  
Logical. Plot the DEM and the cropped section?

max_iter  
Maximum number of random crops to try when allow_NA = FALSE before failing.

**Value**

Digital elevation model of class RasterLayer.

**Note**

Not allowing NAs may increase sampling time for irregular DEMs that contain a lot of NAs; e.g., structure from motion transects.

**Examples**

dem <- dem_sample(horseshoe, L = 2, plot=TRUE)

---

**dem_split**

Split DEM into smaller tiles

**Description**

Split DEM into smaller tiles

**Usage**

dem_split(data, size, parallel = FALSE, ncores = (parallel::detectCores() - 1))
dem_to_points

Arguments

- **data**: Digital elevation model of class RasterLayer.
- **size**: Size of tiles, in the same unit as the RasterLayer.
- **parallel**: Logical. Use parallel processing? Note: parallel must be installed.
- **ncores**: Number of cores to use when parallel = TRUE.

Value

List of RasterLayers.

Examples

```r
L <- habtools::extent(horseshoe)  # size of horseshoe = 8m
size <- 2  # size of target tiles
(L / size)^2  # number of target tiles = 16
dem_list <- dem_split(horseshoe, 2)
length(dem_list)
```

dem_to_points  
_Transform DEM to 3D pointcloud of raster corners_

Description

Transform DEM to 3D pointcloud of raster corners

Usage

```r
dem_to_points(dem, bh = NULL, parallel = FALSE)
```

Arguments

- **dem**: Digital elevation model of class RasterLayer.
- **bh**: Border height from lowest point.
- **parallel**: Logical. Use parallel computation?

Value

A 3D point cloud for raster cell corners.

Examples

```r
dem <- sim_dem(20, 0.5)
raster::plot(dem)
pts <- dem_to_points(dem)
rgl::plot3d(pts)
```
detect_drop

Detect a sudden drop, edge, or overhang in a DEM

description
Detect a sudden drop, edge, or overhang in a DEM

Usage

\texttt{detect\_drop(data, d = 0.1)}

Arguments

\begin{itemize}
  \item \textit{data} \quad \text{DEM of class \texttt{RasterLayer}.}
  \item \textit{d} \quad \text{The threshold height difference to define a drop.}
\end{itemize}

Value

A \texttt{RasterLayer} marking edges. Values indicate maximum height difference of surrounding cells.

Examples

\begin{verbatim}
edges <- detect\_drop(horseshoe, d = 0.2)
raster::plot(horseshoe)
raster::plot(edges)
\end{verbatim}

extent

Calculate extent of a 3D object

Description

This function calculates the extent or largest length of the bounding box of a mesh or a DEM.

Usage

\texttt{extent(data)}

Arguments

\begin{itemize}
  \item \textit{data} \quad \text{Digital elevation model of class \texttt{RasterLayer} or a triangular mesh of class \texttt{mesh3d}.}
\end{itemize}

Value

A value, the extent of the mesh or DEM.
Note

There are several extent function in other packages, including the raster package. Therefore it is recommended to use the package namespace, see examples below.

Examples

```r
habtools::extent(mcap)
habtools::extent(horseshoe)
```

---

**fd**

*Calculate fractal dimension*

**Description**

Calculate fractal dimension

**Usage**

```r
fd(data, method, lvec, keep_data = FALSE, diagnose = FALSE, ...)
```

**Arguments**

- `data`: Digital elevation model of class RasterLayer or a triangular mesh of class mesh3d.
- `method`: If data is a RasterLayer, possible methods are: "hvar", "area", "sd", and "cubes" (defaults to "hvar"). If data is a mesh3d, possible methods are "cubes" and "area" (defaults to "cubes").
- `lvec`: Vector of scales to use for calculation.
- `keep_data`: Logical. Keep data? Default is FALSE.
- `diagnose`: Logical. Show diagnostic plot and metrics?
- `...`: Arguments from method-specific fd_ functions.

**Details**

Calculates fractal dimension using the specified method. Note that methods are distinctly different and should not be mixed when comparing values for multiple objects. See `fd_hvar()`, `fd_area()`, `fd_cubes()`, `fd_sd()` for details about each method. If `lvec` is not specified, a default based on resolution, extent, and method will be used. The cubes method is not recommended if the height range is much smaller than the extent of a 3d object or DEM, which is typically the case for DEMs. Most objects and surfaces are not perfectly fractal. It is recommended to investigate scale transitions by setting `diagnose` to TRUE.

**Value**

A value for fractal dimension, typically between 2 and 3 or a list if `keep_data` = TRUE.


**fd_area**

Calculate fractal dimension using the surface area method

**Description**

Calculate fractal dimension using the surface area method

**Usage**

```r
fd_area(data, lvec = NULL, keep_data = FALSE, plot = FALSE, scale = FALSE)
```

**Arguments**

- **data**: DEM of class "RasterLayer" or mesh of class "mesh3d".
- **lvec**: Vector of scales to use for calculation.
- **keep_data**: Logical. Keep data? Default is FALSE.
- **plot**: Logical. Plot surface with area resolutions superimposed? Defaults to FALSE.
- **scale**: Logical. Rescale height values to fit the extent? Only relevant for DEMs. Defaults to FALSE.

**Details**

This function calculates fractal dimension using the area method. Based on values in `lvec`, the DEM or mesh is reprojected to varying scales. Fractal dimension is defined as $2 - s$ with $s$ being the slope of the regression between the log-transformed surface areas across scales and the log-transformed scales. Considerate bias is introduced if scales approach the extent of the object due to an edge effect. Therefore, this approach is only appropriate when the object is large relative to the scales of interest to be used as `lvec`. Diagnostic plots may help visualize whether bias is present for the scales chosen (i.e. points do not fall on a straight line).
Value

A value for fractal dimension, typically between 2 and 3 or a list if keep_data = TRUE.

Examples

```r
fd_area(horseshoe, lvec = c(0.125, 0.25, 0.5))

# Look at diagnostic plot
fdata <- fd_area(horseshoe, lvec = c(0.05, 0.1, 0.2, 0.4), keep_data = TRUE)
fdata <- fd_diagnose(fdata)
# points fall on straight line

fdata <- fd_area(horseshoe, lvec = c(0.5, 1, 2, 4), keep_data = TRUE)
fdata <- fd_diagnose(fdata)
# points fall on hollow curve, indicating that lvec includes values that
# are too high.
```

---

**fd_boxes**

*Calculate fractal dimension using the box counting method*

**Description**

Calculate fractal dimension using the box counting method

**Usage**

```r
fd_boxes(data, lvec, keep_data = FALSE, plot = FALSE)
```

**Arguments**

- `data`:
  A data frame in which the first two columns are x and y coordinates, respectively.

- `lvec`:
  The scales to use for calculation (i.e. box sizes).

- `keep_data`:
  Logical. Keep calculation data? Default = TRUE.

- `plot`:
  Logical. Plot the shape with box sizes superimposed? Defaults to FALSE.

**Details**

This function calculates fractal dimension using the box counting method. If lvec is not specified, a default based on resolution and extent will be used. Based on lvec, boxes of different sizes are defined and the function counts boxes that capture the outline of the shape. It is recommended to specify the maximum value of lvec so that the largest box encapsulates the entire object. The smallest scale included in lvec should not be smaller than the resolution of your object.

**Value**

A value for fractal dimension, typically between 1 and 2 or a list if keep_data = TRUE.
Examples

```r
mcap_2d <- mesh_to_2d(mcap)
fd_boxes(mcap_2d, plot = TRUE, keep_data = TRUE)
fd_boxes(mcap_2d, lvec = c(0.05, 0.1, 0.2, 0.4), plot = TRUE)
```

---

**fd_cubes**

*Calculate fractal dimension using the cube counting method*

**Description**

Calculate fractal dimension using the cube counting method

**Usage**

```r
fd_cubes(data, lvec = NULL, plot = FALSE, keep_data = FALSE, scale = FALSE)
```

**Arguments**

- `data`: An object of class RasterLayer or mesh3d.
- `lvec`: Vector of scales to use for calculation (i.e. cube sizes).
- `plot`: Planar representation of cubes superimposed on 3D mesh or DEM for visualizing lvec. Default = FALSE.
- `keep_data`: Logical. Keep calculation data? Default = TRUE.
- `scale`: Logical. Rescale height values to the extent? Only relevant for RasterLayer objects. (Defaults to FALSE).

**Details**

This function calculates fractal dimension using the cube counting method. If lvec is not specified, a default based on resolution and extent will be used. Based on lvec, cubes of different sizes are defined and the function counts mesh points that fall within each cube. It is recommended to specify the maximum value of lvec so that the largest box encapsulates the entire object. The smallest scale included in lvec should not be smaller than the resolution of your object.

**Value**

A value for fractal dimension, typically between 2 and 3 or a list if keep_data = TRUE.

**See Also**

`fd()`
Examples

```r
fd_cubes(mcap, keep_data = TRUE, plot = TRUE)
fd_cubes(mcap, lvec = c(0.05, 0.1, 0.25, 0.5), plot = TRUE)

dem <- dem_crop(horseshoe, x0 = -469, y0 = 1267, L = 2, plot = TRUE)
fd_cubes(dem, plot = TRUE, keep_data = TRUE)
fd_cubes(dem, plot = TRUE, keep_data = TRUE, scale = TRUE)
```

---

### fd_diagnose

#### Description

Diagnoses fractal dimension variation across neighboring scales.

#### Usage

```r
fd_diagnose(data, keep_data = TRUE)
```

#### Arguments

- **data**: Output of `fd()` with option `keep_data = TRUE`.
- **keep_data**: Logical. Keep diagnostics data?

#### Value

A list with fractal dimension across scales, mean fractal dimension, and sd of fractal dimensions across scales.

#### Examples

```r
fd_data <- fd(horseshoe, lvec = c(0.05, 0.1, 0.2, 0.4), method = "area", keep_data = TRUE)
fd_diagnose(fd_data)
fd_diagnose(fd_data, keep_data = FALSE)
```
Calculate fractal Dimension using the height variation method

Description
Calculate fractal Dimension using the height variation method

Usage

\[
\text{fd_hvar}(\text{data, lvec, regmethod = "mean", keep_data = FALSE, plot = FALSE, parallel = FALSE, ncores = (parallel::detectCores() - 1)}
\]

Arguments

data Digital elevation model of class RasterLayer or dataframe (output of hvar function)
lvec Vector of scales to use for calculation.
regmethod Method to use for linear regression between scale (lvec) and height range. One of raw (all data), mean (default) median or ends (minimum and maximum scale only)
keep_data Keep the data used for fd calculation? defaults to FALSE
plot Logical. Show plot of scales relative to data?
parallel Logical. Use parallel processing? Note: parallel must be installed.
defaults to FALSE
ncores Number of cores to use when parallel = TRUE.

Details
Calculates fractal dimension using the height variation regression. If lvec is not specified, a default based on resolution and extent will be used. data can be a DEM or a dataframe with columns labeled l and h for grid cell length and height range of that cell, respectively (output of hvar()). A rule of thumb is that l should range an order of magnitude. However, large ranges also average-out fractal dimension of a surface that might have phase transitions, and therefore a thorough exploration of height ranges is suggested using the plot. regmethod specifies whether data is summarized by taking the mean or median of height ranges across scales or all data is used. regmethod "raw" is not recommended because the regression will give much more weight to the lower scales that include more points and likely underestimate \( D \).
Value

A value for fractal dimension, typically between 2 and 3 or a list if keep_data = TRUE.

Examples

dem <- habtools::dem_crop(horseshoe, x0 = -469, y0 = 1267, L = 2, plot = TRUE)
fd_hvar(dem, lvec = c(0.125, 0.25, 0.5, 1, 2))
fd_hvar(dem, regmethod = "mean", plot = TRUE, keep_data = TRUE)
fd_hvar(dem, regmethod = "median", plot = TRUE, keep_data = TRUE)
fd_hvar(dem)

fd_sd

Calculate fractal Dimension using the standard deviation method

Description

Calculate fractal Dimension using the standard deviation method

Usage

fd_sd(
  data,
  lvec,
  regmethod = "mean",
  keep_data = FALSE,
  plot = FALSE,
  parallel = FALSE,
  ncores = (parallel::detectCores() - 1)
)

Arguments

data Digital elevation model of class RasterLayer.
lvec Vector of scales to use for calculation.
regmethod Method to use for linear regression between scale (lvec) and height range. One of raw (all data), mean (default) median or ends (minimum and maximum scale only)
keep_data Logical. Keep the data used for fd calculation? Defaults to FALSE.
plot Logical. Show plot of scales relative to data?
parallel Logical. Use parallel processing? Note: parallel must be installed.
ncores Number of cores to use when parallel = TRUE.
Details

Calculates fractal dimension using the standard deviation method, an analogue of the variation method, but using the standard deviation in height per grid cell instead of the full height range. If lvec is not specified, a default based on resolution and extent will be used. A rule of thumb is that lvec should range at least an order of magnitude. However, large ranges also average-out fractal dimension of a surface that might have phase transitions, and therefore a thorough exploration of height ranges is suggested using the plot. regmethod specifies whether data is summarized by taking the mean or median of height ranges across scales or all data is used. regmethod "raw" is not recommended because the regression will give much more weight to the lower scales that include more points and likely underestimate D.

Value

A value for fractal dimension, typically between 2 and 3 or a list if keep_data = TRUE.

Examples

dem <- habtools::dem_crop(horseshoe, x0 = -469, y0 = 1267, L = 2, plot = TRUE)
fd_sd(dem, lvec = c(0.125, 0.25, 0.5, 1, 2))

horseshoe  Horseshoe reef

Description

A digital elevation model (DEM) of a reef patch in the Great Barrier Reef.

Usage

horseshoe

Format

A 800 by 800 digital elevation model (of class RasterLayer).

Values  depth
Resolution  0.01 m
Extent  8 m ...

Examples

raster::plot(habtools::horseshoe)
**hr**  
*Calculate height range*

**Description**  
Calculates the distance between the lowest and highest point in a 3D object.

**Usage**  
hr(data)

**Arguments**  
- **data**  
  A RasterLayer or mesh3d object.

**Value**  
Value of height range.

**Examples**  
- # for a DEM  
  hr(horseshoe)
- # for a 3D mesh  
  hr(mcap)

---

**hvar**  
*Calculate height variation in cells at different scales*

**Description**  
This is a helper function used for calculating fractal dimension using the height variation and standard deviation methods.

**Usage**  
hvar(  
  data,  
  lvec = NULL,  
  parallel = FALSE,  
  ncores = (parallel::detectCores() - 1)
)
**mcap**

**Arguments**

- **data**: Digital elevation model of class `RasterLayer`.
- **lvec**: Scales to use for calculation.
- **parallel**: Logical. Use parallel processing? Note: parallel must be installed.
- **ncores**: Number of cores to use when parallel = TRUE.

**Value**

A data.frame containing height ranges of cells at different scales.

**Examples**

```
hvar(horseshoe, lvec = c(1, 2, 4, 8))
```

---

**mcap**  
*Montipora capitata*

**Description**

A laser scan of a coral colony.

**Usage**

```
mcap
```

**Format**

mesh3d object with 5568 vertices, 10939 triangles.

**Examples**

```
library(rgl)
plot3d(mcap)
```
Description
A remeshed version of mcap with resolution = 0.005.

Usage
mcap2

Format
mesh3d object.

Examples
library(rgl)
plot3d(mcap2)

Description
Turns a 3D Mesh file into an xy data frame.

Usage
mesh_to_2d(mesh, L0 = NULL, plot = FALSE, silent = TRUE)

Arguments
mesh A mesh3d object.
L0 (Optional) The desired DEM resolution in same units at the 3D mesh.
plot logical. Plot the output?
silent logical. Defaults to not showing warnings.

Details
The function uses the vertices of the mesh object and projects them on the XY plane. Then, only points that define the perimeter of the shape are maintained.

Value
A data frame.
mesh_to_dem

Examples

mcap_2d <- mesh_to_2d(mcap, plot = TRUE)

geometry::polyarea(mcap_2d$x, mcap_2d$y) # area
planar(mcap)

perimeter(mcap_2d) # perimeter
circularity(mcap_2d) # circularity
fd_boxes(mcap_2d) # fractal dimension

---

mesh_to_dem

Transform 3D mesh to DEM

Description

Turns a 3D mesh file into a Digital Elevation Model (DEM) of class RasterLayer format.

Usage

mesh_to_dem(mesh, res, fill = TRUE)

Arguments

mesh A mesh3d object.
res (Optional) The desired DEM resolution in same units at the 3D mesh.
fill Logical. Fill NA values in raster with minimum value?

Details

The function rasterizes uses the vertices of the mesh file. If resolution is not given, it is calculated by finding the maximum nearest neighbor of vertices projected on the xy plane. fill is used when irregular 3D meshes result in NA values in raster cells. The default is to fill these cells with the minimum, non-NA raster value.

Value

A dem of class RasterLayer.

Examples

dem <- mesh_to_dem(mcap)
raster::plot(dem)

dem <- mesh_to_dem(mcap, res = 0.05)
raster::plot(dem)

# Don't fill empty raster cells
dem <- mesh_to_dem(mcap, res = 0.02, fill = FALSE)
raster::plot(dem)

define mesh_to_points

**Description**

Transform mesh to 3D point cloud

**Usage**

```r
mesh_to_points(mesh)
```

**Arguments**

- `mesh`: A triangular mesh of class `mesh3d`.

**Value**

A data frame with XYZ coordinates.

---

**mid_find**

*Find midpoint of a DEM*

**Description**

Find midpoint of a DEM

**Usage**

```r
mid_find(data)
```

**Arguments**

- `data`: A DEM in RasterLayer format.

**Value**

A data frame with x and y midpoints.

**Examples**

```r
mid_find(horseshoe)
```
packing

Calculate packing of 3D object

Description
The ratio of the surface area of the object and the surface area of the convex hull around the object.

Usage
packing(mesh)

Arguments
- mesh A triangular mesh of class mesh3d.

Value
Value of packing.

Examples
packing(mcap)

perimeter
Calculate perimeter of a 2D shape

Description
Calculates the perimeter of a 2D shape.

Usage
perimeter(data, keep_data = FALSE)

Arguments
- data A data frame with the first two columns ordered x and y coordinates.
- keep_data Logical. Keep lengths of all segments of the perimeter? Defaults to FALSE.

Value
The perimeter.
Examples

```r
mcap_2d <- mesh_to_2d(mcap)
plot(mcap_2d)

perimeter(mcap_2d)

r <- 1 # radius
circ <- sim_circle(r=r) # simulate xy coordinates for a circle of radius 1
plot(circ, asp=1)
perimeter(circ)

2 * pi * r # Note xy resolution affects output
```

---

**planar**

Calculates planar area of a mesh

Description

Calculates planar area of a mesh

Usage

```r
planar(mesh, L0, silent = FALSE)
```

Arguments

- `mesh` A triangular mesh of class mesh3d.
- `L0` Resolution of the planar area. Is set to the resolution of the mesh when left empty.
- `silent` Logical. Suppress messages and warnings?

Value

A value for planar area.

Examples

```r
planar(mcap)
```
rdh

Calculate rugosity, fractal dimension, and height for a DEM

Description

Calculate rugosity, fractal dimension, and height for a DEM

Usage

rdh(
  data,
  lvec,
  method_fd = "hvar",
  method_rg = "area",
  parallel = FALSE,
  ncores = (parallel::detectCores() - 1),
  ...
)

Arguments

data A dem of class RasterLayer.
lvec Scales to use for calculation.
method_fd method for the calculation of rugosity and fractal dimension. Can be "hvar",
  "sd", "cubes", or "area". Defaults to "hvar".
method_rg Method to be used for the rugosity calculation. Defaults to "area".
parallel Logical. Use parallel processing? Defaults to FALSE.
ncores Number of cores to use if parallel = TRUE.
... Additional arguments see fd().

Details

Uses area method for rugosity and hvar method for fractal dimension calculations as default.

Value

A dataframe with the three complexity metrics.

See Also

fd()
rg()
hr()
Examples

dem <- dem_sample(horseshoe, L = 1)
rdh(dem, lvec = c(0.125, 0.25, 0.5, 1))

Description

Calculates either rugosity, fractal dimension or height range based on the other two variables.

Usage

rdh_theory(R, D, H, L, L0)

Arguments

R, D, H  Two of the three variables to calculate the third.
L        Extent.
L0       Resolution.

Details

This function uses the geometric plane equation from Torres-Pulliza et al. (2020) to calculate one of rugosity, fractal dimension or height range based on the other two variables.

Value

A value corresponding one of the three variables not given to the function.

References


Examples

rdh_theory(R=4, H=1, L=1, L0=0.01)
rdh_theory(D=2.36928, H=1, L=1, L0=0.01)
rdh_theory(D=2.36928, R=4, L=1, L0=0.01)
Calculate rugosity

Description

Rugosity is defined as the surface area divided by the planar area. For digital elevation models, there are two methods: "hvar" and "area". The "hvar" method for calculating rugosity is described in Torres-Pulliza et al. (2004) and is based on height variations. The "area" method uses the `sp::surfaceArea()` function and is detailed in Jenness (2004). Method is ignored if data is a mesh3D object. In that case the function uses `Rvcg::vcgArea()` to calculate surface area of a triangular mesh of class mesh3d.

Usage

```r
rg(data, L0, method = "area", parallel = FALSE, ncores = (parallel::detectCores() - 1))
```

Arguments

- `data` Digital elevation model of class RasterLayer or a triangular mesh of class mesh3d.
- `L0` Grain or resolution of calculation.
- `method` If data is a RasterLayer methods "hvar" or "area" are allowed. Defaults to "hvar".
- `parallel` Logical. Use parallel processing? Defaults to FALSE.
- `ncores` Number of cores to use if parallel = TRUE. (Defaults to number of available cores - 1)

Value

Rugosity value

References


Examples

rg(horseshoe, L0 = 0.1)
rg(mcap, L0 = 0.01)

---

sa_triangle  
**Calculate surface area of triangle**

Description

Calculates the surface area of a triangle based on a set of XYZCoords.

Usage

`sa_triangle(XYZcoords)`

Arguments

`XYZcoords`  
A data frame with XYZ coordinates of three points in following order: X1,X2,X3,Y1,Y2,Y3,Z1,Z2,Z3.

Value

The surface area of the triangle.

Examples

`sa_triangle(c(X1 = 1, X2 = 2, X3 = 3 , Y1 = 1, Y2 = 2, Y3 = 1, Z1 = 1, Z2 = 1, Z3 = 1))`

---

scale_area  
**Re-scale mesh based on a fixed area**

Description

Re-scale mesh based on a fixed area

Usage

`scale_area(mesh, target_area = 1)`

Arguments

`mesh`  
A triangular mesh of class mesh3d.

`target_area`  
The target area of the scaled 3D mesh. Defaults to 1.

Value

A mesh with area = target_area (1 as default).
**scale_volume**

Re-scale mesh based on a fixed volume of 1

**Description**

Re-scale mesh based on a fixed volume of 1

**Usage**

```r
scale_volume(mesh)
```

**Arguments**

- `mesh`: A triangular mesh of class `mesh3d`.

**Value**

A mesh with volume = 1.

**Examples**

```r
Rvcg::vcgVolume(mcap)
mcap_scaled <- scale_volume(mcap)
Rvcg::vcgVolume(mcap_scaled)
```

**set_origin**

Set the origin of a mesh

**Description**

Transforms XYZ coordinates relative to a chosen origin

Transforms coordinates so that the origin lies at the reference vertex (defaults to the minimum of x, y, and z coordinates).

**Usage**

```r
set_origin(mesh, reference = NULL)
```
Arguments

- mesh: A triangular mesh of class mesh3d.
- reference: Vector containing coordinates of the reference vertex. If left empty, this will default to the minimum of x, y, and z.

Value

- mesh3d object

Examples

```r
mesh <- set_origin(mcap)
```

---

Simulate a circle

**Description**

Simulates xy coordinates for a circle of given radius. Created for package testing purposes, but might be useful for others.

**Usage**

```r
sim_circle(r = 1, n = 100, mid = c(0, 0))
```

**Arguments**

- **r**: Radius of the circle (default 1).
- **n**: Number of xy coordinates defining the circle (default 100).
- **mid**: Mid point of the circle (default 0, 0).

**Value**

- A data frame of n xy-coordinates.

**Examples**

```r
circ <- sim_circle()
plot(circ)
circularity(circ)
perimeter(circ)
```
Simulates a fractal DEM

Description

Simulates z-values based on the Diamond-square algorithm.

Usage

```r
sim_dem(
  L,
  smoothness,
  H,
  R,
  plot = FALSE,
  prop = 0.1,
  n = 100,
  method = "area",
  parallel = FALSE
)
```

Arguments

- `L`: The extent.
- `smoothness`: A value between 0.0 and 1.0 (lower values produce rougher DEM).
- `H`: Desired height range (optional).
- `R`: Desired rugosity value (optional).
- `plot`: Logical. Plot the simulated DEM during simulation? Only relevant if R is provided.
- `prop`: Proportion of cells that undergo smoothing at each iteration when R is provided.
- `n`: Number of iterations to try and reach desired R. Recommended to adapt R and H instead of increasing n if simulation fails.
- `method`: The method to be used for rugosity calculation in case R is given. Can be "hvar" or "area".
- `parallel`: Logical. Use parallel processing? Defaults to FALSE. Only relevant if method = "hvar".

Details

Warning: this function gets slow for n > 128. If H is provided, the simulated DEM is rescaled based on the value for H. If R is provided, a DEM is simulated using the same algorithm based on R, H, and the predicted D based on `rdh_theory()`, while smoothness is ignored. From that first simulated DEM, R is calculated and the DEM undergoes smoothing at each iteration until the rugosity approximates the inputted R. Argument prop defined the proportion of random cells of the DEM that are smoothed by averaging the z values of cell and neighboring cells at each iteration.
Caution: When R is provided, the DEM may become increasingly less fractal as it is modified at each iteration.

Value

Digital elevation model of class RasterLayer.

Examples

library(raster)
dem <- sim_dem(L = 32, smoothness = 0.5)
plot(dem)
dem <- sim_dem(L = 32, smoothness = 0.2, H = 20)
plot(dem)

sma

Calculate second moment of area

Description

Calculates the 2nd moment of surface area about the origin by multiplying the surface area of each triangle in the mesh by its distance from the origin (should be set to the attachment point of the mesh). The sum of these values is the 2nd moment of area. This metric is size-dependent so to compare moments in terms of shape only, set scale = TRUE.

Usage

sma(mesh, axis = "z", scale = FALSE, origin = TRUE)

Arguments

mesh A triangular mesh of class mesh3d.
axis The axis along which to calculate the second moment of area. z is the default.
scale Logical. Scale the object to have a volume = 1? Default = FALSE
origin Logical. Set the origin to the bottom left corner of bounding box? Default = TRUE.

Value

SMA value.

Examples

sma(mcap)
sma(mcap, scale = TRUE)
smv

Calculate second moment of volume

Description
Calculates the 2nd moment of volume (SMV) by multiplying the volume of each triangle in the mesh by its centroids’ distance from the origin (should be set to the attachment point of the mesh). The sum of these values is the 2nd moment of volume. Axis is z by default, meaning it will calculate the vertical second moment, but this can be changed if needed. This metric is size-dependent so to compare moments in terms of shape only, set scale = TRUE.

Usage
smv(mesh, axis = "z", scale = FALSE, origin = TRUE)

Arguments
- mesh: A triangular mesh of class mesh3d.
- axis: The axis along which to calculate the second moment of volume z is the default.
- scale: Logical. Scale the object to have a volume = 1? Default = TRUE.
- origin: Logical. Set the origin to the bottom left corner of bounding box? Default = FALSE

Value
SMV value.

Examples
smv(mcap)

sphericity
Calculate sphericity of a 3D object

Description
Calculates the ratio of the surface area of a sphere with the same volume as the object and the surface area of the object.

Usage
sphericity(mesh)

Arguments
- mesh: A triangular mesh of class mesh3d.
Value

Sphericity value.

See Also

circularity()

Examples

sphericity(mcap)

data DEM in RasterLayer format, mesh3d object or data frame with xy coordinates.

Value

Surface area value.

References


Examples

surface_area(mcap)
surface_area(horseshoe)
surface_area(mesh_to_2d(mcap))
svol_triangle

**svol_triangle**  Calculate signed volume of triangle

**Description**
Calculates the signed volume of a triangle based on a set of XYZCoords. Signed volume means that volumes can take on a negative value depending on whether the surface normal of the triangle is facing towards or away from the origin. When all positive and negative volumes are integrated across the entire mesh, these values cancel out so that the final volume is an approximation of the total volume of the mesh.

**Usage**
svol_triangle(XYZCoords)

**Arguments**
- **XYZCoords**  A dataframe with XYZ coordinates of three points in following order: X1,X2,X3,Y1,Y2,Y3,Z1,Z2,Z3

**Value**
Value for the signed volume of a triangle.

**Examples**
svol_triangle(c(X1 = 1, X2 = 2, X3 = 3, Y1 = 1, Y2 = 2, Y3 = 1, Z1 = 1, Z2 = 1, Z3 = 1))
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