

# Package: rTLS (via r-universe)

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**Type** Package

**Title** Tools to Process Point Clouds Derived from Terrestrial Laser Scanning

**Version** 0.2.6.1

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**Description** A set of tools to process and calculate metrics on point clouds derived from terrestrial LiDAR (Light Detection and Ranging; TLS). Its creation is based on key aspects of the TLS application in forestry and ecology. Currently, the main routines are based on filtering, neighboring features of points, voxelization, canopy structure, and the creation of artificial stands. It is written using data.table and C++ language and in most of the functions it is possible to use parallel processing to speed-up the routines.

**License** GPL (>= 3)

**URL** <https://github.com/Antguz/rTLS>

**BugReports** <https://github.com/Antguz/rTLS/issues>

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rTLS-package

*rTLS: Tools to Process Point Clouds Derived From Terrestrial Laser Scanning*

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## Description

rTLS is a package that compiles a set of tools to process and calculate metrics on point clouds derived from terrestrial LiDAR (Light Detection and Ranging). Its creation is based on key aspects of the TLS application in forestry. Currently, the main routines are based on filtering, neighboring features of points, voxelization, optimal sphere or voxel size, and the creation of artificial stands. rTLS is written using data.table and C++ language and in most of the functions it is possible to use parallel processing to speed-up the routines.

## Author(s)

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## See Also

Useful links:

- <https://github.com/Antguz/rTLS>
- Report bugs at <https://github.com/Antguz/rTLS/issues>

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artificial\_stand

*Artificial Forest Stand*

---

## Description

Create an artificial forest stand of a given area using tree point clouds.

**Usage**

```

artificial_stand(
  files,
  n.trees,
  dimension,
  coordinates = NULL,
  sample = TRUE,
  replace = TRUE,
  overlap = NULL,
  rotation = TRUE,
  degrees = NULL,
  n_attempts = 100,
  progress = TRUE,
  plot = TRUE,
  ...
)

```

**Arguments**

files	A character vector describing the file name or path of the tree point cloud to use. Those files most contain three columns representing the *XYZ* coordinates of a given point cloud.
n.trees	A positive numeric vector describing the number of point clouds to use.
dimension	A positive numeric vector of length two describing the width and length of the future forest stand.
coordinates	A data.table of two columns and with n rows equal to n.trees describing the basal *XYZ* coordinates of the point clouds in the future stand. If NULL, it uses random basal coordinates based on stand dimension. NULL as default.
sample	Logical. If TRUE, it performs a sample of the files to determine the order to build the artificial stand. If FALSE, it use the file order described in files. TRUE as default.
replace	Logical. If TRUE, it performs a sample selection with a replacement if sample = TRUE to determine the order to build the artificial stand. Useful if the n.trees is lower than length(files). TRUE as default.
overlap	A positive numeric vector between 0 and 100 describing the overlap percentage of a given the tree crowns in the future forest stand. If NULL, the degree of overlap is not controlled.
rotation	Logical. If TRUE, it performs a rotation in yaw axis of the point cloud. TRUE as default.
degrees	A positive numeric vector describing the degrees of rotation of the point clouds in the future stand. The length(degree) should be the same as n.trees. If NULL, it creates random degrees of rotation for each n.trees.
n_attempts	A positive numeric vector of length one describing the number of attempts to provide random coordinates until a tree met the overlap criteria. This needs to be used if coordinate = NULL and overlap != NULL. n_attempts = 100 as default.

progress	Logical, if TRUE displays a graphical progress bar. TRUE as default.
plot	Logical. If TRUE, it provides visual tracking of the distribution of each tree in the artificial stand. This can not be exported as a return object.
...	Parameters passed to <code>fread</code> for the reading of files.

### Details

When `coordinates = NULL`, `artificial_stand` adds, in sequence, random coordinates to each files in the future stand based on the crown area overlap. That is, first a tree from files is randomly located within the stand dimation, then a second tree from files will be located in the future stand based on the crown area overlap from the previous tree, and so on. If during the random location a given tree does not meet the requirements of `overlap`, new random coordinates will be provided until the requirements are met.

Since `artificial_stand` will try to add tree to the stand until the requirements are met, this could lead to an infinite loop if the stand dimation is small or if the trees on files are large or many `n.trees`. Therefore, the use of `n_attempts` is recommended to avoid this scenario.

### Value

A list which contain a `data.table` (Trees) with the information of the point clouds used and their current coordinates in the stand, and another `data.table` with that compile all the point clouds used.

### Author(s)

J. Antonio Guzmán Q.

### See Also

[voxels\\_counting](#)

### Examples

```
## Import an example point cloud
path <- system.file("extdata", "pc_tree.txt", package = "rTLS")

#Creates a stand of 4 trees with 10% of overlap
files <- rep(path, 4)
artificial_stand(files, n.trees = 4, dimension = c(15, 15), overlap = 10)

#Creates a stand of 4 trees with their locations
location <- data.table(X = c(5, 10, 10, 5), Y = c(5, 5, 10, 10))
artificial_stand(files, n.trees = 4, dimension = c(15, 15), coordinates = location)
```

---

canopy\_structure      *Canopy Structure*

---

### Description

Estimates the canopy structure from a discrete returns scan from different TLS.

### Usage

```
canopy_structure(
  TLS.type,
  scan,
  zenith.range,
  zenith.rings,
  azimuth.range,
  vertical.resolution,
  TLS.pulse.counts,
  TLS.resolution = NULL,
  TLS.coordinates = c(0, 0, 0),
  TLS.frame = NULL,
  TLS.angles = NULL,
  threads = 1
)
```

### Arguments

TLS.type	A character describing is the TLS used. It most be one of "single" return, "multiple" return, or "fixed.angle" scanner.
scan	If TLS.type is equal to "single" or "fixed.angle", a data.table with three columns describing *XYZ* coordinates of the discrete return. If TLS.type is equal to "multiple", a data.table with four columns describing *XYZ* coordinates and the target count pulses. Currently, "fixed.angle" present errors, use with discretion.
zenith.range	If TLS.type is equal to "single" or "multiple", a numeric vector of length two describing the min and max range of the zenith angle to use. Theoretically, the max range should be lower than 90 degrees.
zenith.rings	If TLS.type is equal to "single" or "multiple", a numeric vector of length one describing the number of zenith rings to use between zenith.range. This is used to estimate the frequency of laser shots from the scanner and returns in scan. If TLS.type = "fixed.angle", zenith.rings = 1 be default.
azimuth.range	A numeric vector of length two describing the range of the azimuth angle to use. Theoretically, it should be between 0 and 360 degrees.
vertical.resolution	A numeric vector of length one describing the vertical resolution to extract the vertical profiles. Low values lead to more variable profiles. The scale used needs to be in congruence with the scale of scan.

TLS.pulse.counts	If TLS.type is equal to "single" or "multiple", a numeric vector of length two describing the horizontal and vertical pulse counts of the scanner. If TLS.type is equal to "fixed.angle", a numeric vector of length one describing the horizontal pulse counts resolution. Preferred parameter over TLS.resolution to estimate the number of pulses.
TLS.resolution	If TLS.pulse.counts = NULL, the code use the angles resolution to estimate the pulse counts in a given TLS.frame. If TLS.type is equal to "single" or "multiple", a numeric vector of length two describing the horizontal and vertical angle resolution of the scanner. If TLS.type is equal to "fixed.angle", a numeric vector of length one describing the horizontal angle resolution.
TLS.coordinates	A numeric vector of length three describing the scanner coordinates within scan. It assumes that the coordinates are $c(X = 0, Y = 0, Z = 0)$ for default.
TLS.frame	If TLS.type is equal to "single" or "multiple", a numeric vector of length four describing the min and max of the zenith and azimuth angle of the scanner frame. If TLS.type = "fixed.angle", a numeric vector of length three describing the fixed zenith angle and the min and max of the azimuth angle of the scanner frame. If NULL, it assumes that a complete hemisphere ( $c(\text{zenith.min} = 0, \text{zenith.max} = 90, \text{azimuth.min} = 0, \text{azimuth.max} = 360)$ ), or a cone projection ( $c(\text{zenith} = 57.5, \text{azimuth.min} = 0, \text{azimuth.max} = 360)$ ) depending on TLS.type.
TLS.angles	A numeric vector of length three describing the roll (*X*), pitch (*Y*), and yaw (*Z*) angles of the scanner during the scan. If NULL, it assumes that there is no need to to correction of angles. This needs to be used if TLS.type is equal to "single" or "multiple", since it assumes that "fixed.angle" scanner is previously balanced. NULL as default.
threads	An integer specifying the number of threads to use. Experiment to see what works best for your data on your hardware.

## Details

Since scan describes discrete returns measured by the TLS, canopy\_structre first simulates the number of pulses emitted based on Danson et al. (2007). The simulated pulses are created based on the TLS properties (TLS.pulse.counts, TLS.resolution, TLS.frame) assuming that the scanner is perfectly balance. Then these pulses are rotated (`rotate3D`) based on the TLS.angles roll, pitch, and yaw, and move to TLS.coordintates to simulate the positioning of the scanner during the scan. Rotated simulated-pulses of interest and scan returns are then extracted based on the zenith.range and azimuth.range for a given number of zenith.rings, azimuth.rings and vertical profiles. The probability of gap (Pgap) is then estimated using the frequency of pulses and returns. For TLS.type = "multiple", the frequency of returns is estimated using the sum of  $1/\text{target count}$  following Lovell et al. (2011).

Using the Pgap estimated per each zenith ring and vertical profile, canopy\_structure then estimates the accumulative  $L(z)$  profiles based on the closest zenith ring to 57.5 (hinge region) and, if TLS.type = "fixed.angle", the  $f(z)$  or commonly named PAVD based on the ratio of the derivative of  $L(z)$  and height ( $z$ ) following Jupp et al. 2009 (Equation 18). If TLS.type is equal to "single" or "multiple", canopy\_structure also estimates the normalized average weighted  $L/LAI$ , and then

PAVD based on the L (hinge angle) at the highest height (LAI) and the ratio between the derivative of L/LAI (average weighted) and the derivative of z (Jupp et al. 2009; Equation 21).

Jupp et al. 2009 excludes the zero zenith or first ring to conduct the average weighted L/LAI estimations, `canopy_structure` does not exclude these sections since it depends on the regions of interest of the user. Therefore, user should consider this difference since it may introduce more variability to profile estimations.

### Value

For any TLS.type, it returns a data.table with the height profiles defined by `vertical.resolution`, the gap probability based on the `zenith.range` and `zenith.rings`, and the accumulative L(z) profiles based on the closest zenith ring to 57.5 degrees (hinge angle). If TLS.type is equal to "fixed.angle", it returns f(z) or commonly named PAVD based on the ratio of the derivative of L(z) and the derivative of height (z). If TLS.type is equal to "single" or "multiple", it returns the normalized average weighting L/LAI, and PAVD: based on the L (hinge angle) at the highest height and the ratio between the derivative of L/LAI average weighted and the derivative of z.

### Author(s)

J. Antonio Guzmán Q.

### References

- Danson F.M., Hetherington D., Morsdorf F., Koetz B., Allgower B. 2007. Forest canopy gap fraction from terrestrial laser scanning. *IEEE Geosci. Remote Sensing Letters* 4:157-160. doi: 10.1109/LGRS.2006.887064
- Lovell J.L., Jupp D.L.B., van Gersel E., Jimenez-Berni J., Hopkinson C., Chasmer L. 2011. Foliage profiles from ground based waveform and discrete point LiDAR. In: *SilviLaser 2011*, Hobart, Australia, 16–20 October 2011.
- Jupp D.L.B., Culvenor D.S., Lovell J.L., Newnham G.J., Strahler A.H., Woodcock C.E. 2009. Estimating forest LAI profiles and structural parameters using a ground-based laser called "Echidna®". *Tree Physiology* 29(2): 171-181. doi: 10.1093/treephys/tpn022

### Examples

```
data(TLS_scan)
#Using a multiple return file
#Select the four columns required
TLS_scan <- TLS_scan[, 1:4]

#This will take a while#
canopy_structure(TLS.type = "multiple",
                scan = TLS_scan,
                zenith.range = c(50, 70),
                zenith.rings = 4,
                azimuth.range = c(0, 360),
                vertical.resolution = 0.25,
                TLS.pulse.counts = c(2082, 580),
                TLS.frame = c(30, 130.024, 0, 359.90),
```

```

        TLS.angles = c(1.026, 0.760, -110.019))

#Using a single return file

data(TLS_scan)
#Subset to first return observations
TLS_scan <- TLS_scan[Target_index == 1, 1:3]

#This will take a while#
canopy_structure(TLS.type = "single",
                 scan = TLS_scan,
                 zenith.range = c(50, 70),
                 zenith.rings = 4,
                 azimuth.range = c(0, 360),
                 vertical.resolution = 0.25,
                 TLS.pulse.counts = c(2082, 580),
                 TLS.frame = c(30, 130.024, 0, 359.90),
                 TLS.angles = c(1.026, 0.760, -110.019))

```

---

cartesian\_to\_polar      *Cartesian to Polar Coordinates*

---

## Description

Convert from East-North-Up cartesian coordinates to polar coordinates.

## Usage

```
cartesian_to_polar(cartesian, anchor = c(0, 0, 0), digits = NULL)
```

## Arguments

cartesian	A data.table with three columns describing the *XYZ* coordinates of a point cloud.
anchor	A numeric vector of length three which describe the *XYZ* anchor coordinate for reference to get the polar coordinates. It assumes that the reference coordinates are $c(X = 0, Y = 0, Z = 0)$ as default.
digits	A numeric vector of length 1 describing the decimal numbers to <a href="#">round</a> the zenith and azimuth angles. If NULL, <a href="#">round</a> does not apply. NULL as default.

## Details

It assumes that the positive \*Z\* axis is the reference vector for the zenith angle. Likewise, it assumes that the \*Y\* axis is the north-south direction (positive to negative) for the azimuth angle. If a point from cartesian presents the same \*XY\* coordinates than anchor, angles returns NA.

**Value**

A data.table with the zenith and azimuth angles (degrees), and the distance to the anchor coordinate.

**Author(s)**

J. Antonio Guzmán Q.

**See Also**

[polar\\_to\\_cartesian](#)

**Examples**

```
data(pc_tree)
cartesian_to_polar(pc_tree)
anchor <- c(1, 1, 1)
cartesian_to_polar(pc_tree, anchor)
```

---

circleRANSAC

*Adaptive RANSAC Circle Fitting*

---

**Description**

Adaptive random sample consensus for circle fitting.

**Usage**

```
circleRANSAC(
  cloud,
  fpoints,
  pconf,
  poutlier,
  max_iterations,
  threads = 1L,
  plot = TRUE
)
```

**Arguments**

cloud	A data.table with *XY* coordinates in the first two columns.
fpoints	A numeric vector between 0 and 1 representing the fraction of point samples that will be used during each iteration.
pconf	A numeric vector between 0 and 1 describing the confidence threshold to consider a point in a given fitted circle outlier or inlier.

poutlier	A numeric vector of length two describing the proportion of outliers to consider inside or outside of the pconf threshold.
max_iterations	An integer specifying the number of iterations. If NULL, the number of iterations are automatically estimated using pconf, 1 - poutlier, and 1 - fpoints; see details.
threads	An integer specifying the number of threads to use for parallel processing. Experiment to see what works best for your data on your hardware.
plot	Logical. If TRUE, it provides visual representation of the fitted circle.

### Value

A data.table with the \*XY\* coordinate information of the circle center, the radius, the error based on the least squares fit, and the proportion of inliers.

### Author(s)

J. Antonio Guzmán Q.

### See Also

[tree\\_metrics](#), [trunk\\_volume](#)

### Examples

```
#Point cloud
data("pc_tree")

#Subset region at at breast height
sub <- pc_tree[between(Z, 1.25, 1.35),]

#Fit circle
circleRANSAC(sub, fpoints = 0.2, pconf = 0.95, poutlier = c(0.5, 0.5), max_iterations = 100)
```

---

euclidean\_distance      *Euclidean Distance Between 3D points*

---

### Description

Estimate the distance between a point and a group of point.

### Usage

```
euclidean_distance(point, cloud, threads = 1L)
```

**Arguments**

point	A numeric vector of length three describing the *XYZ* coordinates.
cloud	A data.table with *XYZ* coordinates in the first three columns representing a point cloud.
threads	An integer specifying the number of threads to use for parallel processing. Experiment to see what works best for your data on your hardware.

**Value**

A numeric vector describing of point to each row of cloud.

**Author(s)**

J. Antonio Guzmán Q.

**Examples**

```
data("pc_tree")  
euclidean_distance(point = c(0, 0, 0), pc_tree)
```

---

filter

*Filtering of Point Clouds*

---

**Description**

Filtering of point clouds using different methods

**Usage**

```
filter(  
  cloud,  
  method,  
  radius,  
  min_neighbours,  
  k,  
  nSigma,  
  edge_length,  
  distance = "euclidean",  
  threads = 1L,  
  verbose = FALSE,  
  progress = FALSE,  
  ...  
)
```

**Arguments**

<code>cloud</code>	A <code>data.table</code> contain three columns representing the *XYZ* coordinates.
<code>method</code>	A filtering method to use. It must be "SOR", "min_neighbors", or "min_neighbours".
<code>radius</code>	A numeric vector representing the radius of the sphere to consider. This needs to be used if <code>method = "voxel_center"</code> .
<code>min_neighbours</code>	An integer representing the minimum number of neighbors to keep a given point. This needs to be used if <code>method = "min_n"</code> .
<code>k</code>	An integer vector representing the number of neighbors to consider. This needs be used if <code>method = "SOR"</code> .
<code>nSigma</code>	A numeric vector representing the standard deviation multiplier. This needs to be used if <code>method = "SOR"</code> .
<code>edge_length</code>	A positive numeric vector with the voxel-edge length for the x, y, and z coordinates. This needs to be used if <code>method = "voxel_center"</code> .
<code>distance</code>	Type of distance to calculate. "euclidean" as default. Look <code>hnsw_knn</code> for more options.
<code>threads</code>	An integer specifying the number of threads to use for parallel processing. Experiment to see what works best for your data on your hardware.
<code>verbose</code>	If TRUE, log messages to the console.
<code>progress</code>	If TRUE, log a progress bar when <code>verbose = TRUE</code> . Tracking progress could cause a small overhead.
<code>...</code>	Arguments passed to <code>hnsw_build</code> and <code>hnsw_search</code> .

**Value**

A `data.table` with the filtered points

**Author(s)**

J. Antonio Guzmán Q.

**Examples**

```
#Load data
data("pc_tree")

#Move pc_tree for comparison
pc_compare <- pc_tree
pc_compare$X <- pc_compare$X - 7

#SOR filter
r1 <- filter(pc_tree, method = "SOR", k = 30, nSigma = 1)
rgl::plot3d(r1, col = "red") #Filter
rgl::points3d(pc_compare, col = "black") #Original

#min_neighbours filter
```

```

r2 <- filter(pc_tree, "min_neighbors", radius = 0.02, min_neighbours = 20)
rgl::plot3d(r2, col = "red") #Filter
rgl::points3d(pc_compare, col = "black") #Original

#voxel_center filter
r3 <- filter(pc_tree, method = "voxel_center", edge_length = 0.1)
rgl::plot3d(r3, col = "red") #Filter
rgl::points3d(pc_compare, col = "black") #Original

```

---

geometry\_features      *Geometry features of Neighboring Points.*

---

## Description

Estimate geometry features of neighboring points in a cloud.

## Usage

```

geometry_features(
  cloud,
  method,
  radius,
  k,
  max_neighbour,
  distance = "euclidean",
  target = FALSE,
  threads = 1L,
  verbose = FALSE,
  progress = TRUE,
  ...
)

```

## Arguments

cloud	A data.table with *XYZ* coordinates in the first three columns.
method	A character string specifying the method to estimate the neighbors. It must be one of "radius_search" or "knn".
radius	A numeric vector representing the radius for search to consider. This needs be used if method = "radius_search".
k	An integer vector representing the number of neighbors to consider. This needs be used if method = "knn".
max_neighbour	An integer specifying the maximum number of points to look around each query point for a given radius. This needs be used if method = "radius_search".
distance	Type of distance to calculate. "euclidean" as default. Look hnsw_knn for more options.

target	Logic. If TRUE, it consider the each target point for the calculations of geometry features.
threads	An integer specifying the number of threads to use for parallel processing. Experiment to see what works best for your data on your hardware.
verbose	If TRUE, log messages to the console.
progress	If TRUE, log a progress bar when verbose = TRUE. Tracking progress could cause a small overhead.
...	Arguments passed to hnsw_build and hnsw_search.

### Details

The function returns the geometry features of the neighboring points of a given point in `cloud`. Geometry features are represented by the relative values of the eigenvalues derived from a covariance matrix of the neighboring points. Geometry features are not estimated on target points with less than 3 neighboring points.

### Value

A array describing the point of the cloud in rows, the relative eigenvalues in columns, and the radius or k per slide. If `method = "radius_search"`, it add in the first column the number of neighboring points.

### Author(s)

J. Antonio Guzmán Q.

### Examples

```
#Create cloud
example <- data.table(X = runif(200, min=0, max=10),
                    Y = runif(200, min=0, max=10),
                    Z = runif(200, min=0, max=10))

#Using knn method with two different k
k_test <- c(5, 10)
geometry_features(example, method = "knn", k = k_test)

#Using radius search method with two different radius
radius_test <- c(3, 4)
geometry_features(example, method = "radius_search", radius = radius_test, max_neighbour = 200)
```

---

knn *K Nearest Neighbors*


---

**Description**

Adapted K nearest neighbors based on RcppHNSW

**Usage**

```
knn(
  query,
  ref,
  k,
  distance = "euclidean",
  same = FALSE,
  threads = 1L,
  verbose = FALSE,
  progress = FALSE,
  ...
)
```

**Arguments**

query	A data.table containing the set of query points where each row represent a point and each column a given coordinate.
ref	A numeric containing the set of reference points where each row represent a point and each column a given coordinate.
k	An integer describing the number of nearest neighbors to search for.
distance	Type of distance to calculate. "euclidean" as default. Look hnsw_knn for more options.
same	Logic. If TRUE, it delete neighbors with distance of 0, useful when the k search is based on the same query.
threads	An integer specifying the number of threads to use for parallel processing. Experiment to see what works best for your data on your hardware.
verbose	If TRUE, log messages to the console.
progress	If TRUE, log a progress bar when verbose = TRUE. Tracking progress could cause a small overhead.
...	Arguments passed to hnsw_build and hnsw_search.

**Details**

This function is based on hnswlib C++ library (Malkov & Yashunin 2016) and its bindings for R (RcppHNSW; Melville 2020) for a fast estimation of neighbors points. It is adapted to simplify the workflow within rTLS. If you use this function, please consider cite the C++ library and RcppHNSW package.

**Value**

A data.table with three columns describing the indices of the query, ref, and k neighbors and the distances.

**Author(s)**

J. Antonio Guzmán Q.

**References**

Malkov, Y. A., & Yashunin, D. A. (2016). Efficient and robust approximate nearest neighbor search using Hierarchical Navigable Small World graphs. arXiv preprint arXiv:1603.09320.

**See Also**

[radius\\_search](#)

**Examples**

```
#Point cloud
data("pc_tree")

#knn search using k = 3
knn(pc_tree, pc_tree, k = 3, same = TRUE)
```

---

line\_AABB

*Line-AABB*

---

**Description**

Intersection of a line by an Axis-Aligned Bounding Box.

**Usage**

```
line_AABB(orig, end, AABB_min, AABB_max)
```

**Arguments**

orig	A data.table with the describing *XYZ* coordinates of the the start path of a line.
end	A data.table with the describing *XYZ* coordinates of the the end path of a line.
AABB_min	A numeric vector with the minimum *XYZ* coordinates of the AABB
AABB_max	A numeric vector with the maximum *XYZ* coordinates of the AABB.

## Details

The interaction of a line with a AABB may result in five scenarios: i) the line is not intercepted by a AAABB (0), ii) the origin and end of the line falls within the AABB (1), iii) the origin point of the line falls within the AABB both not the end point (2), iv) the end point of the line falls within the AABB both not the origin point (3), and v) the line is intercepted by the AABB (4).

## Value

An numeric vector of length two, describing if the line was intercepted or not, and the length of the intercepted line within in the AABB. See details.

## Author(s)

J. Antonio Guzmán Q.

## See Also

[lines\\_interception](#), [voxels](#),

## Examples

```
#Create origins and end paths
orig <- data.table(X = c(0, 0, 0, 0, 0),
                  Y = c(-0.45, -0.25, 0, 0.25, 0.45),
                  Z = c(-1, -0.25, 0, -1, -1))

end <- data.table(X = c(0, 0, 0, 0, 0),
                 Y = c(-0.45, -0.25, 0, 0.25, 0.45),
                 Z = c(-0.75, 0.25, 1, 0, 1))

#Create the AABB
AABB <- matrix(c(0, 0, 0), ncol = 3)
edge_length <- c(1, 1, 1)

AABB_min <- c(AABB[1, 1] - edge_length[1]/2,
              AABB[1, 2] - edge_length[2]/2,
              AABB[1, 3] - edge_length[3]/2)

AABB_max <- c(AABB[1, 1] + edge_length[1]/2,
              AABB[1, 2] + edge_length[2]/2,
              AABB[1, 3] + edge_length[3]/2)

#Plot
cube <- rgl::cube3d()
cube <- rgl::scale3d(cube, edge_length[1]/2,
                    edge_length[2]/2,
                    edge_length[3]/2)
box <- rgl::translate3d(cube, AABB[1, 1], AABB[1, 2], AABB[1, 3])
rgl::shade3d(box, col= "green", alpha = 0.6)
rgl::points3d(orig, size = 4, col = "black")
rgl::points3d(end, size = 4, col = "red")
```

```
#Line no intercepted
rgl::lines3d(c(orig[1, 1], end[1, 1]),
             c(orig[1, 2], end[1, 2]),
             c(orig[1, 3], end[1, 3]), col = "grey")

line_AABB(orig[1,], end[1,], AABB_min, AABB_max)

#Both ends falls inside
rgl::lines3d(c(orig[2, 1], end[2, 1]),
             c(orig[2, 2], end[2, 2]),
             c(orig[2, 3], end[2, 3]), col = "red")

line_AABB(orig[2,], end[2,], AABB_min, AABB_max)

#Oring falls inside, but not the end.
rgl::lines3d(c(orig[3, 1], end[3, 1]),
             c(orig[3, 2], end[3, 2]),
             c(orig[3, 3], end[3, 3]), col = "blue")

line_AABB(orig[3,], end[3,], AABB_min, AABB_max)

#End falls inside, but not the orig
rgl::lines3d(c(orig[4, 1], end[4, 1]),
             c(orig[4, 2], end[4, 2]),
             c(orig[4, 3], end[4, 3]), col = "green")

line_AABB(orig[4,], end[4,], AABB_min, AABB_max)

#Some segments of the line are intercepted
rgl::lines3d(c(orig[5, 1], end[5, 1]),
             c(orig[5, 2], end[5, 2]),
             c(orig[5, 3], end[5, 3]), col = "black")

line_AABB(orig[5,], end[5,], AABB_min, AABB_max)
```

---

lines\_interception      *Intersection of Lines by AABBs*

---

## Description

Intersection of lines by several Axis-Aligned Bounding Boxes.

## Usage

```
lines_interception(orig, end, AABBs, edge_length, threads = 1, progress = TRUE)
```

**Arguments**

orig	A data.table with the describing *XYZ* coordinates of the the start path of the rays.
end	A data.table with the describing *XYZ* coordinates of the the end path of the rays.
AABBs	A data.table with *XYZ* coordinates of the center of AABBs.
edge_length	A positive numeric vector with the AABB length edge for the X, Y, and Z coordinates.
threads	An integer $\geq 0$ describing the number of threads to use. This need to be used if parallel = TRUE.
progress	Logical, if TRUE displays a graphical progress bar. TRUE as default.

**Value**

It returns a data.table with nine columns: 1-5 columns with the counts for the code of intersection (see [line\\_AABB](#)), and 6-9 columns with sum the path length of intersection. The number of rows match with nrow(AABBs).

**Author(s)**

J. Antonio Guzmán Q.

**See Also**

[line\\_AABB](#), [voxels](#)

**Examples**

```
#Create points with paths
n <- 20
orig <- data.table(X = runif(n, min = -5, max = 5),
                  Y = runif(n, min = -5, max = 5),
                  Z = runif(n, min = -5, max = 5))

end <- data.table(X = runif(n, min = -5, max = 5),
                 Y = runif(n, min = -5, max = 5),
                 Z = runif(n, min = -5, max = 5))

#Create a potential AABB
AABBs <- data.table(X = 0, Y = 0, Z = 0)
edge_length <- c(2, 2, 2)

#Plot

cube <- rgl::cube3d()
cube <- rgl::scale3d(cube,
                    edge_length[1]/2,
                    edge_length[2]/2,
                    edge_length[3]/2)
```

```

box <- rgl::translate3d(cube, AABBs[[1]], AABBs[[2]], AABBs[[3]])
rgl::shade3d(box, col= "green", alpha = 0.6)
rgl::points3d(orig, size = 5, col = "black")
rgl::points3d(end, size = 5, col = "red")

for(i in 1:nrow(orig)) {
  rgl::lines3d(c(orig[[1]][i], end[[1]][i]),
               c(orig[[2]][i], end[[2]][i]),
               c(orig[[3]][i], end[[3]][i]), col = "grey")
}

#Estimation
lines_interception(orig, end, AABBs, edge_length, progress = FALSE)

```

---

min_distance	<i>Minimum Distance Between Points</i>
--------------	--

---

### Description

Estimate the minimum distance between points in a point cloud.

### Usage

```

min_distance(
  cloud,
  distance = "euclidean",
  threads = 1L,
  verbose = FALSE,
  progress = FALSE,
  ...
)

```

### Arguments

cloud	A data.table with *XYZ* coordinates in the first three columns representing a point cloud.
distance	Type of distance to calculate. "euclidean" as default. Look hnsw_knn for more options.
threads	An integer specifying the number of threads to use for parallel processing. Experiment to see what works best for your data on your hardware.
verbose	If TRUE, log messages to the console.
progress	If TRUE, log a progress bar when verbose = TRUE. Tracking progress could cause a small overhead.
...	Arguments passed to hnsw_build and hnsw_search.

**Value**

A numeric vector describing the minimum distance between points.

**Author(s)**

J. Antonio Guzmán Q.

**Examples**

```
data("pc_tree")

#Estimate the minimum distance of a sample o 100 points
min_distance(pc_tree)
```

---

pc\_tree

*A Tree Point Cloud*

---

**Description**

A data.table from a point cloud of a tree with a spatial point resolution of 0.05 mm.

**Usage**

```
data(pc_tree)
```

**Format**

A data.table with three columns, which are:

**X** the "X" coordinate

**Y** the "Y" coordinate

**Z** the "Z" coordinate

A data.table where the rows represent the points and the three columns represent the \*XYZ\* coordinates.

**References**

Guzman, Sharp, Alencastro, Sanchez-Azofeifa. 2018. To be published.

**Examples**

```
data(pc_tree)
head(pc_tree)
```

## Description

The plot method for objects of class "voxels" created using the [voxels](#) function.

## Usage

```
plot_voxels(  
  voxels,  
  add.points = TRUE,  
  add.voxels = TRUE,  
  border = TRUE,  
  points.size = 1,  
  points.col = "black",  
  fill.col = "forestgreen",  
  line.lwd = 0.5,  
  line.col = "black",  
  alpha = 0.1,  
  ...  
)
```

## Arguments

voxels	Object of class "voxels" from <a href="#">voxels</a> .
add.points	Logical, if TRUE it adds the original points used to perform the voxelization. TRUE as default.
add.voxels	Logical, if TRUE it adds the voxels created. TRUE as default.
border	Logical, if TRUE it adds a line on the borders of each voxel. TRUE as default.
points.size	The points size, a positive number to use if plot add.points = TRUE.
points.col	A character defining the color of the points to use.
fill.col	A character vector defining the color to fill the voxels, it could be a range of colors or a solid color.
line.lwd	The line width, a positive number, defaulting to 0.5.
line.col	A character defining the color of the border lines to use.
alpha	A positive numeric vector describing the transparency of the voxels to fill. This value must be between 0.0 (fully transparent) .. 1.0 (opaque).
...	General arguments passed to <a href="#">plot3d</a> .

## Value

A 3D plot of a point cloud and voxels.

**Author(s)**

J. Antonio Guzmán Q.

**See Also**

[voxels](#), [voxels\\_counting](#), [summary\\_voxels](#)

**Examples**

```
data("pc_tree")

###Create cubes of a size of 7x7x3.5.
vox <- voxels(pc_tree, edge_length = c(7, 7, 3.5))
plot_voxels(vox)
```

---

polar\_to\_cartesian      *Polar to Cartesian Coordinates*

---

**Description**

Convert from polar to cartesian coordinates.

**Usage**

```
polar_to_cartesian(polar, threads = 1, digits = NULL)
```

**Arguments**

polar	A data.table with three columns describing the zenith, azimuth, and distance of a point to the center.
threads	An integer vector describing the number of threads for parallel processing. Default 1.
digits	A numeric vector of length 1 describing the decimal numbers to <a href="#">round</a> the cartesian coordinates. If NULL, <a href="#">round</a> does not apply. NULL as default.

**Value**

A data.table with three columns describing the \*XYZ\* of the cartesian coordinates.

**Author(s)**

J. Antonio Guzmán Q.

**See Also**

[cartesian\\_to\\_polar](#)

**Examples**

```
#Creates a hemisphere of points each 2 degrees

zenith <- seq(0, 90, 2)
azimuth <- seq(0, 360, 2)
hemi <- CJ(zenith, azimuth)
hemi$distance <- 1
hemicloud <- polar_to_cartesian(hemi)
rgl::plot3d(hemicloud)
```

---

radius_search	<i>Radius Search of Points</i>
---------------	--------------------------------

---

**Description**

Adapted radius searching of points based on RcppHNSW

**Usage**

```
radius_search(
  query,
  ref,
  radius,
  max_neighbour,
  distance = "euclidean",
  same = FALSE,
  threads = 1L,
  verbose = FALSE,
  progress = FALSE,
  ...
)
```

**Arguments**

query	A data.table containing the set of query points where each row represent a point and each column a given coordinate.
ref	A numeric containing the set of reference points where each row represent a point and each column a given coordinate.
radius	A numeric describing maximum euclidean distance form the each query points in which a point can be consider a neighbor.
max_neighbour	An integer specifying the maximum number of ref points to look around to consider for a given radius.
distance	Type of distance to calculate. "euclidean" as default. Look hnsw_knn for more options.

same	Logic. If TRUE, it delete neighbors with distance of 0, useful when the k search is based on the same query.
threads	An integer specifying the number of threads to use for parallel processing. Experiment to see what works best for your data on your hardware.
verbose	If TRUE, log messages to the console.
progress	If TRUE, log a progress bar when verbose = TRUE. Tracking progress could cause a small overhead.
...	Arguments passed to hnsw_build and hnsw_search.

### Details

This function is based on hnswlib C++ library (Malkov & Yashunin 2016) and its bindings for R (RcppHNSW; Melville 2020) for a fast estimation of neighbors points. It is adapted to simplify the workflow within rTLS. If you use this function, please consider cite the C++ library and RcppHNSW package.

### Value

A data.table with three columns describing the indices of the query and ref points and the distances.

### Author(s)

J. Antonio Guzmán Q.

### References

Malkov, Y. A., & Yashunin, D. A. (2016). Efficient and robust approximate nearest neighbor search using Hierarchical Navigable Small World graphs. arXiv preprint arXiv:1603.09320.

### See Also

[radius\\_search](#)

### Examples

```
#Point cloud  
data("pc_tree")
```

```
#Radius search of 1  
radius_search(pc_tree, pc_tree, radius = 1, max_neighbour = 100)
```

---

rotate2D	<i>Rotate a plane of coordinates</i>
----------	--------------------------------------

---

**Description**

Rotate a plane of coordinates to a given angle.

**Usage**

```
rotate2D(plane, angle, threads = 1)
```

**Arguments**

plane	A data.table with two columns describing the plane of coordinates.
angle	A numeric vector describing the degrees of rotation.
threads	An integer specifying the number of threads to use. Experiment to see what works best for your data on your hardware.

**Value**

A data.table with the rotation applied to plane.

**Author(s)**

J. Antonio Guzmán Q.

**Examples**

```
data(pc_tree)

plot(pc_tree[,1:2])

#Rotate in 45 degrees using Z axis of the cloud
plot(rotate2D(pc_tree[,1:2], angle = 45))
```

---

rotate3D	<i>Rotate a Point Cloud</i>
----------	-----------------------------

---

**Description**

Rotate point cloud based on the roll, pitch, and yaw angles.

**Usage**

```
rotate3D(cloud, roll = 0, pitch = 0, yaw = 0, threads = 1)
```

**Arguments**

cloud	A data.table with three columns describing the *XYZ* coordinates of a point cloud.
roll	A numeric vector describing the degrees of rotation angles for roll (*X*).
pitch	A numeric vector describing the degrees of rotation angles for pitch (*Y*).
yaw	A numeric vector describing the degrees of rotation angles for yaw (*Z*). for the roll, pitch, and yaw.
threads	An integer specifying the number of threads to use. Experiment to see what works best for your data on your hardware.

**Details**

The \*XYZ\* coordinates are transformed to E-N-U coordinates (ENU system, East-North-Up).

**Value**

A data.table with the rotation applied to cloud.

**Author(s)**

J. Antonio Guzmán Q.

**Examples**

```
data(pc_tree)
rgl::plot3d(pc_tree)
rgl::plot3d(rotate3D(pc_tree, roll = 45, pitch = 45, yaw = 0))
```

---

stand\_counting

*Stand Counting*


---

**Description**

Applies the [voxels\\_counting](#) function on a grid base point cloud.

**Usage**

```
stand_counting(
  cloud,
  xy.res,
  z.res = NULL,
  points.min = NULL,
  min_size,
  edge_sizes = NULL,
  length_out = 10,
```

```

bootstrap = FALSE,
R = NULL,
progress = TRUE,
parallel = FALSE,
threads = NULL
)

```

### Arguments

cloud	A data.table of a point cloud with xyz coordinates in the first three columns.
xy.res	A positive numeric vector describing the grid resolution of the xy coordinates to perform.
z.res	A positive numeric vector of length 1 describing the vertical resolution. If z.res = NULL vertical profiles are not used.
points.min	A positive numeric vector of length 1 minimum number of points to retain a sub-grid.
min_size	A positive numeric vector of length 1 describing the minimum cube edge length to perform. This is required if edge_sizes = NULL.
edge_sizes	A positive numeric vector describing the edge length of the different cubes to perform within each subgrid when z.res = NULL. If edge_sizes = NULL, it uses the maximum range of values for the xyz coordinates.
length_out	A positive integer of length 1 indicating the number of different edge lengths to use for each subgrid. This is required if edge_sizes = NULL.
bootstrap	Logical. If TRUE, it computes a bootstrap on the H index calculations. FALSE as default.
R	A positive integer of length 1 indicating the number of bootstrap replicates. This need to be used if bootstrap = TRUE.
progress	Logical, if TRUE displays a graphical progress bar. TRUE as default.
parallel	Logical, if TRUE it uses a parallel processing for the voxelization. FALSE as default.
threads	An integer $\geq 0$ describing the number of threads to use. This need to be used if parallel = TRUE.

### Value

A data.table with the summary of the voxels per grid created with their features.

### Author(s)

J. Antonio Guzmán Q.

### See Also

[voxels\\_counting](#), [voxels](#), [summary\\_voxels](#)

**Examples**

```

data(pc_tree)

#Applying stand_counting.

stand_counting(pc_tree, xy.res = c(4, 4), min_size = 3)

#Applying stand_counting using bootstrap in the H index.

stand_counting(pc_tree,
               xy.res = c(4, 4),
               min_size = 3,
               bootstrap = TRUE,
               R = 10)

```

---

summary\_voxels

*Voxels Summary*


---

**Description**

Create a summary objects of class "voxels" created using the [voxels](#).

**Usage**

```
summary_voxels(voxels, edge_length = NULL, bootstrap = FALSE, R = NULL)
```

**Arguments**

voxels	An object of class <code>voxels</code> created using the <code>voxels()</code> function or a <code>data.table</code> describing the voxels coordinates and their number of points produced using <code>voxels()</code> .
edge_length	A positive numeric vector with the voxel-edge length for the x, y, and z coordinates. This need to be used if <code>class(voxels) != "voxels"</code> . It use the same dimensional scale of the point cloud.
bootstrap	Logical, if TRUE it computes a bootstrap on the H index calculations. FALSE as default.
R	A positive integer of length 1 indicating the number of bootstrap replicates. This need to be used if <code>bootstrap = TRUE</code> .

**Details**

The function provides 12 main statistics of the voxels. Specifically, the first three columns represent the edge length of the voxels, the following three columns (ei. N\_voxels, Volume, Surface) describe the number of voxels created, the total volume that they represent, and the surface area that they cover. Following columns represent the mean (Density\_mean) and sd (Density\_sd) of the density of points per voxel (e.g. points/m<sup>2</sup>). Columns 9:12 provide metrics calculated using the Shannon Index. Specifically, H describe the entropy, H\_max the maximum entropy, Equitativity the ratio between H and Hmax, and Negentropy describe the product of Hmax - H. If bootstrap = TRUE four more columns are created (13:16). These represent the mean and sd of the H index estimated using bootstrap (H\_boot\_mean and H\_boot\_sd), the Equitativity\_boot as the ratio of the ratio between H\_boot\_sd and Hmax, and Negentropy\_boot as the product Hmax - H\_boot\_mean.

**Value**

A data.table with with the summary of voxels.

**Author(s)**

J. Antonio Guzmán Q.

**See Also**

[voxels](#), [voxels\\_counting](#), [plot\\_voxels](#)

**Examples**

```
data("pc_tree")

#Apply a summary on a object of class "voxels" using bootstrap with 1000 replicates.
vox <- voxels(pc_tree, edge_length = c(0.5, 0.5, 0.5))
summary_voxels(vox, bootstrap = TRUE, R = 1000)

#Apply a summary on a product from 'voxels' using bootstrap with 1000 replicates.
vox <- voxels(pc_tree, edge_length = c(0.5, 0.5, 0.5), obj.voxels = FALSE)
summary_voxels(vox, edge_length = c(0.5, 0.5, 0.5), bootstrap = TRUE, R = 1000)
```

---

TLS\_scan

*A TLS scan*

---

**Description**

A data.table from a TLS scan.

**Usage**

```
data(TLS_scan)
```

**Format**

A `data.table` with five columns, which are:

**X** the "X" coordinate

**Y** the "Y" coordinate

**Z** the "Z" coordinate

**Target\_count** The number of received by the same laser shot

**Target\_index** The rank of the returned pulse in the target count of received by the same laser shot

A `data.table` where the rows represent the pulse returns and the three columns represent the \*XYZ\* coordinates, and the target count and index.

**Details**

A TLS scan conducted using a Reigel VZ400i with a vertical and horizontal resolution of 0.048 and 0.622 degrees (2082 and 580 lines, respectively). The scanner has frame of scanning between 30 and 130.024 degrees zenith and 0 and 359.90 degrees azimuth. At the moment of the scan the roll, pitch, and yaw of the scanner were 1.026, 0.746, -110.019, respectively. The scanner coordinates in this scan are  $x = 0$ ,  $y = 0$ ,  $z = 0$ .

**Examples**

```
data(TLS_scan)
head(TLS_scan)
```

---

tree\_metrics

*Tree Metrics*

---

**Description**

Estimate the tree height, crown area, and the diameter at breast height of a tree point cloud

**Usage**

```
tree_metrics(cloud, region.diameter = NULL, relocateZ = TRUE)
```

**Arguments**

cloud	A <code>data.table</code> of the target point with three columns of the *XYZ* coordinates.
region.diameter	A numeric vector of length 2 indicating the lower and higher region to subset the point cloud and get the diameter. If <code>region.diameter = NULL</code> , it use <code>c(1.25, 1.35)</code> . <code>NULL</code> as default.
relocateZ	Logical, if <code>TRUE</code> it relocates the *Z* coordinates to a minimum coordinate of zero based on the current <code>min(cloud[, 3])</code> . Useful if the base value (*Z*) of a tree point cloud is not topography corrected.

**Details**

The tree height is estimated based on the maximum value of *\*Z\**, the crown area is calculated applying a convex hull on the point cloud, while the DBH is calculated extracting the area of the convex hull on the subset of points between `region.diameter`, and then estimating the diameter of a circle. For another estimation of DBH try [circleRANSAC](#) or for irregular trucks try [trunk\\_volume](#).

**Value**

A `data.table` with the tree height, crown area, and diameter

**Author(s)**

J. Antonio Guzman Q. and Ronny Hernandez

**See Also**

[circleRANSAC](#), [trunk\\_volume](#)

**Examples**

```
data("pc_tree")
tree_metrics(pc_tree)
```

---

trunk_volume	<i>Tree Trunk Volume</i>
--------------	--------------------------

---

**Description**

Estimates the tree trunk volume of a point cloud using the [ashape3d](#) package.

**Usage**

```
trunk_volume(cloud, max.height = NULL, alpha = 0.2, plot = TRUE, ...)
```

**Arguments**

<code>cloud</code>	A <code>data.table</code> with three columns representing the <i>*XYZ*</i> coordinates of a point cloud.
<code>max.height</code>	A numeric vector to contemplate points in the cloud lower than a specific height. If <code>NULL</code> , it performs the alpha-shape on the entire point cloud.
<code>alpha</code>	A numeric vector of length one passed to <code>ashape3d</code> to describes alpha. <code>alpha = 0.20</code> as default since it seems to provide better estimations of the trunk volume. However, the alpha value may depends on the resolution of the point cloud.
<code>plot</code>	Logical. If <code>TRUE</code> , it uses <code>plot.ashape3d</code> to represent the alpha-shape.
<code>...</code>	General arguments passed to <code>ashape3d</code> .

**Details**

This is an adaptation of the code develop by Lafarge & Pateiro-Lopez (2017) based on Edelsbrunner & Mucke (1994) for the quick extraction of the tree trunk volume. Therefore, if you use this code we kindly suggest to cite these documents in your research.

**Value**

A numeric vector with the estimated trunk volume.

**Author(s)**

J. Antonio Guzmán Q.

**References**

Lafarge, T., Pateiro-Lopez, B. (2017). Implementation of the 3D Alpha-Shape for the Reconstruction of 3D Sets from a Point Cloud. Available at <https://CRAN.R-project.org/package=alphashape3d>.

Edelsbrunner, H., Mucke, E. P. (1994). Three-Dimensional Alpha Shapes. ACM Transactions on Graphics, 13(1), pp.43-72.

**See Also**

[tree\\_metrics](#), [circleRANSAC](#)

**Examples**

```
data("pc_tree")

#Estimates the trunk volume of a height lower than 1.75.
trunk_volume(pc_tree, max.height = 1.75)
```

---

voxels

*Voxelization of a Point Cloud*

---

**Description**

Create cubes of a given distance in a point cloud though their voxelization. It use a modify version of the code used in Greaves et al. 2015.

**Usage**

```
voxels(cloud, edge_length, threads = 1L, obj.voxels = TRUE)
```

## Arguments

<code>cloud</code>	A <code>data.table</code> with <code>*XYZ*</code> coordinates in the first three columns.
<code>edge_length</code>	A positive numeric vector with the voxel-edge length for the x, y, and z coordinates. It use the same dimensional scale of the point cloud.
<code>threads</code>	An integer specifying the number of threads to use for parallel processing. Experiment to see what works best for your data on your hardware.
<code>obj.voxels</code>	Logical. If <code>obj.voxel = TRUE</code> , it returns an object of class "voxels", If <code>obj.voxel = FALSE</code> , it returns a <code>data.table</code> with the coordinates of the voxels created and the number of points in each voxel. TRUE as default.

## Details

Voxels are created from the negative to the positive `*XYZ*` coordinates.

## Value

If `obj.voxels == TRUE`, it return an object of class "voxels" which contain a list with the points used to create the voxels, the parameter `edge_length`, and the voxels created. If `FALSE`, it returns a `data.table` with the coordinates of the voxels created and the number of points in each voxel.

## Author(s)

J. Antonio Guzmán Q.

## References

Greaves, H. E., Vierling, L. A., Eitel, J. U., Boelman, N. T., Magney, T. S., Prager, C. M., & Griffin, K. L. (2015). Estimating aboveground biomass and leaf area of low-stature Arctic shrubs with terrestrial LiDAR. *Remote Sensing of Environment*, 164, 26-35.

## See Also

[voxels\\_counting](#), [plot\\_voxels](#), [summary\\_voxels](#)

## Examples

```
data("pc_tree")

###Create cube of a size of 0.5.
voxels(pc_tree, edge_length = c(0.5, 0.5, 0.5))
```

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voxels_counting	<i>Voxels Counting</i>
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### Description

Creates cube like voxels of different size on a point cloud using the `voxels` function, and then return a `summary_voxels` of their features.

### Usage

```
voxels_counting(
  cloud,
  edge_sizes = NULL,
  min_size,
  length_out = 10,
  bootstrap = FALSE,
  R = NULL,
  progress = TRUE,
  parallel = FALSE,
  threads = NULL
)
```

### Arguments

<code>cloud</code>	A data.table with xyz coordinates of the point clouds in the first three columns.
<code>edge_sizes</code>	A positive numeric vector describing the edge length of the different cubes to perform. If NULL, it use edge sizes by default based on the largest range of XYZ and <code>min_size</code> .
<code>min_size</code>	A positive numeric vector of length 1 describing the minimum cube edge length to perform. This is required if <code>edge_sizes = NULL</code> .
<code>length_out</code>	A positive interger of length 1 indicating the number of different edge lengths to use. This is required if <code>edge_sizes = NULL</code> .
<code>bootstrap</code>	Logical. If TRUE, it computes a bootstrap on the H index calculations. FALSE as default.
<code>R</code>	A positive integer of length 1 indicating the number of bootstrap replicates. This need to be used if <code>bootstrap = TRUE</code> .
<code>progress</code>	Logical, if TRUE displays a graphical progress bar. TRUE as default.
<code>parallel</code>	Logical, if TRUE it uses a parallel processing for the voxelization. FALSE as default.
<code>threads</code>	An integer $\geq 0$ describing the number of threads to use. This need to be used if <code>parallel = TRUE</code> .

### Value

A data.table with the summary of the voxels created with their features.

**Author(s)**

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**See Also**

[voxels](#), [summary\\_voxels](#), [plot\\_voxels](#)

**Examples**

```
data(pc_tree)

#Applying voxels counting.
voxels_counting(pc_tree, min_size = 2)

#Voxels counting using bootstrap on the H indexes with 1000 repetitions.
voxels_counting(pc_tree, min_size = 2, bootstrap = TRUE, R = 1000)
```

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