Package ‘ecochange’

Type   Package
Title  Integrating Ecosystem Remote Sensing Products to Derive EBV Indicators
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Description Essential Biodiversity Variables (EBV) are state variables with dimensions on time, space, and biological organization that document biodiversity change. Freely available ecosystem remote sensing products (ERSP) are downloaded and integrated with data for national or regional domains to derive indicators for EBV in the class ecosystem structure (Pereira et al., 2013) <doi:10.1126/science.1229931>, including horizontal ecosystem extents, fragmentation, and information-theory indices. To process ERSP, users must provide a polygon or geographic administrative data map. Downloadable ERSP include Global Surface Water (Peckel et al., 2016) <doi:10.1038/nature20584>, Forest Change (Hansen et al., 2013) <doi:10.1126/science.1244693>, and Continuous Tree Cover data (Sexton et al., 2013) <doi:10.1080/17538947.2013.786146>.
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**Description**

This function is a wrapper of `cellStats` used to compute statistics for spatial indicators in the EBV class ecosystem structure. To derive the spatial indicators see functions `echanges` and `sampleIndicator`.

**Usage**

```r
EBVstats(ccp, stats, ...
```

**Arguments**

- `ccp` `echanges`, or RasterStack or NULL. If NULL then NULL is returned.
- `stats` character. vector of stats defined in `cellStats`. If missing then six summary statistics, including 'mean', 'sd', 'min', 'max', are computed.
- `...` Additional arguments in `cellStats`

**Value**

tibble.

**Author(s)**

Wilson Lara Henao <wilarhen@gmail.com> [aut, cre], Victor Gutierrez-Velez [aut], Ivan Gonzalez [ctb], Maria C. Londono [ctb]
Examples

```r
## RasterBrick of structural Essential Biodiversity Variables
## covering the extent of a location in the northern Amazon basin
## (Colombia) is imported:
path. <- system.file('amazon.grd', package = 'ecochange')
amazon <- brick(path.)

## Changes in layers of tree-canopy cover (TC) are computed by
## processing the `amazon' brick:
def <- echanges(amazon, eco = 'TC',
                change = 'lossyear',
                eco_range = c(1,80),
                get_unaffected = TRUE,
                binary_output = FALSE,
                mc.cores = 2)

## Function 'EBVstats' is used to compute ecosystem statistics
st_amazon <- EBVstats(def)

## A plot of the `st_amazon' object
plot.EBVstats(st_amazon,
              cex = 1.5,
              xlab = 'Year',
              ylab = 'Canopy cover (%)',
              main = 'Ecosystem changes',
              sub = 'Northern Amazon',
              fill = 'Layer')
```

description

This function produces ecosystem-change maps by masking cell values in a layer of ecosystem changes over a target set of ecosystem variables. The function allows focusing the ecosystem-change analysis on a species distribution range.

Usage

```r
echanges(ps, eco = names(ps[[1:(nayers(ps) - 1)]]),
          change = names(ps[[nayers(ps)]]),
          sp_dist, eco_range = c(1,
                                  100),
          change_vals = 1:19,
          sp_dist_range = c(1, 1),
          spread = TRUE,
          get_unaffected = TRUE,
          binary_output = FALSE,
          noDataValue = 0,
          mc.cores = round(detectCores() *
                          0.6, 0), ...)
```
Arguments

- **ps**: RasterStack or SpatialPolygonsDataFrame. Stack of spatial data, including the target ecosystem variables, a layer of changes, and an alternative layer of a species distribution range. This argument can also be a polygon geometry used to integrate such spatial data via implementation of `rsp2ebv`; see the ellipsis term below.

- **eco**: character. Regular expression matching names of a subset of layers representing the target ecosystem variables. Default matches names of the first 1:(n-1) layers in `ps`.

- **change**: character. Name of the layer of ecosystem changes. Default matches the name of the last layer in `ps`.

- **sp_dist**: character. Name of an alternative layer representing a species distribution range. If missing then this argument is ignored.

- **eco_range**: numeric. Range of values in the target ecosystem variable.

- **change_vals**: numeric. Vector of values in the layer of ecosystem changes.

- **sp_dist_range**: numeric. Range of values in the alternative layer of species distribution range. This argument is ignored if `sp_dist` is missing.

- **spread**: logical. Spread representation of ecosystem changes. Users do not need to change this argument. It is used by other routines to fastening computation of ecosystem horizontal extents. If FALSE then the function mask cell values in the target ecosystem variables over the layer of ecosystem changes. Default TRUE.

- **get_unaffected**: logical. Extract unaffected areas. If FALSE then pixel values of the ecological variable across the changed areas are extracted. Default TRUE.

- **binary_output**: logical. Produce binary outputs (masks). If FALSE then ranges of values of the ecological variable are maintained. Default FALSE.

- **noDataValue**: numeric. Output NoDataValue. Default 0.

- **mc.cores**: numeric. The number of cores. Default uses around 60 percent of the CPU capacity.

- **...**: If `ps` is a polygon then additional arguments in `rsp2ebv`.

Value

- Class `echanges`.

Author(s)

Wilson Lara Henao <wilarhen@gmail.com> [aut, cre], Victor Gutierrez-Velez [aut], Ivan Gonzalez [ctb], Maria C. Londono [ctb]

References


**Examples**

```r
## Brick with structural Essential Biodiversity Variables covering the
## extent of a location in the northern Amazon basin (Colombia):
path. <- system.file('amazon.grd', package = 'ecochange')
amazon <- brick(path.)

## Changes in layers of tree-canopy cover (TC) in the 'amazon'
## brick are computed:
def <- echanges(amazon, eco = 'TC',
               change = 'lossyear',
               eco_range = c(1,80),
               get_unaffected = TRUE,
               binary_output = FALSE,
               mc.cores = 2)

## Method 'plot.echanges' allows comparing rasters using a common scale bar:
plot.echanges(def)
```

---

**gaugeIndicator**

### Gauge Biodiversity Indicator

**Description**

This function processes ecosystem-change maps from `echanges` to calculate biodiversity indicators, including ecosystem extent, entropy, fractal dimension, among others. To sample the indicators across fixed-size grids see `sampleIndicator`.

**Usage**

```r
gaugeIndicator(ps, ...,
               metric = "area_ha",
               smp_lsm = list(),
               mc.cores = round(detectCores() * 0.6, 0))
```
Arguments

- **ps**: SpatialPolygonsDataFrame or RasterStack. Polygon geometry used to produce ecosystem-change maps via the implementation of `echanges` or the stack of ecosystem-change maps.

  ... If `ps` is a polygon then additional arguments in `echanges` or `rsp2ebv`.

- **metric**: character. The name of an indicator. Default 'area_ha' computes ecosystem areas (ha) at class level. See the argument 'metric' in `list_lsm` to implement other metrics.

- **smp_lsm**: list. List of arguments in `calculate_lsm`. This argument is ignored when `metric = 'area_ha'`.

- **mc.cores**: numeric. The number of cores. Default uses around 60 percent of the cores.

Details

Coordinate reference system of the spatial units must have metric units UTM. Performance in the computation of ecosystem extents is optimized via the implementation of the function `tabuleRaster`. Indicators other than ecosystem extents are calculated implementing `calculate_lsm`.

Value

Class `Indicator`.

Author(s)

Wilson Lara Henao <wilarhen@gmail.com> [aut, cre], Victor Gutierrez-Velez [aut], Ivan Gonzalez [ctb], Maria C. Londono [ctb]

References


Examples

```r
## RasterBrick of structural Essential Biodiversity Variables
## covering the extent of a location in the northern Amazon basin
## (Colombia) is imported:
path. <- system.file('amazon.grd', package = 'ecochange')
amazon <- brick(path.)
```
## Changes in layers of tree-canopy cover (TC) in the 'amazon' brick are computed:
def <- exchanges(amazon, eco = 'TC',
  change = 'lossyear',
  eco_range = c(1,80),
  get_unaffected = TRUE,
  binary_output = FALSE,
  mc.cores = 2)

## Function 'gaugeIndicator' is used to compute ecosystem areas (default):
am_areas <- gaugeIndicator(def,
  mc.cores = 2)

plot.Indicator(am_areas)

---

**getGADM**

*Get Geographic Administrative Unit*

**Description**

This function is a wrapper of `getData` used to import levels in Geographic Administrative Data Maps (GADM).

**Usage**

```r
getGADM(unit.nm = NULL,
  level = 2, country = "COL",
  path = tempdir())
```

**Arguments**

- **unit.nm** character or NULL. Name of an administrative unit (e.g. municipality), or the name of such an unit plus its corresponding higher-level unit (e.g. department/state). If NULL then a list of unit names is printed.
- **level** numeric. A number between zero and two, indicating any of the levels of administrative subdivisions in GADM (0=country, 1=first administrative subdivision, and 2=second administrative subdivision).
- **country** character. ISO code specifying a country. Default 'COL'.
- **path** character. Path name indicating where the unit will be stored. Default stores the data in a temporary directory.

**Value**

`SpatialPolygonsDataFrame` or character vector of GADM units.
Author(s)
Wilson Lara Henao <wilarhen@gmail.com> [aut, cre], Victor Gutierrez-Velez [aut], Ivan cre Gonzalez [ctb], Maria C. Londono [ctb]

References
https://gadm.org/

Examples
## Printing municipalities of Colombia:

```r
muni <- getGADM(NA)
head(muni)
```

getrsp

*Get remote sensing product*

Description
This function processes the extent of a predefined region of interest (polygon geometry or GADM unit) to download ecosystem remote sensing products (ERSP). Downloadable ERSP include Global Surface Water, Forest Change, and Continuous Tree Cover data. See `listGP`.

Usage
```r
getrsp(roi = NULL, ..., lyrs = NULL, path,
       rewrite.pass = FALSE,
       verify.web = FALSE,
       mc.cores = round(detectCores() * 0.6, 0))
```

Arguments
- **roi** `SpatialPolygonsDataFrame; or character; or NULL`. Region of interest. This can be either 1) a polygon geometry; or 2) the name of a GADM unit (see `getGADM`); or 3) a NULL value. Default NULL makes the function to print a list of GADM units.
- **...** If roi is a GADM unit then additional arguments in `getGADM`.
- **lyrs** `character`. Remote-sensing products. Default NULL makes the function to print a list of Downloadable data. see `listGP`.
- **path** `character`. Path name indicating where the variables are stored. If missing then a folder named as ‘ecochange’ created in a current temporary directory is used.
getrsp

**rewrite.pass**  logical. Rewrite password. Only valid to download new NASA Earth data, see details section.

**verify.web**  logical. Verify in the web whether the URLs used to download the rsp are available. See `getOption('webs')`. Default FALSE.

**mc.cores**  numeric. The number of cores. Default uses around 60 percent of the cores.

**Details**

Downloads of Continuous Tree Cover data require user authentication through the NASA Earth data Login. To obtain a NASA Earth data Login account, please visit: [https://urs.earthdata.nasa.gov/users/new](https://urs.earthdata.nasa.gov/users/new).

**Value**

Path names of the remote sensing products just retrieved, or character vectors suggesting GADM units/Global Products that can be used to download ERSP (see NULL defaults in arguments 'roi' and 'lyrs').

**Author(s)**

Wilson Lara Henao &lt;wilarhen@gmail.com&gt; [aut, cre], Victor Gutierrez-Velez [aut], Ivan Gonzalez [ctb], Maria C. Londono [ctb]

**References**


**Examples**

```r
## Polygon of the Colombian municipality of Cartagena del Chaira:
load(system.file('cchaira_roi.RData', package = 'ecochange'))

## A Global Surface Water layer ('seasonality') which covers the
## extent of the polygon is retrieved:

rsp_cchaira <- getrsp(cchaira_roi,
  lys = 'seasonality', mc.cores = 2, path = tempdir())
file.exists(rsp_cchaira)
```
getWRS  

Get WRS

Description

This function processes regions of interest (a polygon geometry or GADM unit) to find corresponding Landsat Path/Row World Reference System (WRS) polygons. This function is internally implemented by `getrsp`.

Usage

```r
getWRS(roi = NULL, path = tempdir(), ...
```

Arguments

- **roi**  
  SpatialPolygonsDataFrame; or character; or NULL. Region of interest. This can be whether 1) a polygon geometry; or 2) the name of a GADM unit (see `getGADM`); or 3) a NULL value. Default NULL makes the function to print a list of GADM units.

- **path**  
  character. Path name indicating where the WRS data are processed.

- **...**  
  Additional arguments in `getGADM`.

Value

SpatialPolygonsDataFrame, or set of GADM units.

Author(s)

Wilson Lara Henao &lt;wilarhen@gmail.com&gt; [aut, cre], Victor Gutierrez-Velez [aut], Ivan Gonzalez [ctb], Maria C. Londono [ctb]

Examples

```r
load(system.file('cchaira_roi.RData',package = 'ecochange'))

wrs_cchaira <- getWRS(cchaira_roi)
plot(wrs_cchaira)
```
Description

This function prints information about ecosystem remote sensing products that can be downloaded with `getrsp`.

Usage

```r
listGP(layer = TRUE,
       Algorithm = TRUE,
       author = TRUE, funs = FALSE,
       api.code = FALSE)
```

Arguments

- `layer` character. Add column 'layer' to the data.
- `Algorithm` character. Add column 'Algorithm' to the data.
- `author` character. Add column 'author' to the data.
- `funs` character. Add column 'funs' to the data.
- `api.code` character. Add column 'api.code' to the data.

Value

tibble.

Author(s)

Wilson Lara Henao <wilarhen@gmail.com> [aut, cre], Victor Gutierrez-Velez [aut], Ivan Gonzalez [ctb], Maria C. Londono [ctb]

References


Examples

```r
lst <- listGP()
```

---

**plot.EBVstats**

**Visualize EBVstats objects**

**Description**

Plots for objects from `EBVstats` are printed.

**Usage**

```r
## S3 method for class 'EBVstats'
plot(x, y, ...)
```

**Arguments**

- `x`: tibble. Data set of statistics such as that produced by `EBVstats`.
- `y`: character. Color scale. If missing then `grDevices::terrain.colors(n)`, where `n` is the number of layers, is implemented.
- `...`: Arguments to be passed to some plot methods.

**Author(s)**

Wilson Lara Henao <wilarhen@gmail.com> [aut, cre], Victor Gutierrez-Velez [aut], Ivan cre Gonzalez [ctb], Maria C. Londono [ctb]

**Examples**

```r
## RasterBrick of structural Essential Biodiversity Variables
## covering the extent of a location in the northern Amazon basin
## (Colombia) is imported:
path. <- system.file('amazon.grd',package = 'ecochange')
avian <- brick(path.)

## Changes in layers of tree-canopy cover (TC) are computed by
## processing the 'amazon' brick:
def <- echanges(amazon, eco = 'TC',
                 change = 'lossyear',
                 eco_range = c(1,80),
                 get_unaffected = TRUE,
                 binary_output = FALSE,
                 mc.cores = 2)

## Function 'EBVstats' is used to compute ecosystem statistics
st_amazon <- EBVstats(def)
```
plot.echanges

## A plot of the 'st_amazon' object
plot.EBVstats(st_amazon,
  cex = 1.5,
  xlab = 'Year',
  ylab = 'Canopy cover (%)',
  main = 'Ecosystem changes',
  sub = 'Municipality: Cartagena del Chaire',
  fill = 'Layer')

plot.echanges Visualize ecosystem changes

### Description

This function can display level and box plots for objects from `rsp2ebv`, `echanges`, or `sampleIndicator`.

### Usage

```r
## S3 method for class 'echanges'
plot(x, y, ...)
```

### Arguments

- **x**  
  Raster*, or `echanges`. RasterStack object or ecosystem-change representation.

- **y**  
  character. A color palette. If this is missing or the suggest `viridis` is not installed then `terrain.colors` is implemented.

- **...**  
  Graphical arguments:
  - `type`: what type of plot should be drawn: "p" for level plots (default), or "b" for box plots,
  - `cex`: adjustment of sizes for most text values. If missing then `cex = 1`; if a main title is specified then it is increased `1.4*cex`,
  - `xlab`, and `ylab`: titles for the x and y axes,
  - `main`: a text of the main title,
  - `labels`: a string or numeric sequence for the panel titles

### Author(s)

Wilson Lara Henao <wilarhen@gmail.com> [aut, cre], Victor Gutierrez-Velez [aut], Ivan Gonzalez [ctb], Maria C. Londono [ctb]
Examples

```r
## Brick with structural Essential Biodiversity Variables covering the
## extent of a location in the northern Amazon basin (Colombia):
path. <- system.file('amazon.grd', package = 'ecochange')
amazon <- brick(path.)

## Changes in layers of tree-canopy cover (TC) in the 'amazon'
## brick are computed:
def <- echanges(amazon, eco = 'TC',
                  change = 'lossyear',
                  eco_range = c(1,80),
                  get_unaffected = TRUE,
                  binary_output = FALSE,
                  mc.cores = 2)

plot.echanges(def)
```

---

### plot.Indicator

#### Visualize Indicator objects

**Description**

Plots for objects from `gaugeIndicator` are produced.

**Usage**

```r
## S3 method for class 'Indicator'
plot(x, y, ...
```

**Arguments**

- `x`  
  *tibble.* Data set of indicators such as that produced by `gaugeIndicator`.
- `y`  
  *character.* A color palette. If this is missing or the suggest `viridis` is not installed then `terrain.colors` is implemented.
- `...`  
  Graphical arguments:
  - `type`: what type of plot should be drawn: "s" for stacked bar plots (default), or "b" for box plots,
  - `cex`: adjustment of sizes for most text values,
  - `xlab`, and `ylab`: titles for the x and y axes,
  - `main`: a text of the main title,
  - `sub`: a text for the sub title,
  - `labels`: a string or numeric sequence for the x-axis labels,
  - `fill`: a text for the legend title
**Author(s)**

Wilson Lara Henao <wilarhen@gmail.com> [aut, cre], Victor Gutierrez-Velez [aut], Ivan Gonzalez [ctb], Maria C. Londono [ctb]

**Examples**

```r
## RasterBrick of structural Essential Biodiversity Variables
## covering the extent of a location in the northern Amazon basin
## (Colombia) is imported:
path. <- system.file('amazon.grd',package = 'ecochange')
amazon <- brick(path.)

## Changes in layers of tree-canopy cover (TC) are computed by
## processing the 'amazon' brick:
def <- echanges(amazon, eco = 'TC',
change = 'lossyear',
eco_range = c(1,80),
get_unaffected = TRUE,
binary_output = FALSE,
mc.cores = 2)

## Function 'gaugeIndicator' is used to compute ecosystem areas
## (default metric = 'area_ha'):
am_areas <- gaugeIndicator(def,
mc.cores = 2)

## A plot of the 'am_areas' object
plot.Indicator(am_areas,
cex = 1.5,
xlab = 'Year',
ylab = 'Area (ha)',
main = 'Ecosystem changes',
sub = 'Northern amazon',
fill = 'Forest cover (%)')
```

**rsp2ebv**

*Integrate remote sensing products*

**Description**

This function integrates ecosystem remote sensing products and produces raster-data sections with the cell values enclosed in a region of interest.

**Usage**

```r
rsp2ebv(ps = NULL, ...,
lyrs = NULL, path,
sr, ofr = c(30, 30),
```
mc.cores = round(detectCores() * 0.6, 0))

Arguments

- **ps** (SpatialPolygonsDataFrame; or character; or NULL): Region of interest. This can be whether 1) a polygon geometry; or 2) the name of a GADM unit (see `getGADM`); or 3) a NULL value. Default NULL makes the function to print a list of GADM units.

- **...** (Additional arguments in `getGADM` and `getrsp`):

- **lyrs** (character): Remote-sensing products. Default NULL makes the function to print a list of Downloadable data, see `listGP`.

- **path** (character): Path name indicating where the variables are stored. If missing then a folder named as 'ecochange' created in a current temporary directory is used.

- **sr** (character): PROJ.4 description of the target coordinate reference system. If missing then the target layers are projected to metric system UTM.

- **ofr** (numeric): \(c(xres, yres)\). Output file resolution (in target georeferenced units). Default \(c(30, 30)\) m2.

- **mc.cores** (numeric): The number of cores. Default uses around 60 percent of the cores.

Details

This function implements 'sf::gdal_utils' so it assumes the user's machine has a valid GDAL installation.

Value

Class echanges.

Author(s)

Wilson Lara Henao <wilarhen@gmail.com> [aut, cre], Victor Gutierrez-Velez [aut], Ivan Gonzalez [ctb], Maria C. Londono [ctb]

References


## Examples

### A Global Surface Water layer (`'seasonality'`) covering the extent of a Colombian municipality Cartagena del Chaira is formatted into an spatial EBV:

```r
load(system.file('cchaira_roi.RData', package = 'ecochange'))

rsp_cchaira <- getrsp(cchaira_roi, lyrs = 'seasonality', mc.cores = 2, path = tempdir())

file.exists(rsp_cchaira)

season_cchaira <- rsp2ebv(cchaira_roi, lyrs = 'seasonality', path = tempdir())

plot.echanges(season_cchaira)
```

---

### sampleIndicator

#### Sample Biodiversity indicator

This function divides into fixed-size grids each of the scenes of a stack of ecosystem-spatial data and samples a biodiversity indicator by every grid. To compute biodiversity indicators at the class and landscape levels, see `gaugeIndicator`.

#### Usage

```r
sampleIndicator(ps = NULL, ..., metric = "condent", classes = 5, min = 1, max = 100, side, smp_lsm = list(level = "landscape"), mc.cores = round(detectCores() * 0.6, 0))
```

#### Arguments

- `ps` (SpatialPolygonsDataFrame or RasterStack): Polygon geometry used to produce ecosystem-change maps via the implementation of `echanges` or the stack of ecosystem-change maps.
- `...` (If `ps` is a polygon): Additional arguments in `echanges` or `rsp2ebv`.
- `metric` (character): The name of an indicator other than ecosystem extent. This can be cohesion ('cohesion'), conditional entropy ('condent'), perimeter-area fractal dimension ('pafrac'), among others, see package `list_lsm`. Default 'condent'.

---

### Description

This function divides into fixed-size grids each of the scenes of a stack of ecosystem-spatial data and samples a biodiversity indicator by every grid. To compute biodiversity indicators at the class and landscape levels, see `gaugeIndicator`.

---

### Usage

```r
sampleIndicator(ps = NULL, ..., metric = "condent", classes = 5, min = 1, max = 100, side, smp_lsm = list(level = "landscape"), mc.cores = round(detectCores() * 0.6, 0))
```
sampleIndicator

classes numeric; or NULL. Number of evenly spaced classes used to reclassify the layers. Default 5. If NULL then the layers are not reclassified.

min numeric. If classes != NULL then minimum cell value in the layers. Default 1

max numeric. If classes != NULL then maximum cell value in the layers. Default 100

side numeric. Side of the sampling grid (m). If missing the function tries to find a grid size the samples at least a grid with a non-NA value of the indicator.

smp_lsm list. Additional arguments in sample_lsm

mc.cores numeric. The number of cores. Default uses 60 percent of the cores.

Value

Class echanges

Author(s)

Wilson Lara Henao <wilarhen@gmail.com> [aut, cre], Victor Gutierrez-Velez [aut], Ivan Gonzalez [ctb], Maria C. Londono [ctb]

References


Examples

## RasterBrick of structural Essential Biodiversity Variables
## covering the extent of a location in the northern Amazon basin
## (Colombia) is imported:
path. <- system.file('amazon.grd',package = 'ecochange')
amazon <- brick(path.)

## Changes in layers of tree-canopy cover (TC) in the 'amazon'
## brick are computed:
def <- echanges(amazon, eco = 'TC',
    change = 'lossyear',
    eco_range = c(1,80),
    get_unaffected = TRUE,
    binary_output = FALSE,
    mc.cores = 2)

plot.echanges(amazon)
## Function `sampleIndicator` is implemented to sample a metric of conditional entropy (default):
```
def_condent <- sampleIndicator(def, side = 400, mc.cores = 2)
```
```
plot.echones(def_condent, cex = 1.5)
```

---

**Description**

This function generates frequency tables for scenes in ecosystem remote sensing products by wrapping `rasterDT`. The function is mapped by `gaugeIndicator` to optimize computation of ecosystem extents.

**Usage**

```
tabuleRaster(layer = "",
             del0 = TRUE, useNA = "no",
             n256 = FALSE)
```

**Arguments**

- **layer** character. File path to an ERSP scene.
- **del0** logical. Remove the 0-count categories.
- **useNA** logical. Include NA values. This argument is passed to `rasterDT::freqDT`.
- **n256** logical. Do the raster contains less than 256 unique values?

**Value**

data.frame.

**Author(s)**

Ivan Gonzalez [aut], Wilson Lara Henao <wilarhen@gmail.com> [aut, cre], Victor Gutierrez-Velez [aut], Maria C. Londono [ctb]

**Examples**

```
tabuleRaster(raster(volcano), n256 = FALSE)
```

---
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