

Package ‘TOC’

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Type Package

Title Total Operating Characteristic (TOC) Curve

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Description Construction of the Total Operating Characteristic (TOC) Curve and the Re-
ceiver (aka Relative) Operating Characteristic (ROC) Curve for spatial data.

License GPL (>= 2)

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TOC-package	<i>Total Operating Characteristic (TOC) Curve</i>
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Description

Construction of the Total Operating Characteristic (TOC) Curve and the Receiver (aka Relative) Operating Characteristic (ROC) Curve for spatial data.

Details

Package: TOC
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See Also

[TOC](#), [plot](#)

plot

Plot an object of class Toc or Roc

Description

Plot a Total Operating Characteristic (TOC) curve or a Relative Operating Characteristic (ROC)

Usage

```
## S4 method for signature 'Toc'
plot(x, labelThres=FALSE, digits=3, modelLeg="Model", ...)
```

```
## S4 method for signature 'Roc'
plot(x, labelThres=FALSE, digits=3, modelLeg="Model", ...)
```

Arguments

x	An object of class Toc or Roc
labelThres	logical, default to FALSE. If TRUE, thresholds are labeled in the TOC plot
digits	integer indicating the number of decimal places (round) or significant digits (signif) to be used for labeling the thresholds and the numeric axes. Negative values are allowed. See Details in the round function
modelLeg	a character string for labeling the model in the legend
...	additional parameters to be passed to plot

Value

a plot showing the TOC or the ROC curves

See Also

[TOC](#), [ROC](#)

Examples

```
## Not run:
index <- raster("p_built01_suitability_1.rst")
boolean <- raster("BuiltGain1985_1999.rst")
mask <- raster("1985NonBuilt01.rst")
tocd <- TOC(index, boolean, mask)
plot(tocd, main="TOC curve")
rocd <- ROC(index, boolean, mask)
plot(rocd, main="ROC curve")

## End(Not run)

## Not run:
index <- raster("Prob_Map2.rst")
boolean <- raster("Change_Map2b.rst")
mask <- raster("MASK3.rst")
tocd <- TOC(index, boolean, mask, nthres=100)
plot(tocd, main="TOC curve")
rocd <- ROC(index, boolean, mask, nthres=100)
plot(rocd, main="ROC curve")

## End(Not run)
```

ROC	<i>Construct the table for the ROC curve</i>
-----	--

Description

Construct the table for the Relative Operating Characteristic (ROC) curve

Usage

```
ROC(index, boolean, mask=NULL, nthres=NULL, thres=NULL, NAval=0, P=NA, Q=NA,
progress=FALSE)
```

Arguments

index	index Raster map
boolean	boolean Raster map
mask	mask Raster map
nthres	an optional integer indicating the number of equal-interval thresholds to be evaluated for the ROC curve. See Details below
thres	an optional numeric vector of thresholds to be evaluated for the ROC curve. See Details below

NAval	value for nodata (NA values) in the mask map
P	count of reference presence observations in the population
Q	count of reference absence observations in the population
progress	logical; if TRUE, a progress bar is shown

Details

thresholds are calculated as the unique values of the index map after masking out NA values (default option), if neither nthres nor thres is provided. The default option can be time-consuming if the amount of unique values in the index map (after masking out NA values) is large (e.g., > 1000). In the latter case, the user may prefer to enter specified thresholds (with the thres argument), or to indicate the number of equal-interval thresholds to be evaluated for the ROC curve (with the nthres argument)

Value

an object of class Roc containing the ROC table, the area under the curve (AUC), maximum AUC and minimum AUC

See Also

[plot](#)

Examples

```
## Not run:
index <- raster("p_built01_suitability_1.rst")
boolean <- raster("BuiltGain1985_1999.rst")
mask <- raster("1985NonBuilt01.rst")

# all unique values of the index map after applying the mask are used as thresholds (default
# option)
rocd <- ROC(index, boolean, mask)
rocd

# thresholds can also be defined by indicating the number of equal-interval thresholds
rocd <- ROC(index, boolean, mask, nthres=10)
rocd

# A vector of thresholds can be used to define the thresholds
rocd <- ROC(index, boolean, mask, thres=seq(0, 100, by=10))
rocd

## End(Not run)
## Not run:
index <- raster("Prob_Map2.rst")
boolean <- raster("Change_Map2b.rst")
mask <- raster("MASK3.rst")

# thresholds can also be defined by indicating the number of equal-interval thresholds
rocd <- ROC(index, boolean, mask, nthres=100)
rocd

# A vector of thresholds can be used to define the thresholds
```

```

rocd <- ROC(index, boolean, mask, thres=seq(0, 100, by=10))
rocd

## End(Not run)

```

scaling

scale the output TOC values and change units

Description

scale the 'Hits' and 'Hits+FalseAlarms' values in the TOC output table, as well as the prevalence and population, using a scaling factor. Labels for the modified units in the TOC object are changed to newUnits

Usage

```

## S4 method for signature 'Toc'
scaling(x, scalingFactor, newUnits)

```

Arguments

x	an object of class Toc
scalingFactor	numeric value to scale 'Hits' and 'Hits+FalseAlarms' values in the TOC output table, as well as the prevalence and population
newUnits	charater string for the new data units in the TOC object

Value

an object of class TOC

See Also

[TOC](#), [ROC](#)

Examples

```

## Not run:
index <- raster("p_built01_suitability_1.rst")
boolean <- raster("BuiltGain1985_1999.rst")
mask <- raster("1985NonBuilt01.rst")
tocd <- TOC(index, boolean, mask)
plot(tocd)

# scale units from square m to square km
tocd_sqkm <- scaling(tocd, scalingFactor=1000000, newUnits="square km")
plot(tocd_sqkm)

## End(Not run)
## Not run:
index <- raster("Prob_Map2.rst")
boolean <- raster("Change_Map2b.rst")

```

```

mask <- raster("MASK3.rst")
tocd <- TOC(index, boolean, mask, nthres=100)
plot(tocd)

# scale units from square m to square km
tocd_sqkm <- scaling(tocd, scalingFactor=1000000, newUnits="square km")
plot(tocd_sqkm)

## End(Not run)

```

TOC

*Construct the table for the TOC curve***Description**

Construct the table for the Total Operating Characteristic (TOC) curve

Usage

```
TOC(index, boolean, mask=NULL, nthres=NULL, thres=NULL, NAval=0, P=NA, Q=NA,
progress=FALSE)
```

Arguments

index	index Raster map
boolean	boolean Raster map
mask	mask Raster map
nthres	an optional integer indicating the number of equal-interval thresholds to be evaluated for the TOC curve. See Details below
thres	an optional numeric vector of thresholds to be evaluated for the TOC curve. See Details below
NAval	value for nodata (NA values) in the mask map
P	count of reference presence observations in the population
Q	count of reference absence observations in the population
progress	logical; if TRUE, a progress bar is shown

Details

thresholds are calculated as the unique values of the index map after masking out NA values (default option), if neither `nthres` nor `thres` is provided. The default option can be time-consuming if the amount of unique values in the index map (after masking out NA values) is large (e.g., > 1000). In the latter case, the user may prefer to enter specified thresholds (with the `thres` argument), or to indicate the number of equal-interval thresholds to be evaluated for the TOC curve (with the `nthres` argument)

Value

an object of class `Toc` containing the TOC table, the area under the curve (AUC), maximum AUC and minimum AUC, the prevalence, the population and the data units (for data in the TOC table, prevalence and population)

See Also[plot](#)**Examples**

```
## Not run:
index <- raster("p_built01_suitability_1.rst")
boolean <- raster("BuiltGain1985_1999.rst")
mask <- raster("1985NonBuilt01.rst")

# all unique values of the index map after applying the mask are used as thresholds (default
# option)
tocc <- TOC(index, boolean, mask)
tocc

# thresholds can also be defined by indicating the number of equal-interval thresholds
tocc <- TOC(index, boolean, mask, nthres=10)
tocc

# A vector of thresholds can be used to define the thresholds
tocc <- TOC(index, boolean, mask, thres=seq(0, 100, by=10))
tocc

## End(Not run)
## Not run:
index <- raster("Prob_Map2.rst")
boolean <- raster("Change_Map2b.rst")
mask <- raster("MASK3.rst")

# thresholds can also be defined by indicating the number of equal-interval thresholds
tocc <- TOC(index, boolean, mask, nthres=100)
tocc

# A vector of thresholds can be used to define the thresholds
tocc <- TOC(index, boolean, mask, thres=seq(0, 100, by=10))
tocc

## End(Not run)
```

Toc-class

*Toc and Roc classes***Description**

Toc and Roc classes

Objects from the Class

Objects can be created by calls of the form `new("Toc", ...)`, or with the helper functions such as `Toc`.

Slots

Slots for Roc and Toc objects

table: data.frame

AUC: numeric; Area Under the Curve

maxAUC: numeric; maximum AUC

minAUC: numeric; minimum AUC

prevalence: numeric; prevalence

population: numeric; population

units: character; units for data in the TOC table, prevalence and population

Examples

```
showClass("Toc")
```

Toc-methods	<i>Methods for the Roc and Toc classes</i>
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Description

Methods for the Roc and Toc classes

Details

Methods for the Roc and Toc classes

Value

an object of class Toc or Roc

See Also

[TOC](#), [ROC](#)

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