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Kyoto University
Package ‘wvtool’

November 8, 2016

Type Package
Title Image Tools for Automated Wood Identification
Version 1.0
Date 2016-11-07
Author Junji Sugiyama, Kayoko Kobayashi
Maintainer Kayoko Kobayashi <kayoko_kobayashi@rish.kyoto-u.ac.jp>
Description This tool, wood vision tool, is intended to facilitate preprocessing and analyzing 2-
dimensional wood images toward automated recognition. The former includes some ba-
sics such as functions to RGB to grayscale, gray to binary, cropping, rotation(bilinear), med-
dian/mean/Gaussian filter, and Canny/Sobel edge detection. The latter includes gray level co-
ocurrence matrix (GLCM), Haralick parameters, local binary pattern (LBP), higher order lo-
cal autocorrelation (HLAC), Fourier transform (radial and azimuthal integration), and Gabor fil-
tering. The functions are intended to read data using ‘readTIFF(x,info=T)’ from ‘tiff’ pack-
age. The functions in this packages basically assumes the grayscale images as in-
put data, thus the color images should be subjected to the function rgb2gray() be-
fore used for some other functions.
License GPL (>= 2)
NeedsCompilation no
Suggests tiff
Depends R (>= 2.10)
Repository CRAN
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**bin2dec**

**Description**

A function returns decimal number from a sequential binary number. The function is internally used in lbp function.

**Usage**

\[ \text{bin2dec}(x) \]

**Arguments**

- \( x \)  
  A sequence of 0, 1 numbers

**Value**

A decimal number

**See Also**

dec2bin, lbp
camphora

---

### Image Sample Dataset

**Description**

An optical micrographs of Cinnamomum camphora

**Usage**

```r
data("camphora")
```

**Format**

The format is: num [1:486, 1:518] 0.275 0.337 0.765 0.937 0.933 ... - attr(*, "bits.per.sample")= int 8 - attr(*, "samples.per.pixel")= int 1 - attr(*, "sample.format")= chr "uint" - attr(*, "planar.config")= chr "contiguous" - attr(*, "compression")= chr "none" - attr(*, "x.resolution")= num 32 - attr(*, "y.resolution")= num 32 - attr(*, "resolution.unit")= chr "inch" - attr(*, "orientation")= chr "top.left" - attr(*, "color.space")= chr "black is zero"

**Source**

Kyoto University Xylarium Database

**References**

http://database.rish.kyoto-u.ac.jp

**Examples**

```r
data(camphora)
## maybe str(camphora)
```

---

car2pol

---

### Polar Transformer -Cartesian to Polar Coordinates-

**Description**

The function converts images to polar coordinates. The polar transformation is useful for unwarping images which have a generally round object. From power spectrum for example, one may generate radial integration profile or azimuthal intensity distribution. Default is "bilinear" interpolation.

**Usage**

```r
car2pol(x, method="bilinear")
```
Arguments

x A raster image or a matrix.
method "NN" Nearest neighbour method, "bilinear" Bilinear interpolation.

Value

A matrix in polar coordinate system of the requested image

pol.img Radial distance corresponds to the shorter side of requested image, and polar angle covers 0 to 360 degrees.

See Also

integ.profile

Examples

data(camphora)
par(mfrow=c(1,2))
image(rot90(c(camphora)), col=gray(c(0:255)/255), main="camphora, original",
useRaster=TRUE, asp=1, axes=FALSE)
img <- car2pol(camphora, method="bilinear")
image(rot90(c(img)), col=gray(c(0:255)/255), main="camphora, polar (bilinear)",
 xlab="radial distance(pixel)",ylab="angle(deg)", useRaster=TRUE, asp=1, axes=FALSE)

Description

A function labels the connected components in a binary image. For example, it can be used for statistical analysis of tracheids (see examples).

Usage

cc.label(x, connect=8, inv=FALSE, img.show=FALSE, text.size=0.3)

Arguments

x A binary image (A matrix with 0 and 1)
connect 8-connectivity or 4-connectivity. Default is 8-connectivity.
inv inverse the binary image x before labelling. Labelling the connected area with 0 when this is TRUE.
img.show If this is TRUE, the image with labelling numbers are shown.
text.size the size of labelling numbers used when img.show=TRUE.
Details

Labelling the connected components with pixels equal to 1 (white) in a binary image (If pixels equal to 0 (black) should be labelled, select inv=TRUE). The function returns the labelled image and the statistical data of the labelled components.

Value

a list with 2 components (a matrix and a dataframe)

image A matrix with labels
summary A dataframe summarizing the labelled components with 8 following variables.
summary$label labelling numbers, area: area of each component
summary$aveX, summary$aveY center position of each component
summary$dX, summary$dY width and height of each component
summary$edge If the component is on the edge of the image, this value is 1, otherwise 0.

Examples

```r
## Not run:
data(cryptomeria)
ing <- rgb2gray(cryptomeria)
ing.c <- crop(img,300,300)
ing.bin <- gray2bin(img.c, auto=FALSE, th=180)
par(mfrow=c(2,2))
test <- cc.label(img.bin, connect=8, img.show=TRUE)
hist(test$summary$area,main="histogram of area")
hist(test$summary$dX,main="histogram of dX")
hist(test$summary$dY,main="histogram of dY")

## End(Not run)
```

---

crop

**Image cropping**

Description

image cropping from the center.

Usage

crop(x, width=300, height=300, shift=c(0,0))
Arguments

x A raster or a matrix
width width for cropping
height height for cropping
shift shift of the cropped position from the center

Value

A raster or a matrix

Examples

data(camphora)
par(mfrow=c(2,2))
image(rot90(c(camphora)),col=gray(c(0:255)/255), main="original", useRaster=TRUE, axes=FALSE, asp=1)
image(rot90(c(crop(camphora,200,100)),col=gray(c(0:255)/255),
main="cropped from the center", useRaster=TRUE, axes=FALSE, asp=0.5)
image(rot90(c(crop(camphora,200,200)),col=gray(c(0:255)/255),
main="cropped from the center", useRaster=TRUE, axes=FALSE, asp=1)
image(rot90(c(crop(camphora,200,200,shift=c(50,50))),col=gray(c(0:255)/255),
main="cropped from shifted position c(50,50)", useRaster=TRUE, axes=FALSE, asp=1)

Image Sample Dataset

Description

An optical micrographs of Cryptomeria japonica

Usage

data("cryptomeria")

Format

num [1:1079, 1:1000, 1:4] 0.886 0.89 0.863 0.639 0.424 ... - attr(*, "bits.per.sample")= int 8 - attr(*, "samples.per.pixel")= int 4 - attr(*, "sample.format")= chr "uint" - attr(*, "planar.config")= chr "contiguous" - attr(*, "compression")= chr "none" - attr(*, "x.resolution")= num 72 - attr(*, "y.resolution")= num 72 - attr(*, "resolution.unit")= chr "inch" - attr(*, "orientation")= chr "top.left" - attr(*, "artist")= chr "DP" - attr(*, "date.time")= chr "2016:07:29 11:38:28" - attr(*, "color.space")= chr "RGB"

Source

Kyoto University Xylarium Database
References

http://database.rish.kyoto-u.ac.jp

Examples

data(cryptomeria)
## maybe str(cryptomeria)

---

dec2bin  

**Decimal to Binary Conversion**

**Description**

A function returns binary number from decimal number. The function is internally used in lbp function.

**Usage**

```r
dec2bin(x, digit=8)
```

**Arguments**

- `x` A decimal integer.
- `digit` A length of binary sequence.

**Value**

A binary number in array.

**See Also**

bin2dec, lbp

---

dec2bin  

**Canny and Sobel Edge detector.**

**Description**

A function detects edges in images by Canny or Sobel operator. Sobel provides approximate intensity of gradients for each pixels, while Canny provides a binary image with thin edges.

**Usage**

```r
detect(x, thresh1=1, thresh2=15, noise="gaussian", noise.s=3, method="Canny")
```
Arguments

- **x**: A raster image or a matrix
- **thresh1**: low threshold for edge tracking by hysteresis (0-100). Only used for "Canny" edge detector.
- **thresh2**: high threshold for edge tracking by hysteresis (0-100). Only used for "Canny" edge detector.
- **noise**: a method for noise reduction. "gaussian", "median", and "mean" filters are available. Default is "gaussian".
- **noise.s**: filter size for noise reduction (3 or 5). Default is 3.
- **method**: "Canny" and "Sobel" can be selected. Default is "Canny".

Details

Canny edge detector has four steps. 1. noise reduction/ 2. finding the gradient in images by Sobel operator/ 3. Non-maximum suppression/ 4. Hysteresis threshold. When the method "Sobel" is selected, only step 1 and 2 will be done.

Value

A raster or a matrix

See Also

noise.filter

Examples

```r
## Not run:
data(camphora)
data(cryptomeria)
cryptomeria <- rgb2gray(cryptomeria)
img.c1 <- crop(camphora, 200, 200)
img.c2 <- crop(cryptomeria, 300, 300)
par(mfrow=c(2,2))
image(rot90c(edge.detect(img.c1, thresh1=1, thresh2=15, noise="gaussian", noise.s=3, method="Canny")), col=gray(c(0:255)/255), main="Canny", useRaster=TRUE, axes=FALSE, asp=1)
image(rot90c(edge.detect(img.c1, thresh1=1, thresh2=15, noise="gaussian", noise.s=3, method="Sobel")), col=gray(c(0:255)/255), main="Sobel", useRaster=TRUE, axes=FALSE, asp=1)
image(rot90c(edge.detect(img.c2, thresh1=1, thresh2=15, noise="gaussian", noise.s=3, method="Canny")), col=gray(c(0:255)/255), main="Canny", useRaster=TRUE, axes=FALSE, asp=1)
image(rot90c(edge.detect(img.c2, thresh1=1, thresh2=15, noise="gaussian", noise.s=3, method="Sobel")), col=gray(c(0:255)/255), main="Sobel", useRaster=TRUE, axes=FALSE, asp=1)
## End(Not run)
```
**gabor.filter**

**Two Dimensional Gabor Filtering in Frequency Domain**

**Description**

The function provides two dimensional Gabor function proposed by Daugman to model the spatial summation properties in the visual cortex. It returns Gabor filter in real and reciprocal space, and filtered image.

**Usage**

```r
gabor.filter(x, lamda=5, theta=45, bw=1.5, phi=0, asp=1, disp=FALSE)
```

**Arguments**

- `x`: A raster or matrix to be filtered
- `lamda`: Wavelength of the cosine part of Gabor filter kernel in pixel. Real number greater than 2 can be used. However, lamda=2 should not be used with phase offset (phi) = -90 or 90.
- `theta`: The orientation of parallel strips of Gabor function in degree
- `bw`: Half response spatial frequency bandwidth of a Gabor filter. This relates to the ratio sigma/lamda, where sigma is the standard deviation of Gaussian factor of Gabor function.
- `phi`: Phase offset of the cosine part of Gabor filter kernel in degree
- `asp`: Ellipticity of the Gabor function. asp=1 means circular. For asp<1 it gives elongated parallel strip
- `disp`: If this operator is TRUE, original image, gabor filter in real space domain, that in frequency domain, and filtered image will be generated

**Value**

The function provides following four outputs.

- `kernel`: 151x151 gabor filter kernel
- `mask`: A mask with kernel in the center in a space domain
- `freq_mask`: Real part of Fourier transform of the mask in spatial frequency domain
- `filtered_image`: Inversed Fourier transform of FFT(img)*FFT(mask)

**References**

Examples

data(cryptomeria)
img <- rgb2gray(cryptomeria)
img <- crop(img, 300, 300)
# simple example
test <- gabor.filter(x=img, lambda=8, theta=60, bw=1.5, phi=0, asp=0.3, disp=TRUE)
## Not run:
# azimutal intensity distribution with respect to the orientation of Gabor function
par(mfrow=c(2,1))
Integ <- array()
filt.img <- matrix(0, nrow(img), ncol(img))
for ( i in 1:60 ) {
  out <- gabor.filter(x=img, lambda=8, theta=3*i, bw=1.5, phi=0, asp=0.3)
  filt.img <- out$filtered_img + filt.img
  Integ[i] <- sum(out$filtered_img*out$filtered_img)
}
image(rot90c(filt.img), col=gray(c(0:255)/255), axes=FALSE, useRaster=TRUE)
x <- 1:60
plot(3*x, Integ, ty="l", ylab="integrated intensity (a.u.)", xlab="azimuthal angle (deg)")
## End(Not run)

glc

Gray Level Co-occurrence Matrix (glcm)

description

This function supports calculating gray level co-occurrence matrices from a grayscale image (< 8 bit) with requested gray level. The gray level of the source image is read from the attributes data from input TIFF file.

Usage

glc(x, t.level=4, d=1)

Arguments

x A gray scale image or matrix. "x" assumes an output from readTIFF(filename, as.is=T, info=T)

t.level A target grey level for GLCM calculation in bite. The grayscale is truncated linearly.
d Displacement between adjacent i, j points in pixel.

Details

The data in matrix is either inspected as images or subsequently used to calculate Haralick texture features, originally published 15 features(Haralick et al., 1973) and two additionals (Albregsten, 1995).
Value

The gray level cooccurrence matrices of 4 directions (theta) and their average, gray level, and displacement vector are listed.

- glcm: GLCM at theta = "0", "45", "90", "135" degree and "average"
- level: number of gray level
- d: length of displacement vector

References

- Albregtsen F (1995) Statistical texture measures computed from gray level cooccurrence matrices. In: Technical Note, Department of Informatics, University of Oslo, Norway

See Also

gray2bin, rgb2gray, haralick,

Examples

data(camphora)
img <- camphora
par(mfrow=c(1,2))
lev <- 4
theta <- c(1,3)  # "th_0","th_90"
theta_c <- c("th_0","th_90")
dist <- 1
for (i in 1:2) {
tst <- glcm(img,lev,dist)
title <- paste(lev, "bit", " glcm ", theta_c[i], " d=" , dist, sep="")
persp(tst$glcm[i], theta=30, phi=30,main=title, asp=1,
     xlab="i", ylab="j", zlab="probability")
}

Description

A function provides automatic clustering-based thresholding proposed by Ohtsu, and a gray scale image is converted to binary image. Initial histogram and discriminant level of binning are displayed by his=TRUE, dis=TRUE options. A threshold value can be also set manually.
Usage

```
gray2bin(x, auto=TRUE, th=200, his=FALSE, dis=FALSE)
```

Arguments

- `x`: A raster image or a matrix.
- `auto`: set threshold automatically (Otsu method) or manually.
- `th`: a threshold value used when `auto=FALSE`.
- `his`: A histogram of initial gray scale image.
- `dis`: A plot of variation between classes divided by variation within classes.

Value

A requested binary image. Black is zero.

References


See Also

`rgb2gray`

Examples

```
data(camphora)
par(mfrow=c(2,3))
image(rot90c(camphora), col=gray((0:255)/255), main="camphora", asp=1, useRaster=TRUE, axes=FALSE)
out <- gray2bin(camphora, his=TRUE, dis=TRUE)
image(rot90c(out), col=gray((0:255)/255), main="binary image, auto", asp=1, useRaster=TRUE, axes=FALSE)
image(rot90c(gray2bin2(camphora,auto=FALSE,th=100)), col=gray((0:255)/255), main="binary image, thresh=100", asp=1, useRaster=TRUE, axes=FALSE)
image(rot90c(gray2bin2(camphora,auto=FALSE,th=180)), col=gray((0:255)/255), main="binary image, thresh=180", asp=1, useRaster=TRUE, axes=FALSE)
```

Description

A function returns 15 Haralick features for 4 directions, their average and range.

Usage

```
haralick(x)
```
Arguments

x output of glcm() function from a TIFF data

Details

15 outputs are
#1 Angular Second Moment / Homogeniety "asm" #2 Contrast "con" #3 inverse Difference Moment "idm" #4 Entropy "ent" #5 Correlation "cor" #6 Variance in Haralick 1973 "var"
#7 Sum Average "sav" #8 Sum Entropy "sen" #9 Difference Entropy "den" #10 Difference Variance 
"dva" #11 Sum Variance "sva" #12 Information Measures of Correlation "f12" #13 Information Measures of Correlation "f13" #14 Cluster Shade "sha" #15 Cluster prominence "pro", respectively

Value

A matrix of angles and features

References


Albregtsen F (1995) Statistical texture measures computed from gray level cooccurrence matrices. In: Technical Note, Department of Informatics, University of Oslo, Norway


See Also

glm

Examples

data(camphora)
haralick(glm(camphora, 6, 1))

hlac

Higher Order Local Autocorrelation (HLAC)

Description

Feature extraction for practical vision system, whose features are shift-invariant and additive. The function gives zero to the eighth order cases, represented by 223 mask patterns of 3 x 3 within a 2r+1 x 2r+1 (r >=1) displacement region.

Usage

hlac(x, r=1, disp=FALSE)
Arguments

x  A binary or gray image or matrix
r  Displacement vector r for 3 x 3 mask pattern
disp  If TRUE, function saves 223 filtered images in one matrix.

Details

The feature parameter should be a list. The function returns 1, 4, 20, 45, 62, 54, 28, 8, 1 features and corresponding filtered images if disp is TRUE.

Value

HLAC features or the corresponding image with requested HLAC measures.

features  Numerical output of 0 to 8th order masks
mat  A large matrix of 223 images expanded in a row

References


See Also

rgb2gray, gray2bin, glcm, lbp

Examples

# features plot and the corresponding image presentation
data(camphora)
tmp <- hlac(gray2bin(camphora), 2, disp=TRUE)
par(mfrow=c(2,2))
plot(unlist(tmp$features), main="HLAC histogram")
image(rot90c(matrix(tmp$mat[2,],tmp$row,tmp$col)),
col = gray((255:0)/255), main="2", useRaster=TRUE, asp=1, axes=FALSE)
image(rot90c(matrix(tmp$mat[23,],tmp$row,tmp$col)),
col = gray((255:0)/255), main="23", useRaster=TRUE, asp=1, axes=FALSE)
image(rot90c(matrix(tmp$mat[156,],tmp$row,tmp$col)),
col = gray((255:0)/255), main="156", useRaster=TRUE, asp=1, axes=FALSE)
**Simple Integration for Making Profile**

**Description**

A function returns integrated line profile. It crops rectangular area from a requested size and project and integrate pixel values either to horizontal or vertical axis. When used with a matrix in polar coordinate (car2pol) calculated from power spectrum (power.spec) of an image, the function provides radial integration or azimuthal integration that are useful for diffraction analysis.

**Usage**

```r
integ.profile(x, axis="H", h=c(20, 50), v=c(30, 120), disp=FALSE)
```

**Arguments**

- `x`: A raster image or a matrix
- `axis`: Axis to project. `H`: Projection to horizontal axis or radial distance (in polar coordinate). `V`: Projection to vertical axis or azimuthal angle (in polar coordinate).
- `h`: `c(h1,h2)`: A horizontal or radial (in polar coordinate) range for integration.
- `v`: `c(v1,v2)`: A vertical or azimuthal (in polar coordinate) range for integration.
- `disp`: Plot calculated profile. Default is FALSE.

**Details**

The row and column corresponds to horizontal and vertical axes, respectively.

**Value**

An array of requested line profile

**See Also**

`swap.quad`, `car2pol`, `fft`, `Mod`

**Examples**

```r
data("camphora")
img <- camphora
par(mfrow=c(2,2))
image(rot90c(img),col=gray(c(0:255)/255), useRaster=TRUE, main="camphora",asp=1, axes=FALSE)
integ.profile(img, axis="H", h=c(1,nrow(img))), v=c(1,ncol(img)), disp=TRUE)
integ.profile(img, axis="V", h=c(1,nrow(img))), v=c(1,ncol(img)), disp=TRUE)
ps <- log(sw rap.quad(Mod(fft(img))))
pol <- car2pol(ps)
image(rot90c(ps),col=gray(c(0:255)/255), useRaster=TRUE,main="power spectrum",asp=1, axes=FALSE)
```
### lbp

**Description**

Calculate local binary patterns from a grayscale image

**Usage**

```r
lbp(x, r=1)
```

**Arguments**

- `x` A raster image or a matrix
- `r` displacement vector in 8 direction. `r=1` means `c(-1, 0, -1, 1, 1, 1, 0, -1, 0, 1, 1, 1, 0, -1) r=2 means c(-2, 1, -1, 2, 1, 2, 1, -1, 1, -2, -1, 2, 1)`

---

### lbnorm

**Counts 0-1 or 1-0 in a Binary Sequence**

**Description**

A function returns how many 0-1 or 1-0 transitions in a binary sequence. For example, 00010000 is 2 transition and 01010100 is 6 transitions. It is internally used in the lbp function.

**Usage**

```r
lbnorm(seq)
```

**Arguments**

- `seq` A sequential 0, 1 array

**See Also**

- lbp

---

```r
image(rot90(c(pol), col=gray(c(0:255)/255), useRaster=TRUE, main="polar map", asp=1))
integ.profile(pol, axis="H", h=c(10, 200), v=c(0, 90), disp=TRUE)
integ.profile(pol, axis="V", h=c(70, 100), v=c(0, 360), disp=TRUE)
```
Details
The LBP operator was originally designed for texture description. The operator assigns a label to every pixel of an image by thresholding the 3x3-neighborhood of each pixel with the center pixel value and considering the result as a binary number (gives 0 if each pixel is smaller than the center, otherwise 1). Then, the histogram of the labels can be used as a texture descriptor. The circular \((8,r=1)\), and \((8,r=2)\) neighborhoods are considered. The function assumes 8-bit grayscale image as an input.

Value
- \texttt{lbp.u2} a matrix (image) returned from requested LBP u2 operation
- \texttt{lbp.ori} a matrix (image) returned from requested LBP operation

Note
A local binary pattern is called uniform if the binary pattern contains at most two 0-1 or 1-0 transitions. In calculation, the histogram has 58 separate bins for uniform patterns, and all other non-uniform patterns are assigned to one single bin. Thus, the length of the features reduces from 256 to 59. This function returned both the lbp patterns before (lbp.ori) and after (lbp.u2) considering the uniform patterns.

References


See Also
\texttt{rgb2bin, hlac}

Examples
```r
## Not run:
data(camphora)
par(mfrow=c(2,2))
r1 <- lbp(camphora,1)
image(rot90c(r1$lbp.u2),col = gray((0:58)/58), main="lbp.u2 (r=1, 8 points)", useRaster=TRUE, asp=1, axes=FALSE)
image(rot90c(r1$lbp.ori),col = gray((0:255)/255), main="lbp.ori (r=1, 8 points)", useRaster=TRUE, asp=1, axes=FALSE)
hist(r1$lbp.u2,breaks=59, main="Histogram of lbp.u2")
hist(r1$lbp.ori,breaks=256, main="Histogram of lbp.ori")

## End(Not run)
```
noise.filter  

Median, Mean and Gaussian Filter

Description

A function provides three kinds of noise reduction on an image, "median", "mean", and "gaussian". A typical pre-processing step to improve the results of later processing for example, glcm-haralick analysis.

Usage

noise.filter(x, n=3, method="median")

Arguments

x  
A raster image or a matrix

n  
filter size is given by n x n. Default is 3 x 3. Number has to be an odd number. For gaussian filter, only 3 or 5 is available.

method  
"median", "mean", and "gaussian" can be selected. Default is "median".

Value

A raster or a matrix

References


See Also

glc

Examples

data(camphora)
camphora <- crop(camphora,200,200)
par(mfrow=c(2,2))
image(rot90c(noise.filter(camphora,3,"median")),col=gray(c(0:255)/255),
main="median", useRaster=TRUE, axes=FALSE, asp=1)
image(rot90c(noise.filter(camphora,3,"mean")),col=gray(c(0:255)/255),
main="mean", useRaster=TRUE, axes=FALSE, asp=1)
image(rot90c(noise.filter(camphora,3,"gaussian")),col=gray(c(0:255)/255),
main="gaussian", useRaster=TRUE, axes=FALSE, asp=1)
**rgb2gray**  
Convert RGB image to Grayscale

**Description**
A function returns grayscale image with coefficients = c(0.3, 0.59, 0.11).

**Usage**
```r
rgb2gray(x, coefs=c(0.30, 0.59, 0.11))
```

**Arguments**
- `x`: A raster image or a matrix
- `coefs`: R, G, B weights. Default are coefs=c(0.30, 0.59, 0.11)

**Value**
A grayscale image

**See Also**
grey2bin

---

**rot90c**  
Transpose Image 90 Degrees Clockwisely

**Description**
Maybe useful to visualize images read by tiff::readTIFF using graphic::image

**Usage**
```r
rot90c(x)
```

**Arguments**
- `x`: A raster or a matrix
### rotate.matrix

**Image Rotation by Bilinear Interpolation**

**Description**

Three methods to execute rotation by 1) assuming values to destination, 2) obtaining values from the source image by inverse rotation with "nearest neighbor (NN)". 3) previous procedure together with "bilinear interpolation". The default is a rotation with "bilinear Interpolation".

**Usage**

```r
rotate.matrix(x, angle, method="bilinear")
```

**Arguments**

- `x`: A raster image or a matrix
- `angle`: Plus(>0) value to request clockwise rotation, while minus for anticlockwise rotation.
- `method`: "simple" assumes values to destination’, "NN" obtains values from the source image by inverse rotation with "nearest neighbor", and "bilinear" performs the same but with "bilinear interpolation" of the source image. value to request clockwise rotation.

**Details**

Assuming 8-bit grayscale image as an input.

**Value**

A matrix after rotation

**See Also**

- `rgb2gray`

**Examples**

```r
data(camphora)
par(mfrow=c(2,2))
r1 <- rotate.matrix(camphora,15, method="simple")
image(rot90c(r1),asp=1,col=grey(c(0:255)/255), main= "simple", useRaster=TRUE, axes=FALSE)
r2 <- rotate.matrix(camphora,25, method="NN")
image(rot90c(r2),asp=1,col=grey(c(0:255)/255), main="nearest neighbour", useRaster=TRUE, axes=FALSE)
r3 <- rotate.matrix(camphora,35, method="bilinear")
image(rot90c(r3),asp=1,col=grey(c(0:255)/255),main="bilinear interpolation", useRaster=TRUE, axes=FALSE)
```
**swap.quad**

*Swapping Quadrants*

**Description**

A function maybe useful to generates power spectrum from fft output.

**Usage**

```r
swap.quad(x, disp=FALSE, reverse=FALSE)
```

**Arguments**

- `x` output of Mod(fft(imagefile))
- `disp` TRUE requests to draw power spectrum
- `reverse` TRUE should be used when power spectrum of N x M, where one of them is odd number.

**Value**

a matrix of power spectrum

**See Also**

fft, Mod

**Examples**

```r
data(camphora)
data(cryptomeria)
img1 <- camphora
img2 <- rgb2gray(cryptomeria)
par(mfrow=c(2,2))

image(rot90c(img1),col=gray(c(0:255)/255), main="Camphora", asp=1, useRaster=TRUE, axes=FALSE)
o.fft <- Mod(fft(img1))
ps <- swap.quad(o.fft)
image(rot90c(log(ps)),col=gray(c(0:255)/255), main="power spectrum", asp=1, useRaster=TRUE, axes=FALSE)

image(rot90c(img2),col=gray(c(0:255)/255), main="Cryptomeria", asp=1, useRaster=TRUE, axes=FALSE)
image(rot90c(log(swap.quad(Mod(fft(img2))))),col=gray(c(0:255)/255),
main="power spectrum", asp=1, useRaster=TRUE, axes=FALSE)
```
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