

# Package ‘EBImage’

October 9, 2015

**Version** 4.10.1

**Title** Image processing and analysis toolbox for R

**Encoding** UTF-8

**Author** Andrzej Oleś, Gregoire Pau, Mike Smith, Oleg Sklyar, Wolfgang Huber, with contributions from Joseph Barry and Philip A. Marais

**Maintainer** Andrzej Oleś <andrzej.oles@embl.de>

**Depends**

**Imports** BiocGenerics (>= 0.7.1), methods, graphics, grDevices, stats, abind, tiff, jpeg, png, locfit, fftwtools (>= 0.9-7)

**Suggests** BiocStyle

**Description** EBImage provides general purpose functionality for image processing and analysis. In the context of (high-throughput) microscopy-based cellular assays, EBImage offers tools to segment cells and extract quantitative cellular descriptors. This allows the automation of such tasks using the R programming language and facilitates the use of other tools in the R environment for signal processing, statistical modeling, machine learning and data visualization.

**License** LGPL

**LazyLoad** true

**biocViews** Visualization

**NeedsCompilation** yes

## R topics documented:

bwlabel . . . . .	2
channel . . . . .	3
colorLabels . . . . .	5
combine . . . . .	6
computeFeatures . . . . .	7
display . . . . .	10
distmap . . . . .	11
drawCircle . . . . .	12

EImage	13
EImage-defunct	16
equalize	16
fillHull	17
filter2	18
floodFill	19
gblur	20
Image	21
io	23
localCurvature	25
medianFilter	27
morphology	28
normalize	30
ocontour	31
otsu	32
paintObjects	33
propagate	34
resize	36
rmObjects	38
stackObjects	39
thresh	41
tile	42
transpose	43
watershed	44
<b>Index</b>	<b>46</b>

---

 bwlabel

*Binary segmentation*


---

### Description

Labels connected (connected sets) objects in a binary image.

### Usage

`bwlabel(x)`

### Arguments

`x` An Image object or an array. `x` is considered as a binary image, whose pixels of value 0 are considered as background ones and other pixels as foreground ones.

### Details

All pixels for each connected set of foreground (non-zero) pixels in `x` are set to a unique increasing integer, starting from 1. Hence, `max(x)` gives the number of connected objects in `x`.

**Value**

A Grayscale Image object or an array, containing the labelled version of *x*.

**Author(s)**

Gregoire Pau, 2009

**See Also**

[computeFeatures](#), [propagate](#), [watershed](#), [paintObjects](#), [colorLabels](#)

**Examples**

```
## simple example
x = readImage(system.file('images', 'shapes.png', package='EBImage'))
x = x[110:512,1:130]
display(x, title='Binary')
y = bwlabel(x)
display(normalize(y), title='Segmented')

## read nuclei images
x = readImage(system.file('images', 'nuclei.tif', package='EBImage'))
display(x)

## computes binary mask
y = thresh(x, 10, 10, 0.05)
y = opening(y, makeBrush(5, shape='disc'))
display(y, title='Cell nuclei binary mask')

## bwlabel
z = bwlabel(y)
display(normalize(z), title='Cell nuclei')
nb nuclei = apply(z, 3, max)
cat('Number of nuclei=', paste(nb nuclei, collapse=', '), '\n')

## recolor nuclei in colors
cols = c('black', sample(rainbow(max(z))))
zrainbow = Image(cols[1+z], dim=dim(z))
display(zrainbow, title='Cell nuclei (recolor)')
```

---

channel

*Color and image color mode conversions*

---

**Description**

channel handles color space conversions between image modes. `rgbImage` combines Grayscale images into a Color one. `toRGB` is a wrapper function for convenient grayscale to RGB color space conversion; the call `toRGB(x)` returns the result of `channel(x, 'rgb')`.

## Usage

```
channel(x, mode)
rgbImage(red, green, blue)
toRGB(x)
```

## Arguments

x	An Image object or an array.
mode	A character value specifying the target mode for conversion. See Details.
red, green, blue	Image objects in Grayscale color mode or arrays of the same dimension. If missing, a black image will be used.

## Details

Conversion modes:

rgb Converts a Grayscale image or an array into a Color image, replicating RGB channels.

gray, grey Converts a Color image into a Grayscale image, using uniform 1/3 RGB weights.

luminance Luminance-preserving Color to Grayscale conversion using CIE 1931 luminance weights:  $0.2126 * R + 0.7152 * G + 0.0722 * B$ .

red, green, blue Extracts the red, green or blue channel from a Color image. Returns a Grayscale image.

asred, asgreen, asblue Converts a Grayscale image or an array into a Color image of the specified hue.

NOTE: channel changes the pixel intensities, unlike colorMode which just changes the way that EBIImage renders an image.

## Value

An Image object or an array.

## Author(s)

Oleg Sklyar, <osklyar@ebi.ac.uk>

## See Also

[colorMode](#)

## Examples

```
x = readImage(system.file("images", "shapes.png", package="EBImage"))
display(x)
y = channel(x, 'asgreen')
display(y)
```

```
## rgbImage
x = readImage(system.file('images', 'nuclei.tif', package='EBImage'))
y = readImage(system.file('images', 'cells.tif', package='EBImage'))
display(x, title='Cell nuclei')
display(y, title='Cell bodies')

cells = rgbImage(green=1.5*y, blue=x)
display(cells, title='Cells')
```

---

colorLabels

*Color Code Labels*

---

### Description

Color codes the labels of object masks by a random permutation.

### Usage

```
colorLabels(x, normalize = TRUE)
```

### Arguments

x	an Image object in Grayscale color mode or an array containing object masks. Object masks are sets of pixels with the same unique integer value
normalize	if TRUE normalizes the resulting color image

### Details

Performs color coding of object masks, which are typically obtained using the `bwlabel` function. Each label from `x` is assigned an unique color. The colors are distributed among the labels using a random permutation. If `normalize` is set to `TRUE` the intensity values of the resulting image are mapped to the `[0,1]` range.

### Value

An Image object containing color coded objects of `x`.

### Author(s)

Bernd Fischer, Andrzej Oles, 2013-2014

### See Also

[bwlabel](#), [normalize](#)

## Examples

```
x = readImage(system.file('images', 'shapes.png', package='EImage'))
x = x[110:512,1:130]
y = bwlabel(x)
z = colorLabels(y)
display(z, title='Colored segmentation')
```

---

combine

*Combining images*

---

## Description

Merges images to create image sequences.

## Usage

```
combine(x, y, ...)
```

## Arguments

x	An Image object, an array, or a list of Image objects and arrays.
y	An Image object or an array.
...	Image objects or arrays.

## Details

The function `combine` uses `abind` to merge multi-dimensional arrays along the dimension depending on the color mode of `x`. If `x` is a Grayscale image or an array, image objects are combined along the third dimension, whereas when `x` is a Color image they are combined along the fourth dimension, leaving room on the third dimension for color channels.

## Value

An Image object or an array.

## Author(s)

Gregoire Pau, Andrzej Oles, 2013

## See Also

[Image](#)

**Examples**

```
## combination of color images
img = readImage(system.file("images", "sample-color.png", package="EBImage"))[257:768,,]
x = combine(img, flip(img), flop(img))
display(x, all=TRUE)

## Blurred images
x = resize(img, 128, 128)
xt = list()
for (t in seq(0.1, 5, len=9)) xt=c(xt, list(gblur(x, s=t)))
xt = combine(xt)
display(xt, title='Blurred images', all=TRUE)
```

---

computeFeatures	<i>Compute object features</i>
-----------------	--------------------------------

---

**Description**

Computes morphological and texture features from image objects.

**Usage**

```
computeFeatures(x, ref, methods.noref=c("computeFeatures.moment", "computeFeatures.shape"),
  methods.ref=c("computeFeatures.basic", "computeFeatures.moment", "computeFeatures.haralick"),
  xname="x", refnames, properties=FALSE, expandRef=standardExpandRef, ...)
computeFeatures.basic(x, ref, properties=FALSE, basic.quantiles=c(0.01, 0.05, 0.5, 0.95, 0.99), xs, .
computeFeatures.shape(x, properties=FALSE, xs, ...)
computeFeatures.moment(x, ref, properties=FALSE, xs, ...)
computeFeatures.haralick(x, ref, properties=FALSE, haralick.nbins=32, haralick.scales=c(1, 2), xs, .
standardExpandRef(ref, refnames)
```

**Arguments**

x	An Image object or an array containing labelled objects. Labelled objects are pixel sets with the same unique integer value.
ref	A matrix or a list of matrices, containing the intensity values of the reference objects.
methods.noref	A character vector containing the function names to be called to compute features without reference intensities. Default is computeFeatures.moment and computeFeatures.shape.
methods.ref	A character vector containing the function names to be called to compute features with reference intensities. Default is computeFeatures.basic, computeFeatures.moment and computeFeatures.haralick.
xname	A character string naming the object layer. Default is x.
refnames	A character vector naming the reference intensity layers. Default are the names of ref, if present. If not, reference intensity layers are named using lower-case letters.

<code>properties</code>	A logical. If FALSE, the default, the function returns the feature matrix. If TRUE, the function returns feature properties.
<code>expandRef</code>	A function used to expand the reference images. Default is <code>standardExpandRef</code> . See Details.
<code>basic.quantiles</code>	A numerical vector indicating the quantiles to compute.
<code>haralick.nbins</code>	An integer indicating the number of bins using to compute the Haralick matrix. See Details.
<code>haralick.scales</code>	A integer vector indicating the number of scales to use to compute the Haralick features.
<code>xs</code>	An optional temporary object created by <code>computeFeatures</code> used for performance considerations.
<code>...</code>	Optional arguments passed to the feature computation functions.

### Details

Features are named `x.y.f`, where `x` is the object layer, `y` the reference image layer and `f` the feature name. Examples include `cell.dna.mean`, indicating mean DNA intensity computed in the cell or `nucleus.tubulin.cx`, indicating the `x` center of mass of tubulin computed in the nucleus region.

The function `computeFeatures` computes sets of features. Features are organized in 4 sets, each computed by a different function. The function `computeFeatures.basic` computes spatial-independent statistics on pixel intensities:

- `b.mean`: mean intensity
- `b.sd`: standard deviation intensity
- `b.mad`: mad intensity
- `b.q*`: quantile intensity

The function `computeFeatures.shape` computes features that quantify object shape:

- `s.area`: area size (in pixels)
- `s.perimeter`: perimeter (in pixels)
- `s.radius.mean`: mean radius (in pixels)
- `s.radius.sd`: standard deviation of the mean radius (in pixels)
- `s.radius.max`: max radius (in pixels)
- `s.radius.min`: min radius (in pixels)

The function `computeFeatures.moment` computes features related to object image moments, which can be computed with or without reference intensities:

- `m.cx`: center of mass `x` (in pixels)
- `m.cy`: center of mass `y` (in pixels)
- `m.majoraxis`: elliptical fit major axis (in pixels)
- `m.eccentricity`: elliptical eccentricity defined by  $\sqrt{1 - \text{minoraxis}^2 / \text{majoraxis}^2}$ . Circle eccentricity is 0 and straight line eccentricity is 1.



- m.theta: object angle (in radians)

The function `computeFeatures.haralick` computes features that quantify pixel texture. Features are named according to Haralick's original paper.

### Value

If `properties` is FALSE (by default), `computeFeatures` returns a matrix of `n` cells times `p` features, where `p` depends of the options given to the function. Returns NULL if no object is present.

If `properties` is TRUE, `computeFeatures` returns a matrix of `p` features times 2 properties (translation and rotation invariance). Feature properties are useful to filter out features that may not be needed for specific tasks, e.g. cell position when doing cell classification.

### Author(s)

Gregoire Pau, <gregoire.pau@embl.de>, 2011

### References

R. M. Haralick, K Shanmugam and Its' Hak Deinstein (1979). *Textural Features for Image Classification*. IEEE Transactions on Systems, Man and Cybernetics.

### See Also

[bwlabel](#), [propagate](#)

### Examples

```
## load and segment nucleus
y = readImage(system.file("images", "nuclei.tif", package="EBImage"))[, , 1]
x = thresh(y, 10, 10, 0.05)
x = opening(x, makeBrush(5, shape='disc'))
x = bwlabel(x)
display(y, title="Cell nuclei")
display(x, title="Segmented nuclei")

## compute shape features
fts = computeFeatures.shape(x)
fts

## compute features
ft = computeFeatures(x, y, xname="nucleus")
cat("median features are:\n")
apply(ft, 2, median)

## compute feature properties
ftp = computeFeatures(x, y, properties=TRUE, xname="nucleus")
ftp
```

---

display	<i>Image Display</i>
---------	----------------------

---

**Description**

Display images using an interactive JavaScript viewer or R's built-in graphics capabilities.

**Usage**

```
display(x,
        title = deparse(substitute(x), width.cutoff = 500L, nlines = 1),
        method,
        frame, all = FALSE)
```

**Arguments**

<code>x</code>	an Image object or an array.
<code>title</code>	a character string used as a window title.
<code>method</code>	the method used to display images, can be <code>browser</code> or <code>raster</code> . Defaults to <code>browser</code> when R is used interactively and <code>raster</code> otherwise. The default behavior can be overridden by setting <code>options("EBImage.display")</code> . See Details.
<code>frame</code>	a numeric indicating the frame number; only works in conjunction with <code>method = "raster"</code> and <code>all = FALSE</code> .
<code>all</code>	should all frames of a stacked image be displayed, or just a single frame?

**Details**

The default method used for displaying images depends on whether called from an interactive R session. If `interactive()` returns `TRUE` images are displayed using the `"browser"` method, otherwise the `"raster"` method is used. This behavior can be overridden and the default display method fixed regardless of the interactivity by setting `options("EBImage.display")` to either `"browser"` or `"raster"`.

The `"browser"` method runs an interactive JavaScript image viewer using the default web browser. Multiple windows or tabs can be opened in this way. Pressing `'h'` displays a list of available features along with corresponding mouse and keyboard actions.

The `"raster"` method displays images as raster graphics using R's built-in functions. By default only the first frame of a stacked image is shown; a different frame can be specified using the `frame` argument. When `all=TRUE` all frames are rendered and automatically positioned next to each other in a grid. The user coordinates of the plotting region are set to the image pixel coordinates with the origin  $(0, 0)$  in the upper left corner.

**Value**

Invisible `NULL`.

**Note**

For the "browser" method a compatible web browser with JavaScript enabled is required (e.g. Mozilla Firefox).

**Author(s)**

Andrzej Oles, <andrzej.oles@embl.de>, 2012

**References**

[Mozilla Firefox](#)

**Examples**

```
## Display a single image
x = readImage(system.file("images", "sample-color.png", package="EBImage"))[257:768,,]
display(x, "Sample")

## Display a thresholded sequence ...
y = readImage(system.file("images", "sample.png", package="EBImage"))
yt = list()
for (t in seq(0.1, 5, len=9)) yt=c(yt, list(gblur(y, s=t)))
yt = combine(yt)

## ... using the browser viewer ...
display(yt, "Blurred images")

## ... or using R's build-in raster functions
display(resize(yt, 256, 256), method = "raster", all = TRUE)

## Display the last frame
display(yt, method = "raster", frame = numberOfFrames(yt, type = "render"))
```

---

distmap

*Distance map transform*

---

**Description**

Computes the distance map transform of a binary image. The distance map is a matrix which contains for each pixel the distance to its nearest background pixel.

**Usage**

```
distmap(x, metric=c('euclidean', 'manhattan'))
```

**Arguments**

x	An Image object or an array. x is considered as a binary image, whose pixels of value 0 are considered as background ones and other pixels as foreground ones.
metric	A character indicating which metric to use, L1 distance (manhattan) or L2 distance (euclidean). Default is euclidean.

**Details**

A fast algorithm of complexity  $O(M*N*\log(\max(M,N)))$ , where (M,N) are the dimensions of x, is used to compute the distance map.

**Value**

An Image object or an array, with pixels containing the distances to the nearest background points.

**Author(s)**

Gregoire Pau, <gpau@ebi.ac.uk>, 2008

**References**

M. N. Kolountzakis, K. N. Kutulakos. Fast Computation of the Euclidean Distance Map for Binary Images, Infor. Proc. Letters 43 (1992).

**Examples**

```
x = readImage(system.file("images", "shapes.png", package="EBImage"))
display(x)
dx = distmap(x)
display(dx/10, title='Distance map of x')
```

---

drawCircle

*Draw a circle on an image.*

---

**Description**

Draw a circle on an image.

**Usage**

```
drawCircle(img, x, y, radius, col, fill=FALSE, z=1)
```

**Arguments**

<code>img</code>	An Image object or an array.
<code>x, y, radius</code>	numerics indicating the center and the radius of the circle.
<code>col</code>	A numeric or a character string specifying the color of the circle.
<code>fill</code>	A logical indicating whether the circle should be filled. Default is FALSE.
<code>z</code>	A numeric indicating on which frame of the image the circle should be drawn. Default is 1.

**Value**

An Image object or an array, containing the transformed version of `img`.

**Author(s)**

Gregoire Pau, 2010

**Examples**

```
## Simple white circle
x = matrix(0, nrow=300, ncol=300)
y = drawCircle(x, 100, 200, 47, col=1)
display(y)

## Simple filled yellow circle
x = channel(y, 'rgb')
y = drawCircle(x, 200, 140, 57, col='yellow', fill=TRUE)
display(y)
```

---

EBImage

*Package overview*

---

**Description**

EBImage is an image processing and analysis package for R. Its primary goal is to enable automated analysis of large sets of images such as those obtained in high throughput automated microscopy.

EBImage relies on the Image object to store and process images but also works on multi-dimensional arrays.

**Package content**

Image methods

- Image
- `as.Image`, `is.Image`, `as.raster`
- `colorMode`, `imageData`
- `getFrame`, `numberOfFrames`

#### Image I/O, display

- readImage, writeImage
- display
- image

#### Spatial transforms

- resize, flip, flop, transpose
- rotate, translate, affine

#### Image segmentation, objects manipulation

- thresh, bwlabel, otsu
- watershed, propagate
- ocontour
- paintObjects, rmObjects, reenumerate

#### Image enhancement, filtering

- normalize
- filter2, gblur, medianFilter

#### Morphological operations

- makeBrush
- erode, dilate, opening, closing
- erodeGreyScale, dilateGreyScale, openingGreyScale, closingGreyScale
- whiteTopHatGreyScale, blackTopHatGreyScale, selfcomplementaryTopHatGreyScale
- distmap
- floodFill, fillHull

#### Color space manipulation

- rgbImage, channel, toRGB

#### Image stacking, combining, tiling

- stackObjects
- combine
- tile, untile

#### Drawing on images

- drawCircle

#### Features extraction

- computeFeatures
- computeFeatures.basic, computeFeatures.moment, computeFeatures.shape, computeFeatures.haralick

- standardExpandRef

#### Defunct

- blur, equalize
- drawtext, drawfont
- getFeatures, hullFeatures, zernikeMoments
- edgeProfile, edgeFeatures,
- haralickFeatures, haralickMatrix
- moments, cmoments, smoments, rmoments

#### Authors

Oleg Sklyar, <osklyar@ebi.ac.uk>, Copyright 2005-2007

Gregoire Pau, <gpau@ebi.ac.uk>

Wolfgang Huber, <huber@ebi.ac.uk>

Andrzej Oles, <andrzej.oles@embl.de>

Mike Smith, <msmith@ebi.ac.uk>

European Bioinformatics Institute  
European Molecular Biology Laboratory  
Wellcome Trust Genome Campus  
Hinxton  
Cambridge CB10 1SD  
UK

The code of [propagate](#) is based on the CellProfiler with permission granted to distribute this particular part under LGPL, the corresponding copyright (Jones, Carpenter) applies.

The source code is released under LGPL (see the LICENSE file in the package root for the complete license wording).

This library is free software; you can redistribute it and/or modify it under the terms of the GNU Lesser General Public License as published by the Free Software Foundation; either version 2.1 of the License, or (at your option) any later version. This library is distributed in the hope that it will be useful, but WITHOUT ANY WARRANTY; without even the implied warranty of MERCHANTABILITY or FITNESS FOR A PARTICULAR PURPOSE.

See the GNU Lesser General Public License for more details. For LGPL license wording see <http://www.gnu.org/licenses/lgpl.html>

#### Examples

```
example(readImage)
example(display)
example(rotate)
example(propagate)
```

---

EImage-defunct	<i>EImage defunct functions</i>
----------------	---------------------------------

---

### Description

These following functions are defunct and will be removed in the next Bioconductor release.

### Usage

```
blur(...)
drawtext(...)
drawfont(...)
getFeatures(...)
hullFeatures(...)
zernikeMoments(...)
edgeProfile(...)
edgeFeatures(...)
haralickFeatures(...)
haralickMatrix(...)
moments(...)
rmoments(...)
smoments(...)
cmoments(...)
```

### Arguments

... Defunct arguments.

---

equalize	<i>Histogram Equalization</i>
----------	-------------------------------

---

### Description

Equalize the image histogram to a specified range and number of levels.

### Usage

```
equalize(x, range = c(0, 1), levels = 256)
```

### Arguments

x	an Image object or an array
range	numeric vector of length 2, the output range of the equalized histogram
levels	number of grayscale levels (Grayscale images) or intensity levels of each channel (Color images)



**Details**

Individual channels of Color images and frames of image stacks are equalized separately.

**Value**

An Image object or an array, containing the transformed version of x.

**Author(s)**

Andrzej Oles, <andrzej.oles@embl.de>, 2014

**Examples**

```
x = readImage(system.file('images', 'cells.tif', package='EBImage'))
hist(x)
y = equalize(x)
hist(y)
display(y, title='Equalized Grayscale Image')

x = readImage(system.file('images', 'sample-color.png', package='EBImage'))
hist(x)
y = equalize(x)
hist(y)
display(y, title='Equalized Grayscale Image')
```

---

fillHull

*Fill holes in objects*

---

**Description**

Fill holes in objects.

**Usage**

```
fillHull(x)
```

**Arguments**

x                    An Image object or an array.

**Details**

fillHull fills holes in the objects defined in x, where objects are sets of pixels with the same unique integer value.

**Value**

An Image object or an array, containing the transformed version of x.

**Author(s)**

Gregoire Pau, Oleg Sklyar; 2007

**See Also**

[bwlabel](#)

**Examples**

```
x = readImage(system.file('images', 'nuclei.tif', package='EBImage'))
display(x)

y = thresh(x, 10, 10, 0.05)
display(y, title='Cell nuclei')

y = fillHull(y)
display(y, title='Cell nuclei without holes')
```

---

filter2

*2D Convolution Filter*

---

**Description**

Filters an image using the fast 2D FFT convolution product.

**Usage**

```
filter2(x, filter)
```

**Arguments**

x	An Image object or an array.
filter	An Image object or an array, with odd spatial dimensions. Must contain only one frame.

**Details**

Linear filtering is useful to perform low-pass filtering (to blur images, remove noise...) and high-pass filtering (to detect edges, sharpen images). The function `makeBrush` is useful to generate filters.

Data is reflected around borders.

If x contains multiple frames, the filter will be applied one each frame.

**Value**

An Image object or an array, containing the filtered version of x.

**Author(s)**

Gregoire Pau, <gpau@ebi.ac.uk>

**See Also**

[makeBrush](#), [convolve](#), [fft](#), [blur](#)

**Examples**

```
x = readImage(system.file("images", "sample-color.png", package="EBImage"))
display(x, title='Sample')

## Low-pass disc-shaped filter
f = makeBrush(21, shape='disc', step=FALSE)
display(f, title='Disc filter')
f = f/sum(f)
y = filter2(x, f)
display(y, title='Filtered image')

## High-pass Laplacian filter
la = matrix(1, nc=3, nr=3)
la[2,2] = -8
y = filter2(x, la)
display(y, title='Filtered image')
```

---

floodFill

*Region filling*

---

**Description**

Fill regions in images.

**Usage**

```
floodFill(x, pt, col, tolerance=0)
```

**Arguments**

x	An Image object or an array.
pt	Coordinates of the start filling point.
col	Fill color. This argument should be a numeric for Grayscale images and an R color for Color images.
tolerance	Color tolerance used during the fill.

**Details**

Flood fill is performed using the fast scan line algorithm. Filling starts at pt and grows in connected areas where the absolute difference of the pixels intensities (or colors) remains below tolerance.

**Value**

An Image object or an array, containing the transformed version of x.

**Author(s)**

Gregoire Pau, Oleg Sklyar; 2007

**Examples**

```
x = readImage(system.file("images", "shapes.png", package="EBImage"))
y = floodFill(x, c(67, 146), 0.5)
display(y)
```

```
y = channel(y, 'rgb')
y = floodFill(y, c(48, 78), 'red')
y = floodFill(y, c(156, 52), 'orange')
display(y)
```

```
x = readImage(system.file("images", "sample.png", package="EBImage"))
y = floodFill(x, c(226, 121), 1, tolerance=0.1)
display(y)
```

---

gblur

*Low-pass Gaussian filter*

---

**Description**

Filters an image with a low-pass Gaussian filter.

**Usage**

```
gblur(x, sigma, radius = 2 * ceiling(3 * sigma) + 1)
```

**Arguments**

x	An Image object or an array.
sigma	A numeric denoting the standard deviation of the Gaussian filter used for blurring.
radius	The radius of the filter in pixels. Default is $2 * \text{ceiling}(3 * \text{sigma}) + 1$ .

**Details**

The Gaussian filter is created with the function `makeBrush`.

**Value**

An Image object or an array, containing the filtered version of x.

**Author(s)**

Oleg Sklyar, <osklyar@ebi.ac.uk>, 2005-2007

**See Also**

[filter2](#), [makeBrush](#)

**Examples**

```
x = readImage(system.file("images", "sample.png", package="EBImage"))
display(x)

y = gblur(x, sigma=8)
display(y, title='gblur(x, sigma=8)')
```

---

Image

*Image class*

---

**Description**

EBImage uses the Image class to store and process images. Images are stored as multi-dimensional arrays containing the pixel intensities. Image extends the base class array and uses the colormode slot to store how the color information of the multi-dimensional data is handled.

The colormode slot can be either Grayscale or Color. In either mode, the first two dimensions of the underlying array are understood to be the spatial dimensions of the image. In the Grayscale mode the remaining dimensions contain other image frames. In the Color mode, the third dimension contains color channels of the image, while higher dimensions contain image frames. The number of channels is not limited and can be any number  $\geq 1$ ; these can be, for instance, the red, green, blue and, possibly, alpha channel. Note that grayscale images containing an alpha channel are stored with colormode=Color.

All methods from the EBImage package work either with Image objects or multi-dimensional arrays. In the latter case, the color mode is assumed to be Grayscale.

**Usage**

```
Image(data, dim, colormode)
as.Image(x)
is.Image(x)

## S3 method for class 'Image'
as.array(x, ...)
## S3 method for class 'Image'
as.raster(x, i = 1, ...)

colorMode(y)
colorMode(y) <- value
```

```

imageData(y)
imageData(y) <- value

getFrame(y, i, type = c('total', 'render'))
getFrames(y, i, type = c('total', 'render'))
numberOfFrames(y, type = c('total', 'render'))

```

### Arguments

<code>data</code>	A vector or array containing the pixel intensities of an image. If missing, the default 1x1 zero-filled array is used.
<code>dim</code>	A vector containing the final dimensions of an Image object. If missing, equals to <code>dim(data)</code> .
<code>colormode</code>	A numeric or a character string containing the color mode which can be either Grayscale or Color. If missing, equals to Grayscale.
<code>x</code>	An R object.
<code>y</code>	An Image object or an array.
<code>i</code>	Number(s) of frame(s). A single number in case of <code>getFrame</code> , or a vector of frame numbers for <code>getFrames</code> . If missing all frames are returned.
<code>value</code>	For <code>colorMode</code> , a numeric or a character string containing the color mode which can be either Grayscale or Color. For <code>imageData</code> , an Image object or an array.
<code>type</code>	A character string containing total or render. Default is total.
<code>...</code>	further arguments passed to or from other methods.

### Details

Depending on `type`, `numberOfFrames` returns the total number of frames contained in the object `y` or the number of rendered frames. The total number of frames is independent of the color mode and equals to the product of all the dimensions except the two first ones. The number of rendered frames is equal to the total number of frames in the Grayscale color mode, or to the product of all the dimensions except the three first ones in the Color color mode.

`getFrame` returns the *i*-th frame contained in the image `y`. If `type` is `total`, the function is unaware of the color mode and returns an *xy*-plane. For `type=render`, the function returns the *i*-th image as shown by the `display` function.

### Value

`Image` and `as.Image` return a new Image object.

`is.Image` returns TRUE if `x` is an Image object and FALSE otherwise.

`as.raster` coerces an Image object to its raster representation. For stacked images the *i*-th frame is returned (by default the first one).

`colorMode` returns the color mode of `y` and `colorMode<-` changes the color mode of `y`.

`imageData` returns the array contained in an Image object.

**Author(s)**

Oleg Sklyar, <osklyar@ebi.ac.uk>, 2005-2007

**See Also**

[readImage](#), [writeImage](#), [display](#)

**Examples**

```
s1 = exp(12i*pi*seq(-1, 1, length=300)^2)
y = Image(outer(Im(s1), Re(s1)))
display(normalize(y))

x = Image(rnorm(300*300*3),dim=c(300,300,3), colormode='Color')
display(x)

w = matrix(seq(0, 1, len=300), nc=300, nr=300)
m = abind::abind(w, t(w), along=3)
z = Image(m, colormode='Color')
display(normalize(z))

y = Image(c('red', 'violet', '#ff51a5', 'yellow'), dim=c(71, 71))
display(y)

## colorMode example
x = readImage(system.file('images', 'nuclei.tif', package='EBImage'))
x = x[, ,1:3]
display(x, title='Cell nuclei')
colorMode(x) = Color
display(x, title='Cell nuclei in RGB')
```

---

io

*Image I/O*

---

**Description**

Read and write images from/to files and URLs.

**Usage**

```
readImage(files, type, all = TRUE, names = sub("\\.[^.]*$", "", basename(files)), ...)
writeImage(x, files, type, quality = 100, bits.per.sample, compression = "none", ...)
```

**Arguments**

**files** a character vector of file names or URLs.

**type** image type (optional). Supported values are: jpeg, png, and tiff. If missing, file format is automatically determined by file name extension.

<code>all</code>	logical: when the file contains more than one image should all frames be read, or only the first one?
<code>names</code>	a character vector used for frame names. Should have the same length as <code>files</code> .
<code>x</code>	an Image object or an array.
<code>bits.per.sample</code>	a numeric scalar specifying the number of bits per sample (only for <code>tiff</code> files). Supported values are 8 and 16.
<code>compression</code>	the desired compression algorithm (only for <code>tiff</code> files). For a list of supported values consult the documentation of the <code>writeTIFF</code> function from the <code>tiff</code> package.
<code>quality</code>	a numeric ranging from 1 to 100 (default) controlling the quality of the JPEG output.
<code>...</code>	arguments passed to the corresponding functions from the <code>jpeg</code> , <code>png</code> , and <code>tiff</code> packages.

### Details

`readImage` loads all images from the `files` vector and returns them stacked into a single Image object containing an array of doubles ranging from 0 (black) to 1 (white). All images need to be of the same type and have the same dimensions and color mode. If type is missing, the appropriate file format is determined from file name extension. Color mode is determined automatically based on the number of channels. When the function fails to read an image it skips to the next element of the `files` vector issuing a warning message. Non-local files can be read directly from a valid URL.

`writeImage` writes images into files specified by `files`, were the number of files needs to be equal 1 or the number of frames. Given an image containing multiple frames and a single file name either the whole stack is written into a single TIFF file, or each frame is saved to an individual JPEG/PNG file (for `files = "image.*"` frames are saved into `image-X.*` files, where X equals the frame number less one; for an image containing n frames this results in file names numbered from 0 to n-1).

When writing JPEG files the compression quality can be specified using `quality`. Valid values range from 100 (highest quality) to 1 (lowest quality). For TIFF files additional information about the desired number of bits per sample (`bits.per.sample`) and the compression algorithm (`compression`) can be provided. For a complete list of supported values please consult the documentation of the `tiff` package.

### Value

`readImage` returns a new Image object.

`writeImage` returns an invisible vector of file names.

### Note

Image formats have a limited dynamic range (e.g. JPEG: 8 bit, TIFF: 16 bit) and `writeImage` may cause some loss of accuracy. In specific, writing 16 bit image data to formats other than TIFF will strip the 8 LSB. When writing TIFF files a dynamic range check is performed and an appropriate value of `bits.per.sample` is set automatically.



**Author(s)**

Andrzej Oles, <andrzej.oles@embl.de>, 2012

**See Also**

[Image](#), [display](#), [readJPEG/writeJPEG](#), [readPNG/writePNG](#), [readTIFF/writeTIFF](#)

**Examples**

```
## Read and display an image
f = system.file("images", "sample-color.png", package="EBImage")
x = readImage(f)
display(x)

## Read and display a multi-frame TIFF
y = readImage(system.file("images", "nuclei.tif", package="EBImage"))
display(y)

## Read an image directly from a remote location by specifying its URL
try({
  im = readImage("http://www-huber.embl.de/EBImage/ExampleImages/berlin.tif")
  display(im, title = "Berlin Impressions")
})

## Convert a PNG file into JPEG
tempfile = tempfile("", , ".jpeg")
writeImage(x, tempfile, quality = 85)
cat("Converted '", f, "' into '", tempfile, "'.\n", sep="")

## Save a frame sequence
files = writeImage(y, tempfile("", , ".jpeg"), quality = 85)
cat("Files created: ", files, sep="\n")
```

---

localCurvature

*Local Curvature*

---

**Description**

Computes signed curvature along a line.

**Usage**

```
localCurvature(x, h, maxk)
```

**Arguments**

x	A data frame or matrix of dimensions $N \times 2$ containing the coordinates of the line, where $N$ is the number of points. The points should be ordered according to their position on the line. The columns should contain the x and y coordinates. The curvature calculation is unaffected by any permutation of the columns. Directly accepts a list element from <code>ocontour</code> .
h	Specifies the length of the smoothing window. See <code>locfit::lp</code> for more details.
maxk	See <code>locfit::locfit.raw</code> for details.

**Details**

`localCurvature` fits a local non-parametric smoothing line (polynomial of degree 2) at each point along the line segment, and computes the curvature locally using numerical derivatives.

**Value**

Returns a list containing the contour coordinates `x`, the signed curvature at each point `curvature` and the arc length of the contour `length`.

**Author(s)**

Joseph Barry, Wolfgang Huber, 2013

**See Also**

[ocontour](#)

**Examples**

```
## curvature goes as the inverse of the radius of a circle
range=seq(3.5,33.5,by=2)
plotRange=seq(0.5,33.5,length=100)
circleRes=array(dim=length(range))
names(circleRes)=range
for (i in seq_along(1:length(range))) {
  y=as.Image(makeBrush('disc', size=2*range[i]))
  y=ocontour(y)[[1]]
  circleRes[i]=abs(mean(localCurvature(x=y,h=range[i])$curvature, na.rm=TRUE))
}
plot(range, circleRes, ylim=c(0,max(circleRes, na.rm=TRUE)), xlab='Circle Radius', ylab='Curvature', type='p',
points(plotRange, 1/plotRange, type='l')
```

```
## Calculate curvature
x = readImage(system.file("images", "shapes.png", package="EBImage"))[25:74, 60:109]
x = resize(x, 200)
y = gblur(x, 3) > .3
display(y)

contours = ocontour(bwlabel(y))
c = localCurvature(x=contours[[1]], h=11)
```

```
i = c$curvature >= 0
pos = neg = array(0, dim(x))
pos[c$contour[i,]+1] = c$curvature[i]
neg[c$contour[!i,]+1] = -c$curvature[!i]
display(10*(rgbImage(pos, , neg)), title = "Image curvature")
```

---

**medianFilter***2D constant time median filtering*

---

### Description

Filters a 16-bit image using Perreault's modern constant time median filtering algorithm [1].

### Usage

```
medianFilter(x, size, cacheSize=512)
```

### Arguments

x	An Image object or an array.
size	The sizelength of the square median filter in units of pixels.
cacheSize	The L2 cache size of the system CPU in kB.

### Details

Median filtering is useful as a smoothing technique, e.g. in the removal of speckling noise.

If x contains multiple frames, the filter will be applied on each frame.

### Value

An Image object or an array, containing the filtered version of x.

### Author(s)

Joseph Barry, <joseph.barry@embl.de>, 2012

### References

[1] S. Perreault and P. Hebert, "Median Filtering in Constant Time", IEEE Trans Image Process 16(9), 2389-2394, 2007

### See Also

[makeBrush](#), [fft](#), [gblur](#)

**Examples**

```
x = readImage(system.file("images", "nuclei.tif", package="EBImage"))
display(x, title='Nuclei')
y = medianFilter(x, 5)
display(y, title='Filtered nuclei')
```

---

morphology

---

*Perform morphological operations on images*


---

**Description**

Functions to perform morphological operations on binary images.

**Usage**

```
dilate(x, kern)
erode(x, kern)
opening(x, kern)
closing(x, kern)
dilateGreyScale(x, kern)
erodeGreyScale(x, kern)
openingGreyScale(x, kern)
closingGreyScale(x, kern)
whiteTopHatGreyScale(x, kern)
blackTopHatGreyScale(x, kern)
selfcomplementaryTopHatGreyScale(x, kern)
```

```
makeBrush(size, shape=c('box', 'disc', 'diamond', 'Gaussian', 'line'), step=TRUE, sigma=0.3, angle=45)
```

**Arguments**

x	An Image object or an array. x is considered as a binary image, whose pixels of value 0 are considered as background ones and other pixels as foreground ones.
kern	An Image object or an array, containing the structuring element. kern is considered as a binary image, whose pixels of value 0 are considered as background ones and other pixels as foreground ones.
size	A numeric containing the size of the brush in pixels. This should be an odd number; even numbers are rounded to the next odd one, i.e., size = 4 has the same effect as size = 5. If shape is line, size represents the length of the line.
shape	A character vector indicating the shape of the brush. Can be box, disc, diamond, Gaussian or line. Default is box.
step	a logical indicating if the brush is binary. Default is TRUE. The argument is relevant only for the disc and diamond shapes.
sigma	An optional numeric containing the standard deviation of the Gaussian shape. Default is 0.3.

`angle` An optional numeric containing the angle at which the line should be drawn. The angle is one between the top of the image and the line.

### Details

`dilate` applies the mask positioning its center over every background pixel (0), every pixel which is not covered by the mask is reset to foreground (1).

`erode` applies the mask positioning its center over every foreground pixel (!=0), every pixel which is not covered by the mask is reset to background (0).

`opening` is an erosion followed by a dilation and `closing` is a dilation followed by an erosion. The same goes for the grayscale versions.

`dilateGreyScale` applies the mask positioning its center over every pixel of the Image, the output value of the pixel is the maximum value of the Image covered by the mask.

`erodeGreyScale` applies the mask positioning its center over every pixel of the Image, the output value of the pixel is the minimum value of the Image covered by the mask.

`whiteTopHatGreyScale` subtracts the opening of the Image from the Image

`blackTopHatGreyScale` subtracts the Image from the closing of the Image

`selfcomplementaryTopHatGreyScale` is the sum of a white top-hat and a black top-hat, simplified the difference between closing and opening of the Image

`makeBrush` generates brushes of various sizes and shapes that can be used as structuring elements.

Operations on grayscale images use an implementation of the Urbach-Wilkinson algorithm[1] and can only handle flat (i.e. binary) brushes.

### Value

`dilate`, `erode`, `opening`, `closing`, `dilateGreyScale`, `erodeGreyScale`, `openingGreyScale`, `closingGreyScale`, `whiteTopHatGreyScale`, `blackTopHatGreyScale` and `selfcomplementaryTopHatGreyScale` return the transformed Image object or array, after the corresponding morphological operation.

`makeBrush` generates a 2D matrix containing the desired brush.

### Author(s)

Oleg Sklyar, <osklyar@ebi.ac.uk>, 2006 Ilia Kats, <ilia-kats@gmx.net>, 2012

### References

[1] E. R. Urbach and M.H.F. Wilkinson, "Efficient 2-D grayscale morphological transformations with arbitrary flat structuring elements", IEEE Trans Image Process 17(1), 1-8, 2008

### Examples

```
x = readImage(system.file("images", "shapes.png", package="EBImage"))
kern = makeBrush(5, shape='diamond')

display(x)
display(kern, title='Structuring element')
```

```
display(erode(x, kern), title='Erosion of x')
display(dilate(x, kern), title='Dilatation of x')

## makeBrush
display(makeBrush(99, shape='diamond'))
display(makeBrush(99, shape='disc', step=FALSE))
display(2000*makeBrush(99, shape='Gaussian', sigma=10))
```

---

normalize

*Intensity values linear scaling*

---

### Description

Linearly scale the intensity values of an image to a specified range.

### Usage

```
## S4 method for signature 'Image'
normalize(object, separate=TRUE, ft=c(0,1), inputRange)

## S4 method for signature 'array'
normalize(object, separate=TRUE, ft=c(0,1), inputRange)
```

### Arguments

object	an Image object or an array
separate	if TRUE, normalizes each frame separately
ft	a numeric vector of 2 values, target minimum and maximum intensity values after normalization
inputRange	a numeric vector of 2 values, sets the range of the input intensity values; values exceeding this range are clipped

### Details

normalize performs linear interpolation of the intensity values of an image to the specified range ft. If inputRange is not set the whole dynamic range of the image is used as input. By specifying inputRange the input intensity range of the image can be limited to [min, max]. Values exceeding this range are clipped, i.e. intensities lower/higher than min/max are set to min/max.

### Value

An Image object or an array, containing the transformed version of object.

### Author(s)

Oleg Sklyar, <osklyar@ebi.ac.uk>, 2006-2007 Andrzej Oles, <andrzej.oles@embl.de>, 2013

## Examples

```
x = readImage(system.file('images', 'shapes.png', package='EBImage'))
x = x[110:512,1:130]
y = bwlabel(x)
display(x, title='Original')

print(range(y))
y = normalize(y)
print(range(y))
display(y, title='Segmented')
```

---

ocontour

*Oriented contours*

---

## Description

Computes the oriented contour of objects.

## Usage

```
ocontour(x)
```

## Arguments

**x** An Image object or an array, containing objects. Only integer values are considered. Pixels of value 0 constitute the background. Each object is a set of pixels with the same unique integer value. Objects are assumed connected.

## Value

A list of matrices, containing the coordinates of object oriented contours.

## Author(s)

Gregoire Pau, <gpau@ebi.ac.uk>, 2008

## Examples

```
x = readImage(system.file("images", "shapes.png", package="EBImage"))
x = x[1:120,50:120]
display(x)
oc = ocontour(x)
plot(oc[[1]], type='l')
points(oc[[1]], col=2)
```

---

`otsu`*Calculate Otsu's threshold*

---

**Description**

Returns a threshold value based on Otsu's method, which can be then used to reduce the grayscale image to a binary image.

**Usage**

```
otsu(x, range = c(0, 1), levels = 256)
```

**Arguments**

<code>x</code>	A Grayscale Image object or an array.
<code>range</code>	Numeric vector of length 2 specifying the histogram range used for thresholding.
<code>levels</code>	Number of grayscale levels.

**Details**

Otsu's thresholding method [1] is useful to automatically perform clustering-based image thresholding. The algorithm assumes that the distribution of image pixel intensities follows a bi-modal histogram, and separates those pixels into two classes (e.g. foreground and background). The optimal threshold value is determined by minimizing the combined intra-class variance.

The threshold value is calculated for each image frame separately resulting in a output vector of length equal to the total number of frames in the image.

The default number of `levels` corresponds to the number of gray levels of an 8bit image. It is recommended to adjust this value according to the bit depth of the processed data, i.e. set `levels` to  $2^{16} = 65536$  when working with 16bit images.

**Value**

A vector of length equal to the total number of frames in `x`. Each vector element contains the Otsu's threshold value calculated for the corresponding image frame.

**Author(s)**

Philip A. Marais <philipmarais@gmail.com>, Andrzej Oles <andrzej.oles@embl.de>, 2014

**References**

[1] Nobuyuki Otsu, "A threshold selection method from gray-level histograms". IEEE Trans. Sys., Man., Cyber. 9 (1): 62-66. doi:10.1109/TSMC.1979.4310076 (1979)

**See Also**

[thresh](#)



### Examples

```
x = readImage(system.file("images", "sample.png", package="EBImage"))
display(x)

## threshold using Otsu's method
y = x > otsu(x)
display(y)
```

---

paintObjects	<i>Marks objects in images</i>
--------------	--------------------------------

---

### Description

This function marks objects in images.

### Usage

```
paintObjects(x, tgt, opac=c(1, 1), col=c('red', NA), thick=FALSE, closed=FALSE)
```

### Arguments

x	An Image object in Grayscale color mode or an array containing object masks. Object masks are sets of pixels with the same unique integer value.
tgt	An Image object or an array, containing the intensity values of the objects.
opac	A numeric vector of two opacity values for drawing object boundaries and object bodies. Opacity ranges from 0 to 1, with 0 being fully transparent and 1 fully opaque.
col	A character vector of two R colors for drawing object boundaries and object bodies. By default, object boundaries are painted in red while object bodies are not painted.
thick	A logical indicating whether to use thick boundary contours. Default is FALSE.
closed	A logical indicating whether object contours should be closed along image edges or remain open.

### Value

An Image object or an array, containing the painted version of tgt.

### Author(s)

Oleg Sklyar, <osklyar@ebi.ac.uk>, 2006-2007 Andrzej Oles, <andrzej.oles@embl.de>, 2015

### See Also

[bwlabel](#), [watershed](#), [computeFeatures](#), [colorLabels](#)

## Examples

```

## load images
nuc = readImage(system.file('images', 'nuclei.tif', package='EBImage'))
cel = readImage(system.file('images', 'cells.tif', package='EBImage'))
img = rgbImage(green=cel, blue=nuc)
display(img, title='Cells')

## segment nuclei
nmask = thresh(nuc, 10, 10, 0.05)
nmask = opening(nmask, makeBrush(5, shape='disc'))
nmask = fillHull(nmask)
nmask = bwlabel(nmask)
display(normalize(nmask), title='Cell nuclei mask')

## segment cells, using propagate and nuclei as 'seeds'
ctmask = opening(cel>0.1, makeBrush(5, shape='disc'))
cmask = propagate(cel, nmask, ctmask)
display(normalize(cmask), title='Cell mask')

## using paintObjects to highlight objects
res = paintObjects(cmask, img, col='#ff00ff')
res = paintObjects(nmask, res, col='#ffff00')
display(res, title='Segmented cells')

```

---

propagate

*Voronoi-based segmentation on image manifolds*


---

## Description

Find boundaries between adjacent regions in an image, where seeds have been already identified in the individual regions to be segmented. The method finds the Voronoi region of each seed on a manifold with a metric controlled by local image properties. The method is motivated by the problem of finding the borders of cells in microscopy images, given a labelling of the nuclei in the images.

Algorithm and implementation are from Jones et al. [1].

## Usage

```
propagate(x, seeds, mask=NULL, lambda=1e-4)
```

## Arguments

x	An Image object or an array, containing the image to segment.
seeds	An Image object or an array, containing the seeding objects of the already identified regions.
mask	An optional Image object or an array, containing the binary image mask of the regions that can be segmented. If missing, the whole image is segmented.

lambda            A numeric value. The regularization parameter used in the metric, determining the trade-off between the Euclidean distance in the image plane and the contribution of the gradient of  $x$ . See details.

### Details

The method operates by computing a discretized approximation of the Voronoi regions for given seed points on a Riemann manifold with a metric controlled by local image features.

Under this metric, the infinitesimal distance  $d$  between points  $v$  and  $v+dv$  is defined by:

$$d^2 = ( (t(dv)*g)^2 + lambda*t(dv)*dv ) / (lambda + 1)$$

, where  $g$  is the gradient of image  $x$  at point  $v$ .

lambda controls the weight of the Euclidean distance term. When lambda tends to infinity,  $d$  tends to the Euclidean distance. When lambda tends to 0,  $d$  tends to the intensity gradient of the image.

The gradient is computed on a neighborhood of 3x3 pixels.

Segmentation of the Voronoi regions in the vicinity of flat areas (having a null gradient) with small values of lambda can suffer from artifacts coming from the metric approximation.

### Value

An Image object or an array, containing the labelled objects.

### License

The implementation is based on CellProfiler C++ source code [2, 3]. An LGPL license was granted by Thouis Jones to use this part of CellProfiler's code for the propagate function.

### Author(s)

The original CellProfiler code is from Anne Carpenter <carpenter@wi.mit.edu>, Thouis Jones <thouis@csail.mit.edu>, In Han Kang <intheek@mit.edu>. Responsible for this implementation: Greg Pau.

### References

- [1] T. Jones, A. Carpenter and P. Golland, "Voronoi-Based Segmentation of Cells on Image Manifolds", CVBIA05 (535-543), 2005
- [2] A. Carpenter, T.R. Jones, M.R. Lamprecht, C. Clarke, I.H. Kang, O. Friman, D. Guertin, J.H. Chang, R.A. Lindquist, J. Moffat, P. Golland and D.M. Sabatini, "CellProfiler: image analysis software for identifying and quantifying cell phenotypes", Genome Biology 2006, 7:R100
- [3] CellProfiler: <http://www.cellprofiler.org>

### See Also

[bwlabel](#), [watershed](#)

**Examples**

```

## a paraboloid mountain in a plane
n = 400
x = (n/4)^2 - matrix(
(rep(1:n, times=n) - n/2)^2 + (rep(1:n, each=n) - n/2)^2,
nrow=n, ncol=n)
x = normalize(x)

## 4 seeds
seeds = array(0, dim=c(n,n))
seeds[51:55, 301:305] = 1
seeds[301:305, 101:105] = 2
seeds[201:205, 141:145] = 3
seeds[331:335, 351:355] = 4

lambda = 10^seq(-8, -1, by=1)
segmented = Image(dim=c(dim(x), length(lambda)))

for(i in seq_along(lambda)) {
  prop = propagate(x, seeds, lambda=lambda[i])
  prop = prop/max(prop)
  segmented[, ,i] = prop
}

display(x, title='Image')
display(seeds/max(seeds), title='Seeds')
display(segmented, title="Voronoi regions", all=TRUE)

```

---

 resize

*Spatial linear transformations*


---

**Description**

The following functions perform all spatial linear transforms: reflection, rotation, translation, resizing, and general affine transform.

**Usage**

```

flip(x)
flop(x)
rotate(x, angle, filter = "bilinear", output.dim, ...)
translate(x, v, filter = "none", ...)
resize(x, w, h, filter = "bilinear", output.dim = c(w, h), output.origin = c(0, 0), ...)

affine(x, m, filter = c("bilinear", "none"), output.dim, bg.col = "black")

```

**Arguments**

<code>x</code>	An Image object or an array.
<code>angle</code>	A numeric specifying the image rotation angle in degrees.
<code>v</code>	A vector of 2 numbers denoting the translation vector in pixels.
<code>w, h</code>	Width and height of the resized image. One of these arguments can be missing to enable proportional resizing.
<code>filter</code>	A character string indicating the interpolating sampling filter. Valid values are 'none' or 'bilinear'. See Details.
<code>output.dim</code>	A vector of 2 numbers indicating the dimension of the output image. For <code>affine</code> and <code>translate</code> the default is <code>dim(x)</code> , for <code>resize</code> it equals <code>c(w, h)</code> , and for <code>rotate</code> it defaults to the bounding box size of the rotated image.
<code>output.origin</code>	A vector of 2 numbers indicating the output coordinates of the origin in pixels. Default is <code>c(0, 0)</code> .
<code>m</code>	A 3x2 matrix describing the affine transformation. See Details.
<code>bg.col</code>	Color used to fill the background pixels. The default is "black".
<code>...</code>	Arguments to be passed to the affine function, such as <code>output.dim</code> or <code>bg.col</code>

**Details**

`flip` mirrors `x` across the central horizontal (x-)axis.

`flop` mirrors `x` across the central vertical (y-)axis.

`rotate` rotates the image clockwise by the specified angle around the origin. The rotation origin defaults to the center of the input image and can be changed by modifying the argument `output.origin`.

`resize` resizes the image `x` to desired dimensions. Resizing center is changed by modifying the argument `output.origin`. Zooming, without changing the output dimension, is achieved by setting the arguments `w` and `h` to values different from `output.dim`.

`affine` returns the affine transformation of `x`, where pixels coordinates, denoted by the matrix `px`, are transformed to `cbind(px, 1)%*%m`.

All spatial transformations except `flip` and `flop` are based on the general affine transformation. Spatial interpolation can be one of the following types: `none`, also called nearest neighbor, where the interpolated pixel value equals to the closest pixel value, or `bilinear`, where the interpolated pixel value is computed by bilinear approximation of the 4 neighboring pixels. The `bilinear` filter gives smoother results.

**Value**

An Image object or an array, containing the transformed version of `x`.

**Author(s)**

Gregoire Pau, 2012

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

```
x <- readImage(system.file("images", "sample.png", package="EBImage"))
display(x)

display( flip(x) )
display( flop(x) )
display( resize(x, 128) )
display( rotate(x, 30) )
display( translate(x, c(120, -20)) )

m <- matrix(c(0.6, 0.2, 0, -0.2, 0.3, 300), nrow=3)
display( affine(x, m) )
```

---

**rmObjects***Object removal and re-indexation*

---

**Description**

The rmObjects functions deletes objects from an image by setting their pixel intensity values to 0. reenumerate re-enumerates all objects in an image from 0 (background) to the actual number of objects.

**Usage**

```
rmObjects(x, index, reenumerate = TRUE)

reenumerate(x)
```

**Arguments**

x	An Image object in Grayscale color mode or an array containing object masks. Object masks are sets of pixels with the same unique integer value.
index	A numeric vector (or a list of vectors if x contains multiple frames) containing the indexes of objects to remove in the frame.
reenumerate	Logical, should all the objects in the image be re-indexed afterwards (default).

**Value**

An Image object or an array, containing the new objects.

**Author(s)**

Oleg Sklyar, <osklyar@ebi.ac.uk>, 2006-2007

**See Also**

[bwlabel](#), [watershed](#)

**Examples**

```
## make objects
x = readImage(system.file('images', 'shapes.png', package='EBImage'))
x = x[110:512,1:130]
y = bwlabel(x)

## number of objects found
max(y)

display(normalize(y), title='Objects')

## remove every second letter
objects = list(
  seq.int(from = 2, to = max(y), by = 2),
  seq.int(from = 1, to = max(y), by = 2)
)
z = rmObjects(combine(y, y), objects)

display(normalize(z), title='Object removal')

## the number of objects left in each image
apply(z, 3, max)

## perform object removal without re-enumerating
z = rmObjects(y, objects, reenumerate = FALSE)

## labels of objects left
unique(as.vector(z))[-1L]

## re-index objects
z = reenumerate(z)
unique(as.vector(z))[-1L]
```

---

stackObjects

*Places detected objects into an image stack*

---

**Description**

Places detected objects into an image stack.

**Usage**

```
stackObjects(x, ref, combine=TRUE, bg.col='black', ext)
```

**Arguments**

x	An Image object or an array containing object masks. Object masks are sets of pixels with the same unique integer value.
ref	An Image object or an array, containing the intensity values of the objects.
combine	If x contains multiple images, specifies if the resulting list of image stacks with individual objects should be combined using <code>combine</code> into a single image stack.
bg.col	Background pixel color.
ext	A numeric controlling the size of the output image. If missing, <code>ext</code> is estimated from data. See details.

**Details**

`stackObjects` creates a set of  $n$  images of size  $(2*ext+1, 2*ext+1)$ , where  $n$  is the number of objects in `x`, and places each object of `x` in this set.

If not specified, `ext` is estimated using the 98% quantile of `m.majoraxis/2`, where `m.majoraxis` is the semi-major axis descriptor extracted from `computeFeatures.moment`, taken over all the objects of the image `x`.

**Value**

An Image object containing the stacked objects contained in `x`. If `x` contains multiple images and if `combine` is `TRUE`, `stackObjects` returns a list of Image objects.

**Author(s)**

Oleg Sklyar, <osklyar@ebi.ac.uk>, 2006-2007

**See Also**

[combine](#), [tile](#), [computeFeatures.moment](#)

**Examples**

```
## simple example
x = readImage(system.file('images', 'shapes.png', package='EBImage'))
x = x[110:512,1:130]
y = bwlabel(x)
display(normalize(y), title='Objects')
z = stackObjects(y, normalize(y))
display(z, title='Stacked objects')

## load images
nuc = readImage(system.file('images', 'nuclei.tif', package='EBImage'))
cel = readImage(system.file('images', 'cells.tif', package='EBImage'))
img = rgbImage(green=cel, blue=nuc)
display(img, title='Cells')

## segment nuclei
nmask = thresh(nuc, 10, 10, 0.05)
```



```

nmask = opening(nmask, makeBrush(5, shape='disc'))
nmask = fillHull(bwlabel(nmask))

## segment cells, using propagate and nuclei as 'seeds'
ctmask = opening(cel>0.1, makeBrush(5, shape='disc'))
cmask = propagate(cel, nmask, ctmask)

## using paintObjects to highlight objects
res = paintObjects(cmask, img, col='#ff00ff')
res = paintObjects(nmask, res, col='#ffff00')
display(res, title='Segmented cells')

## stacked cells
st = stackObjects(cmask, img)
display(st, title='Stacked objects')

```

---

 thresh

*Adaptive thresholding*


---

### Description

Thresholds an image using a moving rectangular window.

### Usage

```
thresh(x, w=5, h=5, offset=0.01)
```

### Arguments

x	An Image object or an array.
w, h	Width and height of the moving rectangular window.
offset	Thresholding offset from the averaged value.

### Details

This function returns the binary image resulting from the comparison between an image and its filtered version with a rectangular window. It is equivalent of doing `{f = matrix(1, nc=2*w+1, nr=2*h+1) ; f=f/sum(f) ;` but slightly faster. The function `filter2` provides hence more flexibility than `thresh`.

### Value

An Image object or an array, containing the transformed version of x.

### Author(s)

Oleg Sklyar, <osklyar@ebi.ac.uk>, 2005-2007

**See Also**

filter2

**Examples**

```
x = readImage(system.file('images', 'nuclei.tif', package='EBImage'))
display(x)
y = thresh(x, 10, 10, 0.05)
display(y)
```

---

tile

*Tiling/untiling images*

---

**Description**

Given a sequence of frames, `tile` generates a single image with frames tiled. `untile` is the inverse function and divides an image into a sequence of images.

**Usage**

```
tile(x, nx=10, lwd=1, fg.col="#E4AF2B", bg.col="gray")
untile(x, nim, lwd=1)
```

**Arguments**

<code>x</code>	An Image object, an array or a list of these objects.
<code>nx</code>	The number of tiled images in a row.
<code>lwd</code>	The width of the grid lines between tiled images, can be 0.
<code>fg.col</code>	The color of the grid lines.
<code>bg.col</code>	The color of the background for extra tiles.
<code>nim</code>	A numeric vector of 2 elements for the number of images in both directions.

**Details**

After object segmentation, `tile` is a useful addition to `stackObjects` to have an overview of the segmented objects.

**Value**

An Image object or an array, containing the tiled/untiled version of `x`.

**Author(s)**

Oleg Sklyar, <osklyar@ebi.ac.uk>, 2006-2007

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

```

## make a set of blurred images
img = readImage(system.file("images", "sample-color.png", package="EBImage"))[257:768,,]
x = resize(img, 128, 128)
xt = list()
for (t in seq(0.1, 5, len=9)) xt=c(xt, list(gblur(x, s=t)))
xt = combine(xt)
display(xt, title='Blurred images')

## tile
xt = tile(xt, 3)
display(xt, title='Tiles')

## untile
xu = untile(img, c(3, 3))
display(xu, title='Blocks')

```

---

**transpose***Image Transposition*

---

**Description**

Transposes an image by swapping its first two, i.e., spatial dimensions.

**Usage**

```
transpose(x, coerce = FALSE)
```

**Arguments**

<code>x</code>	an Image object or an array.
<code>coerce</code>	controls the coercion of <code>x</code> . By default the output is of the same class as the input. If <code>coerce = TRUE</code> then <code>x</code> becomes coerced to an array.

**Details**

The transposition of an image is performed by swapping the X and Y indices of its array representation.

**Value**

An Image object or an array, containing `x` with its XY dimensions transposed. When `coerce = TRUE` the output is coerced to an array.

**Note**

transpose is particularly useful when converting between different representations of image data in which the X and Y dimensions are swapped. Typically, in such context only the actual pixel data matters. For performance reasons it is best practice to issue the function directly on an Image object with `coerce = TRUE` rather than to extract its image data first and only then perform the transposition, or to transpose the Image object and coerce it to an array afterwards.

**Author(s)**

Andrzej Oles, <andrzej.oles@embl.de>, 2012

**See Also**

[flip](#), [flop](#), [rotate](#)

**Examples**

```
x = readImage(system.file("images", "sample-color.png", package="EBImage"))
y = transpose(x)

display(x, title='Original')
display(y, title='Transposed')

## performing the transposition of an image twice should result in the original image
z = transpose(y)
identical(x, z)
```

---

watershed

*Watershed transformation and watershed based object detection*

---

**Description**

Watershed transformation and watershed based object detection.

**Usage**

```
watershed(x, tolerance=1, ext=1)
```

**Arguments**

x	An Image object or an array.
tolerance	The minimum height of the object in the units of image intensity between its highest point (seed) and the point where it contacts another object (checked for every contact pixel). If the height is smaller than the tolerance, the object will be combined with one of its neighbors, which is the highest. Tolerance should be chosen according to the range of x. Default value is 1, which is a reasonable value if x comes from <code>distmap</code> .
ext	Radius of the neighborhood in pixels for the detection of neighboring objects. Higher value smooths out small objects.

**Details**

The algorithm identifies and separates objects that stand out of the background (zero). After the water fill, the source image is flipped upside down and the resulting valleys (values with higher intensities) are filled in first until another object or background is met. The deepest valleys (pixels with highest intensity) become indexed first, starting from 1.

The function `bwlabel` is a simpler, faster alternative to segment connected objects from binary images.

**Value**

An Grayscale Image object or an array, containing the labelled version of `x`.

**Author(s)**

Oleg Sklyar, <osklyar@ebi.ac.uk>, 2007

**See Also**

[bwlabel](#), [propagate](#)

**Examples**

```
x = readImage(system.file('images', 'shapes.png', package='EBImage'))
x = x[110:512,1:130]
display(x, title='Binary')
y = distmap(x)
display(normalize(y), title='Distance map')
w = watershed(y)
display(normalize(w), title='Watershed')
```

# Index

- \*Topic **manip**
  - otsu, [32](#)
  - thresh, [41](#)
  - tile, [42](#)
  - watershed, [44](#)
- \*Topic **package**
  - EImage, [13](#)
  - EImage-defunct, [16](#)
- [, Image, ANY, ANY, ANY-method (Image), [21](#)
- [, Image-method (Image), [21](#)
  
- affine (resize), [36](#)
- as.array.Image (Image), [21](#)
- as.Image (Image), [21](#)
- as.raster.Image (Image), [21](#)
  
- blackTopHatGrayscale (morphology), [28](#)
- blackTopHatGreyScale (morphology), [28](#)
- blur, [19](#)
- blur (EImage-defunct), [16](#)
- bwlabel, [2](#), [5](#), [9](#), [18](#), [33](#), [35](#), [39](#), [45](#)
  
- channel, [3](#)
- closing (morphology), [28](#)
- closingGrayscale (morphology), [28](#)
- closingGreyScale (morphology), [28](#)
- cmoments (EImage-defunct), [16](#)
- Color (Image), [21](#)
- colorLabels, [3](#), [5](#), [33](#)
- colorMode, [4](#)
- colorMode (Image), [21](#)
- colormode (Image), [21](#)
- colorMode<- (Image), [21](#)
- combine, [6](#), [40](#)
- combine, array, array-method (combine), [6](#)
- combine, Image, Image-method (combine), [6](#)
- combine, list, missing-method (combine), [6](#)
- combine, matrix, matrix-method (combine), [6](#)
- computeFeatures, [3](#), [7](#), [33](#)
  
- computeFeatures.moment, [40](#)
- convolve, [19](#)
  
- dilate (morphology), [28](#)
- dilateGrayscale (morphology), [28](#)
- dilateGreyScale (morphology), [28](#)
- display, [10](#), [23](#), [25](#)
- distmap, [11](#)
- drawCircle, [12](#)
- drawfont (EImage-defunct), [16](#)
- drawtext (EImage-defunct), [16](#)
  
- EImage, [13](#)
- EImage-defunct, [16](#)
- edgeFeatures (EImage-defunct), [16](#)
- edgeProfile (EImage-defunct), [16](#)
- equalize, [16](#)
- erode (morphology), [28](#)
- erodeGrayscale (morphology), [28](#)
- erodeGreyScale (morphology), [28](#)
  
- fft, [19](#), [27](#)
- fillHull, [17](#)
- filter2, [18](#), [21](#)
- flip, [44](#)
- flip (resize), [36](#)
- floodFill, [19](#)
- flop, [44](#)
- flop (resize), [36](#)
  
- gblur, [20](#), [27](#)
- getFeatures (EImage-defunct), [16](#)
- getFrame (Image), [21](#)
- getFrames (Image), [21](#)
- getNumberOfFrames (Image), [21](#)
- Grayscale (Image), [21](#)
  
- haralickFeatures (EImage-defunct), [16](#)
- haralickMatrix (EImage-defunct), [16](#)
- hist, Image-method (Image), [21](#)
- hullFeatures (EImage-defunct), [16](#)

Image, 6, 21, 25  
image, Image-method (Image), 21  
Image-class (Image), 21  
imageData (Image), 21  
imageData<- (Image), 21  
io, 23  
is.Image (Image), 21  
  
localCurvature, 25  
  
makeBrush, 19, 21, 27  
makeBrush (morphology), 28  
Math2, Image-method (Image), 21  
median.Image (Image), 21  
medianFilter, 27  
moments (EImage-defunct), 16  
morphology, 28  
  
normalize, 5, 30  
normalize, array-method (normalize), 30  
normalize, Image-method (normalize), 30  
normalize, matrix-method (normalize), 30  
numberOfFrames (Image), 21  
  
ocontour, 26, 31  
opening (morphology), 28  
openingGrayscale (morphology), 28  
openingGreyScale (morphology), 28  
Ops, Image, Image-method (Image), 21  
Ops, Image, numeric-method (Image), 21  
Ops, numeric, Image-method (Image), 21  
otsu, 32  
  
paintObjects, 3, 33  
print.Image (Image), 21  
propagate, 3, 9, 15, 34, 45  
  
quantile.Image (Image), 21  
  
readImage, 23  
readImage (io), 23  
readJPEG, 25  
readPNG, 25  
readTIFF, 25  
reenumerate (rmObjects), 38  
resize, 36  
rgbImage (channel), 3  
rmObjects, 38  
rmoments (EImage-defunct), 16  
rotate, 44  
rotate (resize), 36  
  
selfcomplementaryTopHatGrayscale  
  (morphology), 28  
selfcomplementaryTopHatGreyScale  
  (morphology), 28  
show, Image-method (Image), 21  
smoments (EImage-defunct), 16  
stackObjects, 39, 43  
standardExpandRef (computeFeatures), 7  
  
thresh, 32, 41  
tile, 40, 42  
toRGB (channel), 3  
translate (resize), 36  
transpose, 38, 43  
  
untile (tile), 42  
  
watershed, 3, 33, 35, 39, 44  
whiteTopHatGrayscale (morphology), 28  
whiteTopHatGreyScale (morphology), 28  
writeImage, 23  
writeImage (io), 23  
writeJPEG, 25  
writePNG, 25  
writeTIFF, 24, 25  
  
zernikeMoments (EImage-defunct), 16