

# Package ‘imagerExtra’

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**Type** Package

**Title** Extra Image Processing Library Based on ‘imager’

**Version** 1.3.2

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**Description** Provides advanced functions for image processing based on the package ‘imager’.

**License** GPL-3

**Depends** R (>= 2.10.0), imager (>= 0.40.2)

**Imports** fftwtools, magrittr, Rcpp (>= 0.12.14)

**Suggests** testthat (>= 2.0.0), knitr, rmarkdown, tesseract

**URL** <https://github.com/ShotaOchi/imagerExtra>

**BugReports** <https://github.com/ShotaOchi/imagerExtra/issues>

**LinkingTo** Rcpp

**LazyData** true

**RoxygenNote** 6.1.1

**VignetteBuilder** knitr

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**NeedsCompilation** yes

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**Repository** CRAN

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BalanceSimplest	<i>Balance color of image by Simplest Color Balance</i>
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**Description**

Balance color of image by Simplest Color Balance

**Usage**

```
BalanceSimplest(im, sleft, sright, range = c(0, 255))
```

**Arguments**

im	a grayscale image of class cim
sleft	left saturation percentage. sleft can be specified by numeric or string, e.g. 1 and "1%". note that sleft is a percentile.
sright	right saturation percentage. sright can be specified by numeric or string. note that sright is a percentile.
range	this function assumes that the range of pixel values of of input image is [0,255] by default. you may prefer [0,1].

**Value**

a grayscale image of class cim

**Author(s)**

Shota Ochi

**References**

Nicolas Limare, Jose-Luis Lisani, Jean-Michel Morel, Ana Belen Petro, and Catalina Sbert, Simplest Color Balance, Image Processing On Line, 1 (2011), pp. 297-315. <https://doi.org/10.5201/ipol.2011.11mps-scb>

**Examples**

```
dev.new()
par(mfcol = c(1,2))
boats_g <- grayscale(boats)
plot(boats_g, main = "Original")
BalanceSimplest(boats_g, 1, 1) %>% plot(., main = "Simplest Color Balance")
```

---

DCT

*Two Dimensional Discrete Cosine Transformation and Inverse Cosine Transformation*

---

**Description**

DCT2D computes two dimensional discrete cosine transformation. IDCT2D computes two dimensional inverse discrete cosine transformation.

**Usage**

```
DCT2D(imormat, returnmat = FALSE)
```

```
IDCT2D(imormat, returnmat = FALSE)
```

**Arguments**

`imormat` a grayscale image of class `cimg` or a numeric matrix

`returnmat` if `returnmat` is `TRUE`, returns numeric matrix. if `FALSE`, returns a grayscale image of class `cimg`.

**Value**

a grayscale image of class `cimg` or a numeric matrix

**Author(s)**

Shota Ochi

## References

Makhoul, J. (1980). A fast cosine transform in one and two dimensions. IEEE Transactions on Acoustics, Speech, and Signal Processing, 28 (1): 27-34.

## Examples

```
g <- grayscale(boats)
layout(matrix(1:2, 1, 2))
plot(g, main = "Original")
gg <- DCT2D(g) %>% IDCT2D() %>% plot(main = "Transformed")
mean((g - gg)^2)
```

---

DenoiseDCT

*denoise image by DCT denoising*

---

## Description

denoise image by DCT denoising

## Usage

```
DenoiseDCT(im, sdn, flag_dct16x16 = FALSE)
```

## Arguments

im	a grayscale image of class cimg
sdn	standard deviation of Gaussian white noise
flag_dct16x16	flag_dct16x16 determines the size of patches. if TRUE, the size of patches is 16x16. if FALSE, the size of patches is 8x8.

## Value

a grayscale image of class cimg

## Author(s)

Shota Ochi

## References

Guoshen Yu, and Guillermo Sapiro, DCT Image Denoising: a Simple and Effective Image Denoising Algorithm, Image Processing On Line, 1 (2011), pp. 292-296. <https://doi.org/10.5201/ipol.2011.js-dct>

**Examples**

```
dev.new()
par(mfcol = c(1,2))
boats_g <- grayscale(boats)
boats_noisy <- imnoise(dim = dim(boats_g), sd = 0.05) + boats_g
plot(boats_noisy, main = "Noisy Boats")
DenoiseDCT(boats_g, 0.05) %>% plot(., main = "Denoised Boats")
```

dogs

*Photograph of a dog from GAHAG***Description**

This photograph was downloaded from <http://gahag.net/img/201603/03s/gahag-0062116383-1.jpg>. Its size was reduced by half to speed up loading and save space.

**Usage**

dogs

**Format**

an image of class cimg

**Source**

<http://gahag.net/img/201603/03s/gahag-0062116383-1.jpg>

EqualizeADP

*Adaptive Double Plateaus Histogram Equalization***Description**

compute the paramters, `t_down` and `t_up`, and then apply double plateaus histogram equalization.

**Usage**

```
EqualizeADP(im, n = 5, N = 1000, range = c(0, 255),
  returnparam = FALSE)
```

**Arguments**

<code>im</code>	a grayscale image of class cimg
<code>n</code>	window size to determine local maximum
<code>N</code>	the number of subintervals of histogram
<code>range</code>	range of the pixel values of image. this function assumes that the range of pixel values of of an input image is [0,255] by default. you may prefer [0,1].
<code>returnparam</code>	if <code>returnparam</code> is TRUE, returns the computed parameters: <code>t_down</code> and <code>t_up</code> .

**Value**

a grayscale image of class `cimg` or a numeric vector

**Author(s)**

Shota Ochi

**References**

Kun Liang, Yong Ma, Yue Xie, Bo Zhou ,Rui Wang (2012). A new adaptive contrast enhancement algorithm for infrared images based on double plateaus histogram equalization. *Infrared Phys. Technol.* 55, 309-315.

**Examples**

```
g <- grayscale(dogs)
layout(matrix(1:2, 1, 2))
plot(g, main = "Original")
EqualizeADP(g) %>% plot(main = "Contrast Enhanced")
```

---

EqualizeDP

*Double Plateaus Histogram Equalization*

---

**Description**

enhance contrast of image by double plateaus histogram equalization.

**Usage**

```
EqualizeDP(im, t_down, t_up, N = 1000, range = c(0, 255))
```

**Arguments**

<code>im</code>	a grayscale image of class <code>cimg</code>
<code>t_down</code>	lower threshold
<code>t_up</code>	upper threshold
<code>N</code>	the number of subintervals of histogram
<code>range</code>	range of the pixel values of image. this function assumes that the range of pixel values of of an input image is [0,255] by default. you may prefer [0,1].

**Value**

a grayscale image of class `cimg`

**Author(s)**

Shota Ochi

## References

Kun Liang, Yong Ma, Yue Xie, Bo Zhou ,Rui Wang (2012). A new adaptive contrast enhancement algorithm for infrared images based on double plateaus histogram equalization. *Infrared Phys. Technol.* 55, 309-315.

## Examples

```
g <- grayscale(dogs)
layout(matrix(1:2, 1, 2))
plot(g, main = "Original")
EqualizeDP(g, 20, 186) %>% plot(main = "Contrast Enhanced")
```

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EqualizePiecewise	<i>Piecewise Affine Histogram Equalization</i>
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## Description

enhance contrast of image by piecewise affine histogram equalization

## Usage

```
EqualizePiecewise(im, N, smax = 255, smin = 0, range = c(0, 255))
```

## Arguments

im	a grayscale image of class cimg
N	number of subintervals of partition. N controls how the input gray levels will be mapped in the output image. if N is large, Piecewise Affine Equalization and Histogram Equalization are very similar.
smax	maximum value of slopes. if smax is small, contrast enhancement is suppressed.
smin	minimum value of slopes. if smin is large, contrast enhancement is propelled, and saturations occur excessively.
range	range of the pixel values of image. this function assumes that the range of pixel values of of an input image is [0,255] by default. you may prefer [0,1]. if you change range, you should change smax. one example is this (smax = range[2] - range[1]).

## Value

a grayscale image of class cimg

## Author(s)

Shota Ochi

## References

Jose-Luis Lisani, Ana-Belen Petro, and Catalina Sbert, Color and Contrast Enhancement by Controlled Piecewise Affine Histogram Equalization, Image Processing On Line, 2 (2012), pp. 243-265. <https://doi.org/10.5201/ipol.2012.lps-pae>

## Examples

```
dev.new()
par(mfcol = c(1,2))
boats_g <- grayscale(boats)
plot(boats_g, main = "Original")
EqualizePiecewise(boats_g, 10) %>% plot(., main = "Piecewise Affine Equalization")
```

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GetHue	<i>store hue of color image</i>
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---

## Description

store hue of color image

## Usage

```
GetHue(imcol)
```

## Arguments

imcol            a color image of class cimg

## Value

a color image of class cimg

## Author(s)

Shota Ochi

## Examples

```
GetHue(boats)
```

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Grayscale	<i>compute average of RGB channels</i>
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**Description**

compute average of RGB channels

**Usage**

```
Grayscale(imcol)
```

**Arguments**

imcol            a color image of class cimg

**Value**

a grayscale image of class cimg

**Author(s)**

Shota Ochi

**Examples**

```
Grayscale(boats) %>% plot
```

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imagerExtra	<i>imagerExtra: Extra Image Processing Library Based on Imager</i>
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**Description**

imagerExtra is built on imager. imager by Simon Simon Barthelme provides an interface with CImg that is a C++ library for image processing. imager makes functions of CImg accessible from R and adds many utilities for accessing and working with image data from R. imagerExtra provides advanced functions for image processing based on imager.

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 OCR

*Optical Character Recognition with tesseract*


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**Description**

OCR and OCR\_data are wrappers for ocr and ocr\_data of tesseract package. You need to install tesseract package to use these functions.

**Usage**

```
OCR(imorpx, engine = tesseract::tesseract("eng"), HOOCR = FALSE)
```

```
OCR_data(imorpx, engine = tesseract::tesseract("eng"))
```

**Arguments**

imorpx	a grayscale image of class cimg or a pixel set
engine	a tesseract engine. See the reference manual of tesseract for detail.
HOOCR	if TRUE return results as HOOCR xml instead of plain text

**Author(s)**

Shota Ochi

**Examples**

```
hello <- DenoisedCT(papers, 0.01) %>% ThresholdAdaptive(., 0.1, range = c(0,1))
if (requireNamespace("tesseract", quietly = TRUE))
{
  OCR(hello) %>% cat
  OCR_data(hello)
}
```

---

 papers

*Photograph of a paper*


---

**Description**

This photograph was filmed by Shota Ochi.

**Usage**

```
papers
```

**Format**

an image of class cimg

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RestoreHue	<i>restore hue of color image</i>
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**Description**

restore hue of color image

**Usage**

```
RestoreHue(im, hueim)
```

**Arguments**

im	a grayscale image of class cim
hueim	a color image of class cim

**Value**

a color image of class cim

**Author(s)**

Shota Ochi

**Examples**

```
g <- Grayscale(boats)
hue <- GetHue(boats)
layout(matrix(1:2, 1, 2))
plot(g, main = "Original")
RestoreHue(g, hue) %>% plot(main="Resotred")
```

---

SegmentCV	<i>Chan-Vese segmentation</i>
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---

**Description**

iterative image segmentation with Chan-Vese model

**Usage**

```
SegmentCV(im, mu = 0.25, nu = 0, lambda1 = 1, lambda2 = 1,
  tol = 1e-04, maxiter = 500, dt = 0.5, initial, returnstep)
```

**Arguments**

im	a grayscale image of class cimg
mu	length penalty
nu	area penalty
lambda1	fit weight inside the cuve
lambda2	fit weight outside the curve
tol	convergence tolerance
maxiter	maximum number of iterations
dt	time step
initial	"interactive" or a grayscale image of class cimg. you can define initial condition as a rectangle shape interactively if initial is "interactive". If initial is a grayscale image of class cimg, pixels whose values are negative will be treated as outside of contour. pixels whose values are non-negative will be treated as inside of contour. checker board condition will be used if initial is not specified.
returnstep	a numeric vector that determines which result will be returned. 0 means initial condition, and 1 means the result after 1 iteration. final result will be returned if returnstep is not specified.

**Value**

a pixel set or a list of lists of numeric and pixel set

**Author(s)**

Shota Ochi

**References**

Pascal Getreuer (2012). Chan-Vese Segmentation. Image Processing On Line 2, 214-224.

**Examples**

```
layout(matrix(1:2, 1, 2))
g <- grayscale(dogs)
plot(g, main = "Original")
SegmentCV(g, lambda2 = 15) %>% plot(main = "Binarized")
```

---

SPE *Correct inhomogeneous background of image by solving Screened Poisson Equation*

---

### Description

Correct inhomogeneous background of image by solving Screened Poisson Equation

### Usage

```
SPE(im, lamda, s = 0.1, range = c(0, 255))
```

### Arguments

im	a grayscale image of class cimg
lamda	this function corrects inhomogeneous background while preserving image details. lamda controls the trade-off. when lamda is too large, this function acts as an edge detector.
s	saturation percentage. this function uses <a href="#">BalanceSimplest</a> . s is used as both left and sright. that's why s can not be over 50%.
range	this function assumes that the range of pixel values of of an input image is [0,255] by default. you may prefer [0,1].

### Value

a grayscale image of class cimg

### Author(s)

Shota Ochi

### References

Jean-Michel Morel, Ana-Belen Petro, and Catalina Sbert, Screened Poisson Equation for Image Contrast Enhancement, Image Processing On Line, 4 (2014), pp. 16-29. <https://doi.org/10.5201/ipol.2014.84>

### Examples

```
dev.new()
par(mfcol = c(1,2))
boats_g <- grayscale(boats)
plot(boats_g, main = "Original")
SPE(boats_g, 0.1) %>% plot(main = "Screened Poisson Equation")
```

---

ThresholdAdaptive      *Local Adaptive Thresholding*

---

## Description

Local Adaptive Thresholding

## Usage

```
ThresholdAdaptive(im, k, windowsize = 17, range = c(0, 255))
```

## Arguments

<code>im</code>	a grayscale image of class <code>cimg</code>
<code>k</code>	a numeric in the range <code>[0,1]</code> . when <code>k</code> is high, local threshold values tend to be lower. when <code>k</code> is low, local threshold value tend to be higher.
<code>windowsize</code>	<code>windowsize</code> controls the number of local neighborhood
<code>range</code>	this function assumes that the range of pixel values of of input image is <code>[0,255]</code> by default. you may prefer <code>[0,1]</code> . Note that <code>range</code> determines the max standard deviation. The max standard deviation plays an important role in this function.

## Value

a pixel set

## Author(s)

Shota Ochi

## References

Faisal Shafait, Daniel Keysers, Thomas M. Breuel, "Efficient implementation of local adaptive thresholding techniques using integral images", Proc. SPIE 6815, Document Recognition and Retrieval XV, 681510 (28 January 2008)

## Examples

```
layout(matrix(1:4, 2, 2))
plot(papers, main = "Original")
threshold(papers) %>% plot(main = "A variant of Otsu")
ThresholdAdaptive(papers, 0, range = c(0,1)) %>% plot(main = "local adaptive (k = 0)")
ThresholdAdaptive(papers, 0.2, range = c(0,1)) %>% plot(main = "local adaptive (k = 0.2)")
```

---

ThresholdFuzzy      *Fuzzy Entropy Image Segmentation*

---

### Description

automatic fuzzy thresholding based on particle swarm optimization

### Usage

```
ThresholdFuzzy(im, n = 50, maxiter = 100, omegamax = 0.9,  
  omegamin = 0.1, c1 = 2, c2 = 2, mutrate = 0.2, vmaxcoef = 0.1,  
  intervalnumber = 1000, returnvalue = FALSE)
```

### Arguments

im	a grayscale image of class cim
n	swarm size
maxiter	maximum iterative time
omegamax	maximum inertia weight
omegamin	minimum inertia weight
c1	acceleration coefficient
c2	acceleration coefficient
mutrate	rate of gaussian mutation
vmaxcoef	coefficient of maximum velocity
intervalnumber	interval number of histogram
returnvalue	if returnvalue is TRUE, returns a threshold value. if FALSE, returns a pixel set.

### Value

a pixel set or a numeric

### Author(s)

Shota Ochi

### References

Linyi Li, Deren Li (2008). Fuzzy entropy image segmentation based on particle swarm optimization. Progress in Natural Science.

### Examples

```
g <- grayscale(boats)  
layout(matrix(1:2, 1, 2))  
plot(g, main = "Original")  
ThresholdFuzzy(g) %>% plot(main = "Fuzzy Thresholding")
```

**Description**

Segments a grayscale image into several gray levels. Multilevel thresholding selection based on the artificial bee colony algorithm is used when thr is not a numeric vector. Preset parameters for fast computing is used when thr is "fast". Preset parameters for precise computing is used when thr is "precise". You can tune the parameters if thr is "manual". Also you can specify the values of thresholds by setting thr as a numeric vector.

**Usage**

```
ThresholdML(im, k, thr = "fast", sn = 30, mcn = 100, limit = 100,
  intervalnumber = 1000, returnvalue = FALSE)
```

**Arguments**

im	a grayscale image of class cimg
k	level of thresholding. k is ignored when thr is a numeric vector.
thr	thresholds, either numeric vector, or "fast", or "precise", or "manual".
sn	population size. sn is ignored except when thr is "manual".
mcn	maximum cycle number. mcn is ignored except when thr is "manual".
limit	abandonment criteria. limit is ignored except when thr is "manual".
intervalnumber	interval number of histogram. intervalnumber is ignored except when thr is "manual".
returnvalue	if returnvalue is TRUE, returns threshold values. if FALSE, returns a grayscale image of class cimg.

**Value**

a grayscale image of class cimg or a numeric vector

**Author(s)**

Shota Ochi

**References**

Ming-HuwiHorng (2011). Multilevel thresholding selection based on the artificial bee colony algorithm for image segmentation. Expert Systems with Applications.

**Examples**

```
g <- grayscale(boats)
ThresholdML(g, k = 2) %>% plot
```

---

ThresholdTriclass      *Iterative Triclass Thresholding*

---

**Description**

compute threshold value by Iterative Triclass Threshold Technique

**Usage**

```
ThresholdTriclass(im, stopval = 0.01, repeatnum, intervalnumber = 1000,  
  returnvalue = FALSE)
```

**Arguments**

im	a grayscale image of class cimg
stopval	value to determine whether stop iteration of triclass thresholding or not. Note that if repeat is set, stop is ignored.
repeatnum	number of repetition of triclass thresholding
intervalnumber	interval number of histogram
returnvalue	if returnvalue is TRUE, ThresholdTriclass returns threshold value. if FALSE, ThresholdTriclass returns pixset.

**Value**

a pixel set or a numeric

**Author(s)**

Shota Ochi

**References**

Cai HM, Yang Z, Cao XH, Xia WM, Xu XY (2014). A New Iterative Triclass Thresholding Technique in Image Segmentation. IEEE TRANSACTIONS ON IMAGE PROCESSING.

**Examples**

```
g <- grayscale(boats)  
layout(matrix(1:4, 2, 2))  
plot(boats, main = "Original")  
plot(g, main = "Grayscale")  
threshold(g) %>% plot(main = "A Variant of Otsu")  
ThresholdTriclass(g) %>% plot(main = "Triclass")
```

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