

Package: WhiteStripe (via r-universe)

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Type Package

Title White Matter Normalization for Magnetic Resonance Images

Version 2.4.3

Description Shinohara (2014) <[doi:10.1016/j.nicl.2014.08.008](https://doi.org/10.1016/j.nicl.2014.08.008)> introduced 'WhiteStripe', an intensity-based normalization of T1 and T2 images, where normal appearing white matter performs well, but requires segmentation. This method performs white matter mean and standard deviation estimates on data that has been rigidly-registered to the 'MNI' template and uses histogram-based methods.

License GPL-3

Depends R (>= 2.10), methods

Imports graphics, stats, utils, oro.nifti (>= 0.5.0), mgcv, neurobase

LazyData true

BugReports <https://github.com/muschellij2/WhiteStripe/issues>

Suggests knitr, rmarkdown

VignetteBuilder knitr

Encoding UTF-8

LazyDataCompression xz

RoxygenNote 7.3.1

Repository <https://muschellij2.r-universe.dev>

RemoteUrl <https://github.com/muschellij2/whitestripe>

RemoteRef HEAD

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download_img_data	<i>Download T1 and T2 data</i>
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Description

Download T1 and T2 data for Examples

Usage

```
download_img_data(lib.loc = NULL)
```

Arguments

lib.loc a character vector with path names of R libraries. Passed to [img_data](#)

Value

Logical indicator if the files were downloaded.

get.deriv.smooth.hist	<i>Gets nth derivative of smoothed histogram</i>
-----------------------	--

Description

This function outputs the nth derivative of a histogram smooth.

Usage

```
get.deriv.smooth.hist(x, coefs, knots, deg = 4, deriv.deg = 1)
```

Arguments

x	values from smooth_hist
coefs	Coefficients from GAM from smooth_hist
knots	Number of knots fit for GAM
deg	Degree of polynomials
deriv.deg	The degree of the derivative.

Value

Derivative of smoothed histogram

Examples

```
data(smoothed_histogram)
dy<-get.deriv.smooth.hist(xvals,
  coefs=s.hist$coefs,
  knots=s.hist$knots,
  deg=s.hist$deg,
  deriv.deg=1)
```

get.first.mode	<i>Get First Peak</i>
----------------	-----------------------

Description

This function grabs the first peak or shoulder.

Usage

```
get.first.mode(x, y, rare.prop = 1/5, verbose = TRUE, remove.tail = TRUE, ...)
```

Arguments

x	values of midpoints from hist
y	values of counts from hist
rare.prop	Proportion used to remove rare intensity tail
verbose	print diagnostic output
remove.tail	Remove rare intensity tail
...	arguments to be passed to smooth_hist

Value

Value of x that is the first peak

Examples

```

data(t1.voi.hist)
system.time({
y = t1.voi.hist$counts
x = t1.voi.hist$mids
x = x[!is.na(y)];
y = y[!is.na(y)]
# 20 used for speed of example
nawm_peak = get.first.mode(x, y, k=20)
plot(t1.voi.hist, border="red")
abline(v=nawm_peak)
})

```

get.largest.mode	<i>Grab largest peak</i>
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Description

This function grabs the largest peak of the histogram

Usage

```
get.largest.mode(x, y, verbose = TRUE, ...)
```

Arguments

x	values of midpoints from hist
y	values of counts from hist
verbose	print diagnostic output
...	arguments to be passed to smooth_hist

Value

Value of x that is the largest peak

Examples

```

data(t2.voi.hist)
system.time({
y = t2.voi.hist$counts
x = t2.voi.hist$mids
x = x[!is.na(y)];
y = y[!is.na(y)]
# 30 used for speed of example
nawm_peak = get.largest.mode(x, y, k=30)
plot(t2.voi.hist, border="red")
abline(v=nawm_peak)
})

```

get.last.mode	<i>Get Last Peak</i>
---------------	----------------------

Description

This function grabs the last peak or shoulder.

Usage

```
get.last.mode(x, y, rare.prop = 1/5, verbose = TRUE, remove.tail = TRUE, ...)
```

Arguments

x	values of midpoints from hist
y	values of counts from hist
rare.prop	Proportion used to remove rare intensity tail
verbose	print diagnostic output
remove.tail	Remove rare intensity tail
...	arguments to be passed to smooth_hist

Value

Value of x that is the last peak

Examples

```
data(t1.voi.hist)
system.time({
  y = t1.voi.hist$counts
  x = t1.voi.hist$mids
  x = x[!is.na(y)];
  y = y[!is.na(y)]
  # 20 used for speed of example
  nawm_peak = get.last.mode(x, y, k=20)
  plot(t1.voi.hist, border="red")
  abline(v=nawm_peak)
})
```

make_img_voi	<i>Make Image VOI</i>
--------------	-----------------------

Description

Creates a VOI of Image for the specified slices

Usage

```
make_img_voi(img, slices = 80:120, na.rm = TRUE, ...)
```

Arguments

img	Image (T1 usually or T2). Array or object of class nifti
slices	Slices to take for the image voi
na.rm	Remove NAs from mean. This is for double checking
...	Arguments passed from other methods (not used)

Value

VOI of image.

s.hist	<i>Smoothed histogram of image</i>
--------	------------------------------------

Description

Smoothed histogram of image

Usage

```
s.hist
```

Format

A GAM from mgcv for x and y from histograms

Examples

```
## Not run:
data(t2.voi.hist)
y = t2.voi.hist$counts
x = t2.voi.hist$mids
x = x[!is.na(y)];
y = y[!is.na(y)]
# 70 used for speed of example
s.hist = smooth_hist(x, y, k=70)

## End(Not run)
```

smooth_hist

*Histogram smoothing for whitestripe***Description**

Uses a generalized additive model (GAM) to smooth a histogram for whitestripe

Usage

```
smooth_hist(
  x,
  y,
  deg = 4,
  k = floor(min(250, length(x)/2)),
  method = "REML",
  ...
)
```

Arguments

x	values of midpoints from hist
y	values of counts from hist
deg	degree of polynomials used
k	Number of knots
method	Method for smoothing for GAM
...	Arguments passed to gam

Value

List of objects: x and y coordinates of histogram, coefficients from GAM, fitted values from GAM, the GAM model, the knots fitted, and degrees of polynomials

See Also

[gam](#)

Examples

```

data(t2.voi.hist)
y = t2.voi.hist$counts
x = t2.voi.hist$mids
x = x[!is.na(y)];
y = y[!is.na(y)]
# 30 used for speed of example
s.hist = smooth_hist(x, y, k=30)
plot(t2.voi.hist, border="red")
lines(s.hist)

```

t1.voi.hist

Histogram of VOI of T1 template image

Description

Histogram of VOI of T1 template image

Usage

```
t1.voi.hist
```

Format

A volume of interest histogram from a T1 image for smoothing

Examples

```

lib.loc = tempdir()
if (download_img_data(lib.loc = lib.loc)){
  t1 = oro.nifti::readNIFTI(system.file("T1Strip.nii.gz", package="WhiteStripe",
                                       lib.loc = lib.loc))

  t1.voi = make_img_voi(t1)
  any(is.na(t1.voi))
  # FALSE
  t1.voi.hist = hist(t1.voi,
                    breaks=2000,
                    plot=FALSE)
}

```

t2.voi.hist	<i>Histogram of VOI of T2 template image</i>
-------------	--

Description

Histogram of VOI of T2 template image

Usage

```
t2.voi.hist
```

Format

A histogram volume of interest from a T2 image for smoothing

Examples

```
## Not run:
lib.loc = tempdir()
if (download_img_data(lib.loc = lib.loc)){
t2 = readNIfTI(system.file("T2Strip.nii.gz", package="WhiteStripe",
lib.loc = lib.loc))
t2.voi = make_img_voi(t2)
any(is.na(t2.voi))
# FALSE
t2.voi.hist = hist(t2.voi,
breaks=2000,
plot=FALSE)
#save(t2.voi.hist, file="data/t2.voi.hist.rda", compress = TRUE,
# compression_level=9)
}

## End(Not run)
```

whitestripe	<i>Performs White Stripe of T1 or T2 Images</i>
-------------	---

Description

Returns the mean/sd of the whitestripe and indices for them on the image

Usage

```
whitestripe(
  img,
  type = c("T1", "T2", "FA", "MD", "first", "last", "largest"),
  breaks = 2000,
  whitestripe.width = 0.05,
  whitestripe.width.l = whitestripe.width,
  whitestripe.width.u = whitestripe.width,
  arr.ind = FALSE,
  verbose = TRUE,
  stripped = FALSE,
  slices = NULL,
  ...
)
```

Arguments

<code>img</code>	Image (T1, T2, FA, or MD). Array or object of class <code>nifti</code>
<code>type</code>	T1, T2, FA, or MD image <code>whitestripe</code>
<code>breaks</code>	Number of breaks passed to <code>hist</code>
<code>whitestripe.width</code>	Radius of the white stripe
<code>whitestripe.width.l</code>	Lower Radius of the white stripe
<code>whitestripe.width.u</code>	Upper Radius of the white stripe
<code>arr.ind</code>	Whether indices should be array notation or not, passed to <code>which</code>
<code>verbose</code>	Print diagnostic information
<code>stripped</code>	Applying to skull-stripped image. NOTE: This does NOT do a subset of slices, as <code>make_img_voi</code> .
<code>slices</code>	slices to use for <code>make_img_voi</code> if only a subset to estimate the distribution.
<code>...</code>	Arguments to be passed to <code>get.last.mode</code>

Details

This function takes in an image and computes a window of the distribution called the white stripe. If you wish to pass in values you have subset, such as single from a skull-stripped image, you can pass in `img` and set the class to `img_voi` (`class(img) = "img_voi"`) and this will not rerun `make_img_voi`.

Value

List of indices of `whitestripe`, last mode of histogram, array/nifti of 0/1 corresponding to the mask, mean of `whitestripe`, standard deviation of `whitestripe`

Examples

```

## Not run:
library(WhiteStripe)
lib.loc = tempdir()
if (WhiteStripe::download_img_data(lib.loc = lib.loc)){
  library(oro.nifti)
  set.seed(1)
  t1 = readNIfTI(system.file("T1Strip.nii.gz", package="WhiteStripe",
  lib.loc = lib.loc))
  t1.ind = whitestripe(t1, "T1")
  set.seed(2)
  t1_2 = readNIfTI(system.file("T1Strip.nii.gz", package="WhiteStripe",
  lib.loc = lib.loc))
  t1_2.ind = whitestripe(t1_2, "T1")
  t1.mask = whitestripe_ind_to_mask(t1, t1.ind$whitestripe.ind)
  t1.mask[t1.mask == 0] = NA
  orthographic(t1, t1.mask, col.y="red")
  t2 = readNIfTI(system.file("T2Strip.nii.gz", package="WhiteStripe",
  lib.loc = lib.loc))
  t2.ind = whitestripe(t2, "T2")
  t2.mask = whitestripe_ind_to_mask(t2, t2.ind$whitestripe.ind)
  t2.mask[t2.mask == 0] = NA
  orthographic(t2, t2.mask, col.y="red")
}

## End(Not run)

```

whitestripe_hybrid *Hybrid WhiteStripe*

Description

Uses t1 and t2 WhiteStripe to get an intersection of the two masks for a hybrid approach

Usage

```
whitestripe_hybrid(t1, t2, ...)
```

Arguments

t1	T1 image, array or class nifti
t2	T2 image, array or class nifti
...	arguments passed to whitestripe

Value

List of indices of overlap mask, and overlap of class array or nifti

See Also

whitestripe

Examples

```
## Not run:
lib.loc = tempdir()
if (download_img_data(lib.loc = lib.loc)){
  t1 = readNIfTI(system.file("T1Strip.nii.gz", package="WhiteStripe",
  lib.loc = lib.loc))
  t2 = readNIfTI(system.file("T2Strip.nii.gz", package="WhiteStripe",
  lib.loc = lib.loc))
  ind = whitestripe_hybrid(t1, t2)
}

## End(Not run)
```

whitestripe_ind_to_mask

WhiteStripe Indices to Mask

Description

Uses WhiteStripe indices to create image mask

Usage

```
whitestripe_ind_to_mask(img, indices, writeimg = FALSE, ...)
```

Arguments

img	Array or class nifti that is corresponds to dimensions of the images the indices were generated from
indices	indices from whitestripe
writeimg	logical to write image or not
...	arguments to passed to writeNIfTI for writing image

Value

Class of array or nifti depending on img input

See Also

whitestripe, whitestripe_hybrid

Examples

```

lib.loc = tempdir()
if (download_img_data(lib.loc = lib.loc)){
  t1 = oro.nifti::readNIFTI(system.file("T1Strip.nii.gz",
                                     package="WhiteStripe",
                                     lib.loc = lib.loc))
  t2 = oro.nifti::readNIFTI(system.file("T2Strip.nii.gz",
                                     package="WhiteStripe",
                                     lib.loc = lib.loc))

  ind = whitestripe_hybrid(t1, t2)
  mask = whitestripe_ind_to_mask(t1, ind$whitestripe.ind)
  oro.nifti::orthographic(mask)
}

```

whitestripe_norm	<i>Normalize Image using white stripe</i>
------------------	---

Description

Taking the indices from white stripe to normalize the intensity values of the brain

Usage

```
whitestripe_norm(img, indices, ...)
```

Arguments

img	Array or object of class nifti
indices	Indices of white stripe from whitestripe . Can also be a mask (indices where mask > 0 are used.)
...	arguments to be passed to mean and sd

Value

Object of same class as img, but normalized

ws_img_data	<i>Return Filenames of T1 and T2 data</i>
-------------	---

Description

Return filenames T1 and T2 data for example and vignettes

Usage

```
ws_img_data(lib.loc = NULL, warn = TRUE)
```

Arguments

lib.loc	a character vector with path names of R libraries. Passed to system.file
warn	Should a warning be printed if the images were not there

Value

Vector of filenames

xvals	<i>Midpoints from VOI histogram</i>
-------	-------------------------------------

Description

Midpoints from VOI histogram

Usage

```
xvals
```

Format

x values from histogram for VOI

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