# Package 'WhiteStripe'

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Type Package

Version 2.4.3

```
Description Shinohara (2014) <doi: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
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```

Title White Matter Normalization for Magnetic Resonance Images

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download\_img\_data

Download T1 and T2 data

# 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.

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```
get.deriv.smooth.hist Gets nth derivative of smoothed histogram
```

### **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)</pre>
```

get.first.mode

Get First Peak

### **Description**

This function grabs the first peak or shoulder.

# Usage

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

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# **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

Grab largest peak

#### **Description**

This function grabs the largest peak of the histogram

# Usage

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

### **Arguments**

```
    values of midpoints from hist
    values of counts from hist
    verbose
    print diagnostic output
    arguments to be passed to smooth_hist
```

get.last.mode 5

### 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

Get Last Peak

# 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

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### **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

Make Image VOI

# 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 7

s.hist

Smoothed histogram of image

# **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",
   ...
)
```

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#### **Arguments**

Χ	values of midpoints from hist
у	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

t2.voi.hist

#### **Examples**

t2.voi.hist

Histogram of VOI of T2 template image

## **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)
```

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whitestripe

Performs White Stripe of T1 or T2 Images

### **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**

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

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#### **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

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### Value

Class of array or nifti depending on img input

#### See Also

```
whitestripe, whitestripe_hybrid
```

### **Examples**

whitestripe\_norm

Normalize Image using white stripe

# **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

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 $ws\_img\_data$ 

Return Filenames of T1 and T2 data

# 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

Midpoints from VOI histogram

# Description

Midpoints from VOI histogram

### Usage

xvals

### **Format**

x values from histogram for VOI

# **Index**

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