

# Package ‘neurobase’

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**Type** Package

**Title** 'Neuroconductor' Base Package with Helper Functions for 'nifti' Objects

**Version** 1.32.4

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**Description** Base package for 'Neuroconductor', which includes many helper functions that interact with objects of class 'nifti', implemented by package 'oro.nifti', for reading/writing and also other manipulation functions.

**Imports** methods, abind, matrixStats, R.utils, graphics, grDevices, stats, RNifti

**Depends** oro.nifti (>= 0.11.3), R (>= 3.2.0)

**License** GPL-2

**Suggests** testthat, ggplot2, knitr, rmarkdown, dplyr, reshape2, httr, covr, brainR

**VignetteBuilder** knitr

**BugReports** <https://github.com/muschelli.j2/neurobase/issues>

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applyEmptyImageDimensions-methods  
*Apply Subsetting from Empty Image Dimensions*

**Description**

Simple wrapper for subsetting an image with indices, dropping empty dimensions.

**Usage**

```

applyEmptyImageDimensions(img, inds, reorient = FALSE, ...)

## S4 method for signature 'nifti'
applyEmptyImageDimensions(img, inds, reorient = FALSE, ...)

## S4 method for signature 'character'
applyEmptyImageDimensions(img, inds, reorient = FALSE, ...)

## S4 method for signature 'factor'
applyEmptyImageDimensions(img, inds, reorient = FALSE, ...)

## S4 method for signature 'list'

```

```

applyEmptyImageDimensions(img, inds, reorient = FALSE, ...)

## S4 method for signature 'array'
applyEmptyImageDimensions(img, inds, reorient = FALSE, ...)

## S4 method for signature 'anlz'
applyEmptyImageDimensions(img, inds, reorient = FALSE, ...)

## S4 method for signature 'ANY'
applyEmptyImageDimensions(img, inds, reorient = FALSE, ...)

apply_empty_dim(img, ...)

```

**Arguments**

<code>img</code>	image, nifti object, or array
<code>inds</code>	indices of subset from <a href="#">getEmptyImageDimensions</a> or <a href="#">dropEmptyImageDimensions</a> .
<code>reorient</code>	Should image be reoriented if a filename?
<code>...</code>	not used

**Value**

Object of class `nifti` or array if `nifti` is not supplied

**Note**

`apply_empty_dim` is a shorthand for `applyEmptyImageDimensions` with all the same arguments.

**See Also**

[getEmptyImageDimensions](#), [dropEmptyImageDimensions](#)

**Examples**

```

set.seed(5)
dims = rep(10, 3)
arr = array(rnorm(prod(dims)), dim = dims)
arr[, ,10] = 0
nim = oro.nifti::nifti(arr)
inds = getEmptyImageDimensions(nim)
inds_arr = getEmptyImageDimensions(arr)
testthat::expect_equal(inds, inds_arr)

out = applyEmptyImageDimensions(nim, inds = inds)
out_arr = applyEmptyImageDimensions(arr, inds = inds)
testthat::expect_equal(out_arr, array(out, dim = dim(out)))
out = apply_empty_dim(nim, inds = inds)

set.seed(5)
dims = rep(10, 3)

```

```

arr = array(rnorm(prod(dims)), dim = dims)
arr[, , 10] = 0
nim = oro.nifti::nifti(arr)
inds = getEmptyImageDimensions(nim)
rnifti = RNifti::asNifti(nim)
ting = tempimg(nim)
limg = list(factor(timg), factor(timg))
apply_empty_dim(nim, inds = inds)
func = function(...) applyEmptyImageDimensions(..., inds = inds)
func(arr)
func(nim)
func(rnifti)
func(timg)
func(limg)

```

---

boxplot.nifti

*Boxplot of Values in an Image*


---

## Description

Computes the boxplot of values of an image with the option for a mask.

## Usage

```
## S3 method for class 'nifti'
boxplot(x, ..., mask)
```

```
## S3 method for class 'anlz'
boxplot(x, ..., mask)
```

## Arguments

x	Object of class nifti
...	Arguments passed to <a href="#">boxplot.default</a>
mask	object to subset the image. If missing, then all values of the image are plotted.

## Value

Output of [boxplot](#)

## Examples

```

img = nifti(array(rnorm(10^3), dim = rep(10, 3)))
mask = img > 0
boxplot(img)
boxplot(img, mask = mask)
boxplot(as.anlz(img))

```

---

breaker	<i>Find Breaks for nifti Image Plotting</i>
---------	---

---

**Description**

Helper function for plotting - returns breaks for `image` plot function for object of class `nifti`

**Usage**

```
breaker(x, zlim, col, breaks = NULL)
```

**Arguments**

<code>x</code>	Object of class <code>nifti</code>
<code>zlim</code>	A user-specified <code>zlim</code> . If <code>NULL</code> , will calculate how <code>ortho2</code> would calculate <code>zlim</code>
<code>col</code>	colors to be plotted. Only used for <code>length(col)</code> , so can be a vector of length cols to be plotted
<code>breaks</code>	if <code>!is.null(breaks)</code> , then will calculate breaks. Otherwise will return this breaks vector

**Value**

Vector of length 2

If `breaks = NULL`, then vector of `length(col) + 1`, otherwise returns breaks

---

checking-methods	<i>Force object to filename</i>
------------------	---------------------------------

---

**Description**

Ensures the output to be a character filename (or vector) from an input image or `nifti`.

**Usage**

```
checking(file, allow_array = FALSE, ...)

## S4 method for signature 'nifti'
checking(file, allow_array = FALSE, ...)

## S4 method for signature 'ANY'
checking(file, allow_array = FALSE, ...)

## S4 method for signature 'character'
checking(file, allow_array = FALSE, ...)

## S4 method for signature 'list'
checking(file, allow_array = FALSE, ...)
```

**Arguments**

file            character or nifti object  
allow\_array    allow arrays to be passed in  
...            options passed to [temping](#)

**Value**

character filename of image or temporary nii, with .nii extension

**Author(s)**

John Muschelli <muschellij2@gmail.com>

---

checknii-methods      *Force object to filename with .nii extension*

---

**Description**

Ensures the output to be a character filename (or vector) from an input image or nifti, but not gzipped and has .nii extension

**Usage**

```
checknii(file, ...)

## S4 method for signature 'nifti'
checknii(file, ...)

## S4 method for signature 'factor'
checknii(file, ...)

## S4 method for signature 'character'
checknii(file, ...)

## S4 method for signature 'list'
checknii(file, ...)

## S4 method for signature 'ANY'
checknii(file, ...)

ensure_nii(file, ...)
```

**Arguments**

file            character or nifti object  
...            options passed to [checking](#)

**Value**

character filename of image or temporary nii, with .nii extension

**Author(s)**

John Muschelli <muschellij2@gmail.com>

**Examples**

```
set.seed(5)
dims = rep(10, 3)
arr = array(rnorm(prod(dims)), dim = dims)
arr[, ,10] = 0
nim = oro.nifti::nifti(arr)
rnifti = RNifti::asNifti(nim)
timg = tempimg(nim)
limg = list(factor(timg), factor(timg))
func = checknii
func(nim)
func(rnifti)
func(timg)
func(limg)
```

---

checkniigz-methods      *Force object to filename with .nii.gz extension*

---

**Description**

Ensures the output to be a character filename (or vector) from an input image or nifti, to be gzipped and has .nii.gz extension

**Usage**

```
checkniigz(file, ...)
```

## S4 method for signature 'nifti'

```
checkniigz(file, ...)
```

## S4 method for signature 'ANY'

```
checkniigz(file, ...)
```

## S4 method for signature 'factor'

```
checkniigz(file, ...)
```

## S4 method for signature 'character'

```
checkniigz(file, ...)
```

## S4 method for signature 'list'



```
checkniigz(file, ...)
```

```
ensure_nii_gz(file, ...)
```

### Arguments

file	character or nifti object
...	options passed to <a href="#">checking</a>

### Value

Character filename of image or temporary nii, with .nii.gz extension

### Author(s)

John Muschelli <muschellij2@gmail.com>

### Examples

```
set.seed(5)
dims = rep(10, 3)
arr = array(rnorm(prod(dims)), dim = dims)
arr[,,10] = 0
nim = oro.nifti::nifti(arr)
rnifti = RNifti::asNifti(nim)
ting = tempimg(nim)
limg = list(factor(ting), factor(ting))
func = checkniigz
func(nim)
func(rnifti)
func(ting)
func(limg)
```

---

check\_mask

*Check Mask is Binary*

---

### Description

Determine if only values in a mask are 0/1

### Usage

```
check_mask(mask, allow.NA = FALSE, allow.array = TRUE)
```

### Arguments

mask	Object of class nifti
allow.NA	allow NAs in the mask
allow.array	if class(mask) is "array", is this OK?

**Value**

Logical indicating if object is binary mask with only 0, 1, and NA if applicable

**Examples**

```
arr = array(rbinom(1000, size = 1, prob = 0.2), dim = c(10,10,10))
nim = oro.nifti::nifti(arr)
check_mask(nim)
```

---

check_mask_fail	<i>Check Mask is Binary, Fail otherwise</i>
-----------------	---

---

**Description**

Determine if only values in a mask are 0/1. Will error otherwise.

**Usage**

```
check_mask_fail(...)
```

**Arguments**

... arguments to pass to [check\\_mask](#)

**Value**

Either will error if conditions not met or an invisible NULL

**Examples**

```
arr = array(rbinom(1000, size = 1, prob = 0.2), dim = c(10,10,10))
nim = oro.nifti::nifti(arr)
check_mask_fail(nim)
```

---

check_nifti-methods	<i>Check if nifti image or read in a nifti image</i>
---------------------	--

---

**Description**

Simple check to see if input is character or of class nifti

**Usage**

```
check_nifti(  
  x,  
  reorient = FALSE,  
  allow.array = FALSE,  
  fast = FALSE,  
  need_header = TRUE,  
  ...  
)  
  
## S4 method for signature 'nifti'  
check_nifti(  
  x,  
  reorient = FALSE,  
  allow.array = FALSE,  
  fast = FALSE,  
  need_header = TRUE,  
  ...  
)  
  
## S4 method for signature 'character'  
check_nifti(  
  x,  
  reorient = FALSE,  
  allow.array = FALSE,  
  fast = FALSE,  
  need_header = TRUE,  
  ...  
)  
  
## S4 method for signature 'factor'  
check_nifti(  
  x,  
  reorient = FALSE,  
  allow.array = FALSE,  
  fast = FALSE,  
  need_header = TRUE,  
  ...  
)  
  
## S4 method for signature 'list'  
check_nifti(  
  x,  
  reorient = FALSE,  
  allow.array = FALSE,  
  fast = FALSE,  
  need_header = TRUE,  
  ...  
)
```

```

)

## S4 method for signature 'array'
check_nifti(
  x,
  reorient = FALSE,
  allow.array = FALSE,
  fast = FALSE,
  need_header = FALSE,
  ...
)

## S4 method for signature 'anlz'
check_nifti(
  x,
  reorient = FALSE,
  allow.array = FALSE,
  fast = FALSE,
  need_header = TRUE,
  ...
)

## S4 method for signature 'ANY'
check_nifti(
  x,
  reorient = FALSE,
  allow.array = FALSE,
  fast = FALSE,
  need_header = TRUE,
  ...
)

```

### Arguments

x	character path of image or an object of class nifti, or array
reorient	(logical) passed to <a href="#">readnii</a> if the image is to be re-oriented
allow.array	(logical) Are array types allowed (TRUE) or should there be an error if the object is not character or class nifti.
fast	if TRUE, then <a href="#">fast_readnii</a> will be used versus <a href="#">readnii</a> if the files need to be read in.
need_header	if TRUE, then an image type with header information will be returned. If not, then an array is fine. Used really only in conjunction with <code>allow.array</code>
...	additional arguments to pass to <a href="#">readnii</a> if relevant

### Value

nifti object or array if `allow.array=TRUE` and `x` is an array

**Author(s)**

John Muschelli <muschellij2@gmail.com>

**See Also**

[readnii](#)

**Examples**

```
x = nifti()
check_nifti(x)
set.seed(5)
dims = rep(10, 4)
arr = array(rpois(prod(dims), lambda = 2), dim = dims)
nim = oro.nifti::nifti(arr)
check_nifti(nim)
check_nifti(as.anlz(nim))
testthat::expect_error(check_nifti(arr, allow.array = FALSE))
tfile = tempimg(nim)
check_nifti(c(tfile, tfile))
check_nifti(list(tfile, tfile))
check_nifti(factor(c(tfile, tfile)))
check_nifti(RNifti::readNifti(tfile))
```

---

check\_nifti\_header-methods

*Check if nifti image or read in a nifti header*

---

**Description**

Simple check to see if input is character or of class nifti and read in the header

**Usage**

```
check_nifti_header(x)

## S4 method for signature 'nifti'
check_nifti_header(x)

## S4 method for signature 'character'
check_nifti_header(x)

## S4 method for signature 'factor'
check_nifti_header(x)

## S4 method for signature 'list'
check_nifti_header(x)
```

```
## S4 method for signature 'array'
check_nifti_header(x)

## S4 method for signature 'anlz'
check_nifti_header(x)

## S4 method for signature 'ANY'
check_nifti_header(x)
```

### Arguments

x                      character path of image or an object of class nifti, or array

### Value

nifti object or character

### Author(s)

John Muschelli <muschellij2@gmail.com>

### Examples

```
set.seed(5)
dims = rep(10, 4)
arr = array(rpois(prod(dims), lambda = 2), dim = dims)
nim = oro.nifti::nifti(arr)
check_nifti_header(nim)
check_nifti_header(as.anlz(nim))
testthat::expect_error(check_nifti_header(arr))
tfile = tempimg(nim)
check_nifti_header(tfile)
check_nifti_header(RNifti::readNifti(tfile))
check_nifti_header(c(tfile, tfile))
check_nifti_header(list(tfile, tfile))
check_nifti_header(factor(tfile))
```

---

check\_outfile

*Check output filename*

---

### Description

This function checks if an output filename is not NULL in conjunction whether the user would like to return an image

### Usage

```
check_outfile(outfile, retimg, fileext = ".nii.gz")
```

**Arguments**

outfile	output filename or NULL
retimg	Should an image be returned
fileext	a non-empty character vector giving the file extension

**Value**

Filename of output file or a temporary filename

---

cog	<i>Image Center of Gravity</i>
-----	--------------------------------

---

**Description**

Find Center of Gravity of Image, after thresholding

**Usage**

```
cog(img, thresh = 0, ceil = FALSE, warn = TRUE)
```

**Arguments**

img	Object of class nifti
thresh	threshold for image, will find $\text{img} > 0$
ceil	Run <a href="#">ceiling</a> to force integers (usu for plotting)
warn	Produce a warning if the image is empty after thresholding

**Value**

Vector of length 3

**Examples**

```
dims = rep(20, 3)
x = array(rnorm(prod(dims)), dim = dims)
img = nifti(x, dim= dims,
datatype = convert.datatype()$FLOAT32, cal.min = min(x),
cal.max = max(x), pixdim = rep(1, 4))
cog(img)
```

---

colorbar                      *Add a colorbar to an ortho2 object*

---

### Description

Adds a series of colors mapped to a value

### Usage

```
colorbar(breaks, col, text.col = "white", labels = TRUE, maxleft = 0.95)
```

### Arguments

breaks	a set of finite numeric breakpoints for the colours (see <a href="#">image</a> )
col	a list of colors (see <a href="#">image</a> )
text.col	axis and text label color
labels	labels for tick marks - see <a href="#">axis</a>
maxleft	Extent the left hand for colorbar

### Value

A plot

### Note

Much of this was taken from `vertical.image.legend` from the `aqfig` package

---

copyNIFTIHeader                      *Copy NIFTI Header to an array*

---

### Description

Copies slots of a `nifti` object to an array. This is useful if you're subsetting 4D data and getting an array out

### Usage

```
copyNIFTIHeader(
  img,
  arr,
  drop_slots = c(".Data", "dim_"),
  drop = TRUE,
  onlylast = TRUE,
  warn = TRUE,
  ...
)
```



**Arguments**

img	object of class nifti to copy header
arr	array to copy header information
drop_slots	Slots not to copy over from header
drop	Should <code>dropImageDimension</code> be called before returning?
onlylast	if drop = TRUE, passed to <code>dropImageDimension</code> , if only the last dimensions should be dropped
warn	if drop = TRUE, passed to <code>dropImageDimension</code> , for warning print out
...	arguments to pass to <code>nifti</code>

**Value**

Object of class nifti the size of arr

**Examples**

```
img = nifti(img = array(rnorm(10^4), dim=rep(10, 4)), dim=rep(10, 4), datatype = 16)
sub = img[,,1:3]
copyNIFTIHeader(img, sub)
sub = img[,,1, drop=FALSE]
copyNIFTIHeader(img, sub)
copyNIFTIHeader(img, sub, drop = FALSE)
```

---

cut.nifti

*Perform Cut on an image*


---

**Description**

Cuts a numeric image into an integer factor, with the option of a mask.

**Usage**

```
## S3 method for class 'nifti'
cut(x, breaks, ..., mask)

## S3 method for class 'anlz'
cut(x, ..., mask)
```

**Arguments**

x	Object of class nifti
breaks	either a numeric vector of two or more unique cut points or a single number (greater than or equal to 2) giving the number of intervals into which x is to be cut. Passed to <code>cut</code>
...	Arguments passed to <code>cut</code>
mask	object to subset the image. If missing, then all values of the image are used

**Value**

Object of class `nifti` with an attribute of levels

**Examples**

```
img = nifti(array(rnorm(10^3), dim = rep(10, 3)))
mask = img > 0
cut(img, mask = mask, breaks = 4)
```

---

datatype

*Change Data type for img*

---

**Description**

Tries to figure out the correct datatype for image. Useful for image masks - makes them binary if

**Usage**

```
datatypeper(
  img,
  type_string = NULL,
  datatype = NULL,
  bitpix = NULL,
  trybyte = TRUE,
  warn = TRUE
)
```

**Arguments**

<code>img</code>	nifti object (or character of filename)
<code>type_string</code>	(NULL) character of datatype and bitpix. Supercedes both datatype and bitpix. If specified <code>convert.datatype[[type_string]]</code> and <code>convert.bitpix[[type_string]]</code> will be used.
<code>datatype</code>	(NULL) character of datatype see <a href="#">convert.datatype</a>
<code>bitpix</code>	(NULL) character of bitpix see <a href="#">convert.bitpix</a>
<code>trybyte</code>	(logical) Should you try to make a byte (UINT8) if image in <code>c(0, 1)</code> ?
<code>warn</code>	Should a warning be issued if defaulting to FLOAT32?

**Value**

object of type `nifti`

**Examples**

```
img = nifti(array(rnorm(10^3, sd = 1000), dim = rep(10, 3)))
ring = round(img)
newnii(datatypeper(ring))
ring = datatypeper(ring, type_string= "FLOAT32")
```

---

density.nifti	<i>Density of Values in an Image</i>
---------------	--------------------------------------

---

**Description**

Computes the density of values of an image with the option for a mask.

**Usage**

```
## S3 method for class 'nifti'  
density(x, ..., mask)  
  
## S3 method for class 'anlz'  
density(x, ..., mask)
```

**Arguments**

x	Object of class nifti
...	Arguments passed to <a href="#">density.default</a>
mask	object to subset the image. If missing, then all values of the image are plotted.

**Value**

Output of [density](#)

**Examples**

```
img = nifti(array(rnorm(10^3), dim = rep(10, 3)))  
mask = img > 0  
density(img, mask = mask)  
density(img)  
  
density(as.anlz(img), mask = mask)  
density(as.anlz(img))
```

---

dicer	<i>Calculate Dice from a Table</i>
-------	------------------------------------

---

**Description**

Simple wrapper to calculate the Dice Coefficient/Similarity Index from a table

**Usage**

```
dicer(tab, verbose = TRUE)
```

**Arguments**

tab	table or matrix that is 2 by 2
verbose	should the Dice be printed before returned?

**Value**

Numeric scalar (one number)

**Examples**

```
tab = matrix(c(1000, 20, 20, 400), ncol = 2)
dicer(tab)
```

---

double_ortho	<i>Double Orthographic Display</i>
--------------	------------------------------------

---

**Description**

Copy of oro.nifti's [orthographic](#) function with some tweaks such as adding L/R designations for left and right

**Usage**

```
double_ortho(
  x,
  y = NULL,
  col.y = gray(0:64/64),
  NA.x = TRUE,
  mfrow = c(2, 4),
  add = FALSE,
  ...
)
```

**Arguments**

x	is an object of class nifti or similar.
y	is an object of class nifti or similar to be set aside x.
col.y	is grayscale (by default).
NA.x	Set any values of 0 in x to NA
mfrow	(numeric) layout of the 3 slices
add	Should the y-plot be added or its own plot? Used in double_ortho
...	other arguments to <a href="#">ortho2</a>

**See Also**

[orthographic](#)

---

`dropEmptyImageDimensions`*Drop Empty Image Dimensions*

---

## Description

Drops dimensions of an image that has all irrelevant values

## Usage

```
dropEmptyImageDimensions(  
  img,  
  value = 0,  
  threshold = 0,  
  other.imgs = NULL,  
  keep_ind = FALSE,  
  reorient = FALSE  
)
```

```
drop_empty_dim(  
  img,  
  value = 0,  
  threshold = 0,  
  other.imgs = NULL,  
  keep_ind = FALSE,  
  reorient = FALSE  
)
```

## Arguments

<code>img</code>	nifti object
<code>value</code>	Value to check against. If zero, then <code>dropEmptyImageDimensions</code> will drop any dimension that has fewer than <code>threshold</code> zeroes. May be a vector of values, matched with <a href="#">match</a>
<code>threshold</code>	Drop dimension if fewer than <code>threshold</code> voxels are in the slice
<code>other.imgs</code>	List of other nifti objects or filenames to apply the same dropping as <code>img</code> .
<code>keep_ind</code>	keep indices in output. Will return list, even if <code>other.imgs</code> not specified
<code>reorient</code>	Should image be reoriented if a filename?

## Value

List of output image indices, and other images if `other.imgs` not specified or `keep_ind = TRUE`. Otherwise object of class `nifti`

**Note**

drop\_empty\_dim is a shorthand for dropEmptyImageDimensions with all the same arguments. Also, NA are set to zero.

**See Also**

[getEmptyImageDimensions](#)

**Examples**

```
set.seed(5)
dims = rep(10, 3)
arr = array(rnorm(prod(dims)), dim = dims)
arr[, , 10] = 0
nim = oro.nifti::nifti(arr)

dnim = dropEmptyImageDimensions(nim, keep_ind = TRUE)
new_nim = dnim$outimg
names(dnim)
dnim = dropEmptyImageDimensions(nim, keep_ind = TRUE, other_imgs = nim)
dims = rep(10, 4)
arr = array(rnorm(prod(dims)), dim = dims)
nim = oro.nifti::nifti(arr)

testthat::expect_error(
{dnim = dropEmptyImageDimensions(nim, keep_ind = TRUE)}
)
```

---

emptyImageDimensionsMask

*Make Mask from Empty Image Dimensions*

---

**Description**

Make a mask of an image that has all irrelevant values

**Usage**

```
emptyImageDimensionsMask(img, ..., reorient = FALSE)
```

```
empty_dim_mask(img, ..., reorient = FALSE)
```

**Arguments**

img	nifti object
...	Arguments to be passed to <a href="#">getEmptyImageDimensions</a> .
reorient	Should image be reoriented if a filename?

**Value**

Object of class `nifti`, with binary values

**Note**

`empty_dim_mask` is a shorthand for `emptyImageDimensionsMask` with all the same arguments.

**See Also**

[getEmptyImageDimensions](#)

**Examples**

```
set.seed(5)
dims = rep(10, 3)
arr = array(rnorm(prod(dims)), dim = dims)
arr[,,10] = 0
nim = oro.nifti::nifti(arr)
out = emptyImageDimensionsMask(nim)
out_arr = emptyImageDimensionsMask(arr)
testthat::expect_equal(out_arr, array(out, dim = dim(out)))
out_arr = empty_dim_mask(arr)
```

---

ensure\_array

*Ensure an array output*

---

**Description**

Forces an array output for manipulation and overall conversion

**Usage**

```
ensure_array(img)
```

**Arguments**

`img` Image object ([nifti](#) or `niftiImage`)

**Value**

Array of same dimensions as image object

**Examples**

```

set.seed(5)
dims = rep(10, 3)
arr = array(rnorm(prod(dims)), dim = dims)
arr[,,10] = 0
nim = oro.nifti::nifti(arr)
rnifti = RNifti::asNifti(nim)
timg = tempimg(nim)
ling = list(factor(timg), factor(timg))
func = ensure_array
func(arr)
func(nim)
func(rnifti)
func(timg)
func(limg[[1]])

```

---

fast\_dice\_tab

*Fast Dice Tabulation*


---

**Description**

Fast Dice Tabulation

**Usage**

```
fast_dice_tab(x, y)
```

```
fast_dice(x, y, verbose = FALSE)
```

**Arguments**

x	A nifti image, filename, or niftiImage
y	A nifti image, filename, or niftiImage
verbose	A logical indicating output

**Value**

A table object

**Examples**

```

library(oro.nifti)
set.seed(20161007)
dims = rep(10, 3)
arr = array(rnorm(10*10*10), dim = dims)
nim = oro.nifti::nifti(arr) > -1
fast_dice_tab(nim, nim)
fast_dice(nim, nim) == 1

```



---

fast_readnii	<i>Reading NIfTI images through RNifti</i>
--------------	--

---

**Description**

This function calls the `readNifti` function from the `RNifti` package, and then converts the image to a `nifti` object

**Usage**

```
fast_readnii(fname, dtype = TRUE, drop_dim = TRUE)
```

**Arguments**

<code>fname</code>	file name of the NIfTI file.
<code>dtype</code>	Should <code>datatyper</code> be run after reading?
<code>drop_dim</code>	Should <code>drop_img_dim</code> be run after reading?

**Value**

A `nifti` object

**Examples**

```
set.seed(5)
dims = rep(10, 4)
arr = array(rpois(prod(dims), lambda = 2), dim = dims)
nim = oro.nifti::nifti(arr)
tfile = tempfile(fileext = ".nii.gz")
write_nifti(nim, tfile)
ring = fast_readnii(tfile)
```

---

file_imgext	<i>Get Image file extension</i>
-------------	---------------------------------

---

**Description**

Get the image file extension, either `.nii`, `.hdr`, `.nii.gz`, or `.hdr.gz`

**Usage**

```
file_imgext(file, withdot = TRUE)
```

**Arguments**

<code>file</code>	Vector of character filenames
<code>withdot</code>	Should the extension begin with <code>."</code> ?

**Value**

Vector of extensions. If withdot = FALSE, then will return nii instead of .nii

---

finite\_img-methods      *Finite Image*

---

**Description**

Simple wrapper for setting non-finite values to zero

**Usage**

```
finite_img(img, replace = 0)

## S4 method for signature 'nifti'
finite_img(img, replace = 0)

## S4 method for signature 'array'
finite_img(img, replace = 0)

## S4 method for signature 'ANY'
finite_img(img, replace = 0)

## S4 method for signature 'character'
finite_img(img, replace = 0)

## S4 method for signature 'list'
finite_img(img, replace = 0)
```

**Arguments**

img	character path of image or an object of class nifti, or list of images
replace	Value to replace non-finite values to

**Value**

nifti object

**Author(s)**

John Muschelli <muschellij2@gmail.com>

**Examples**

```

set.seed(5)
dims = rep(10, 3)
arr = array(rpois(prod(dims), lambda = 2), dim = dims)
arr[c(5, 6, 7, 8)] = c(NA, NaN, Inf, -Inf)
nim = nifti(arr)
finite_img(nim)
finite_img(arr)
tfile = tempimg(nim)
checking(c(tfile, tfile))
checking(list(tfile, tfile))
finite_img(list(tfile, tfile))
finite_img(c(tfile, tfile))
img = RNifti::readNifti(tfile)
checking(img)
img[c(5, 6, 7, 8)] = c(NA, NaN, Inf, -Inf)
stopifnot(!any(c(is.na(c(finite_img(img))))))

```

flip\_img

*Flip Nifti Image***Description**

This image will flip x, y, or z direction

**Usage**

```
flip_img(img, x = FALSE, y = FALSE, z = FALSE, ...)
```

**Arguments**

img	nifti object or character filename
x	(logical) Flip x direction
y	(logical) Flip y direction
z	(logical) Flip z direction
...	Arguments passed to <a href="#">check_nifti</a>

**Value**

Object of class nifti

**Examples**

```

img = random_nifti(rep(15, 3))
flipped = flip_img(img, x = TRUE, y = TRUE, z = TRUE)
img = random_nifti(rep(15, 2))
flipped = flip_img(img, x = TRUE)
testthat::expect_error(flip_img(img, z= TRUE))

```

---

`getEmptyImageDimensions`*Get Empty Image Dimensions*

---

**Description**

Creates a list of indices of an image that has all irrelevant values

**Usage**

```
getEmptyImageDimensions(img, value = 0, threshold = 0, reorient = FALSE, ...)
```

```
get_empty_dim(img, value = 0, threshold = 0, reorient = FALSE, ...)
```

**Arguments**

<code>img</code>	nifti object or array
<code>value</code>	Value to check against. If zero, then <code>getEmptyImageDimensions</code> will include any dimension that has fewer than <code>threshold</code> zeroes. May be a vector of values, matched with <a href="#">match</a>
<code>threshold</code>	Include dimension if fewer than <code>threshold</code> voxels are in the slice
<code>reorient</code>	Should image be reoriented if a filename?
<code>...</code>	additional arguments to pass to <a href="#">check_nifti</a>

**Value**

List of length 3 of indices.

**Note**

`get_empty_dim` is a shorthand for `getEmptyImageDimensions` with all the same arguments. Also, NA are set to zero.

**Examples**

```
arr = array(rbinom(1000, size = 1, prob = 0.2), dim = c(10,10,10))
arr[, ,1] = 0
arr[2:3, ,] = 0
getEmptyImageDimensions(arr)
```

---

hist.nifti	<i>Histogram of Values in an Image</i>
------------	--

---

**Description**

Computes and displays a histogram of the values of an image with the option for a mask.

**Usage**

```
## S3 method for class 'nifti'
hist(x, ..., mask)

## S3 method for class 'anlz'
hist(x, ..., mask)
```

**Arguments**

x	Object of class nifti
...	Arguments passed to <a href="#">hist.default</a>
mask	object to subset the image. If missing, then all values of the image are plotted.

**Value**

Output of [hist](#)

**Examples**

```
img = nifti(array(rnorm(10^3), dim = rep(10, 3)))
mask = img > 0
hist(img, mask = mask)
```

---

images2matrix	<i>Transform set of images to matrix</i>
---------------	--

---

**Description**

Creates a matrix, where the voxels are on the rows and images are on the columns

**Usage**

```
images2matrix(imgs, mask = NULL)
```

**Arguments**

imgs	Vector of files or list of images (niftiImage, array, or nifti)
mask	Binary image to subset the voxels

**Value**

Matrix of  $V$  by  $p$ , where  $V$  is the product of the dimensions of one image or the number of voxels in the mask, and  $p$  is the number of images

**Examples**

```
set.seed(5)
dims = rep(10, 3)
arr = array(rpois(prod(dims), lambda = 2), dim = dims)
nim = oro.nifti::nifti(arr)
imgs = list(nim, arr)
mask = nim > 2
mat1 = images2matrix(imgs)
mat2 = images2matrix(list(nim, nim))
if (packageVersion("oro.nifti") >= package_version("0.10.2")) {
  testthat::expect_equal(mat1, mat2)
} else {
  testthat::expect_error(testthat::expect_equal(mat1, mat2))
}
mat1 = images2matrix(imgs, mask = mask)
mat2 = images2matrix(list(nim, nim), mask)
```

img\_colour\_df

*Convert Image to Data.frame with Colors***Description**

Takes in an image and a color scheme, converts that image into a `data.frame` with the data and a color mapping.

**Usage**

```
img_colour_df(img, zlim = NULL, breaks = NULL, col = gray(0:64/64))

img_color_df(...)
```

**Arguments**

<code>img</code>	an object to be coerced to <code>nifti</code> using <code>check_nifti</code>
<code>zlim</code>	Limits for the domain of the intensities
<code>breaks</code>	Breaks for the intensities to map to colors
<code>col</code>	Colors to map intensities
<code>...</code>	not used

**Value**

A `data.frame` with the first columns being the  $x,y,z$  (maybe  $t$ ) coordinates (named `dim` and the dimension number), a `value` column that contains the intensity information, and a `colour` column representing the color that voxel maps to

**Note**

img\_color\_df is a duplicate of img\_colour\_df

**Examples**

```
img = nifti(array(rnorm(10^3), dim = rep(10, 3)))
df = img_colour_df(img)
df = img_color_df(img)
```

---

img\_indices

*Retrieve Image Indices*


---

**Description**

Extract image xyz indices (in voxels or millimeters), with the option to append the values

**Usage**

```
img_indices(img, mask = NULL, add_values = FALSE, units = c("index", "mm"))
```

**Arguments**

img	Object of class nifti
mask	Mask to be applied for indices the index
add_values	Should the value be column-bound to the matrix
units	Should the indices be in xyz-coordinates or millimeters.

**Value**

Matrix of 3 columns if add\_values = FALSE or 4 columns, otherwise.

**Examples**

```
set.seed(5)
dims = rep(10, 4)
arr = array(rpois(prod(dims), lambda = 2), dim = dims)
nim = oro.nifti::nifti(arr)
ind = img_indices(nim)
ind2 = img_indices(nim, mask = nim > 2)
# 3d example
set.seed(5)
dims = rep(10, 3)
arr = array(rpois(prod(dims), lambda = 2), dim = dims)
nim = oro.nifti::nifti(arr)
ind = img_indices(nim)
ind2 = img_indices(nim, mask = nim > 2)
testthat::expect_equal(colnames(ind2), c("x", "y", "z"))
ind2 = img_indices(nim, mask = nim > 2, add_values = TRUE)
testthat::expect_equal(colnames(ind2), c("x", "y", "z", "value"))
```

---

img\_list\_to\_ts      *Image List to Time Series*

---

**Description**

Turns a a list of 3D images into a 4D time series image

**Usage**

```
img_list_to_ts(imgs, copy_nifti = TRUE, warn = TRUE)
```

**Arguments**

imgs	object of class <code>list</code> , each with 3 dimensions,
copy_nifti	Should a nifti object be returned (TRUE) or a simply array (FALSE). Should only be used for slight speed up when array is adequate
warn	Should a warning be printed if object is not class <code>nifti</code>

**Value**

Object of class `nifti`

**Note**

If the object is not of class `list`, then the object is returned

---

img\_ts\_to\_df      *Image Time Series to Data.frame*

---

**Description**

Turns a 4D time series image to a Data.frame

**Usage**

```
img_ts_to_df(imgs, warn = FALSE)
```

**Arguments**

imgs	object of class <code>nifti</code> with 4 dimensions, aka a 4D time series
warn	Should a warning be printed if object is not class <code>nifti</code> (e.g. a list instead)

**Value**

Matrix of values



---

img_ts_to_list	<i>Image Time Series to list</i>
----------------	----------------------------------

---

## Description

Turns a 4D time series image to a list of 3D images

## Usage

```
img_ts_to_list(imgs, copy_nifti = TRUE, warn = TRUE)
```

## Arguments

imgs	object of class <code>nifti</code> with 4 dimensions, aka a 4D time series
copy_nifti	Should <code>nifti</code> objects be returned (TRUE) or simply arrays (FALSE). Should only be used for slight speed up when array is adequate
warn	Should a warning be printed if object is not class <code>nifti</code>

## Value

List of images

## Note

If the object is not of class `nifti` or have 4 dimensions, then the object is returned

## Examples

```
set.seed(5)
dims = rep(10, 4)
arr = array(rpois(prod(dims), lambda = 2), dim = dims)
nim = oro.nifti::nifti(arr)
simg = img_ts_to_list(nim)
simg_arr = img_ts_to_list(arr)
back = img_list_to_ts(simg_arr)
slist = lapply(simg, function(x) array(x, dim(x)))
testthat::expect_equal(slist, simg_arr)
simg_arr = img_ts_to_matrix(arr)
simg_arr = img_ts_to_df(arr)
```

---

img\_ts\_to\_matrix      *Image Time Series to Matrix*

---

**Description**

Turns a 4D time series image to a Matrix

**Usage**

```
img_ts_to_matrix(imgs, warn = FALSE)
```

**Arguments**

imgs                    object of class `nifti` with 4 dimensions, aka a 4D time series  
warn                    Should a warning be printed if object is not class `nifti` (e.g. a list instead)

**Value**

Matrix of values

---

maskEmptyImageDimensions-methods  
*Apply Masking from Empty Image Dimensions*

---

**Description**

Simple wrapper for replacing indices with a value

**Usage**

```
maskEmptyImageDimensions(img, inds, reorient = FALSE, mask.value = 0, ...)
```

```
## S4 method for signature 'nifti'
```

```
maskEmptyImageDimensions(img, inds, reorient = FALSE, mask.value = 0, ...)
```

```
## S4 method for signature 'character'
```

```
maskEmptyImageDimensions(img, inds, reorient = FALSE, mask.value = 0, ...)
```

```
## S4 method for signature 'factor'
```

```
maskEmptyImageDimensions(img, inds, reorient = FALSE, mask.value = 0, ...)
```

```
## S4 method for signature 'list'
```

```
maskEmptyImageDimensions(img, inds, reorient = FALSE, mask.value = 0, ...)
```

```
## S4 method for signature 'array'
```

```

maskEmptyImageDimensions(img, inds, reorient = FALSE, mask.value = 0, ...)

## S4 method for signature 'anlz'
maskEmptyImageDimensions(img, inds, reorient = FALSE, mask.value = 0, ...)

## S4 method for signature 'ANY'
maskEmptyImageDimensions(img, inds, reorient = FALSE, mask.value = 0, ...)

mask_empty_dim(img, ...)

```

**Arguments**

img	image, nifti object, or array
inds	indices of subset from <a href="#">getEmptyImageDimensions</a> or <a href="#">dropEmptyImageDimensions</a> .
reorient	Should image be reoriented if a filename?
mask.value	Value to replace voxels outside the mask.
...	not used

**Value**

Object of class nifti or array if nifti is not supplied

**Note**

mask\_empty\_dim is a shorthand for maskEmptyImageDimensions with all the same arguments.

**See Also**

[getEmptyImageDimensions](#), [dropEmptyImageDimensions](#)

**Examples**

```

set.seed(5)
dims = rep(10, 3)
arr = array(rnorm(prod(dims)), dim = dims)
arr[, ,10] = 0
nim = oro.nifti::nifti(arr)
inds = getEmptyImageDimensions(nim)
inds_arr = getEmptyImageDimensions(arr)
res = maskEmptyImageDimensions(nim, inds = inds, mask.value = NA)
res2 = maskEmptyImageDimensions(arr, inds = inds_arr, mask.value = NA)
testthat::expect_equal(array(res, dim = dim(res)),
array(res2, dim = dim(res2)))

set.seed(5)
dims = rep(10, 3)
arr = array(rnorm(prod(dims)), dim = dims)
arr[, ,10] = 0
nim = oro.nifti::nifti(arr)
inds = getEmptyImageDimensions(nim)

```

```

rnifti = RNifti::asNifti(nim)
ting = tempimg(nim)
ling = list(factor(ting), factor(ting))
mask_empty_dim(nim, inds = inds)
func = function(...) maskEmptyImageDimensions(..., inds = inds)
func(arr)
func(nim)
func(rnifti)
func(ting)
func(limg)

```

---

mask\_img

*Mask Image*


---

### Description

Takes an image, masks it by mask, and returns an object of class `nifti`

### Usage

```
mask_img(img, mask, allow.NA = TRUE)
```

### Arguments

<code>img</code>	object of class <code>nifti</code>
<code>mask</code>	array or object of class <code>nifti</code> , same dimensions as <code>img</code>
<code>allow.NA</code>	allow NAs in the mask

### Value

Object of class `nifti` with values outside mask set to 0 if mask is 0 and NA if mask is NA and `img` if `mask == 1`

### Examples

```

set.seed(5)
dims = rep(10, 3)
arr = array(rnorm(prod(dims)), dim = dims)
nim = oro.nifti::nifti(arr)
mask = abs(nim) > 1
masked = mask_img(nim, mask)
mask[mask == 0] = NA
na_masked = mask_img(nim, mask, allow.NA = TRUE)

set.seed(5)
dims = rep(10, 3)
arr = array(rnorm(prod(dims)), dim = dims)
arr[, , 10] = 0

```

```

nim = oro.nifti::nifti(arr)
rnifti = RNifti::asNifti(nim)
mask = nim > 0
ting = tempimg(nim)
limg = list(factor(ting), factor(ting))
func = function(...) mask_img(..., mask = mask)
func(arr)
func(nim)
func(rnifti)
func(ting)
lapply(limg, func)

```

---

mask\_vals

*Extract or Replace Values Inside a Mask*


---

## Description

Methods that act on the `.Data` field in a NIFTI/ANALYZE image but only on values inside a mask.

## Usage

```

mask_vals(object, mask)

mask_vals(object, mask) <- value

## S4 replacement method for signature 'nifti'
mask_vals(object, mask) <- value

## S4 replacement method for signature 'anlz'
mask_vals(object, mask) <- value

## S4 replacement method for signature 'array'
mask_vals(object, mask) <- value

```

## Arguments

<code>object</code>	is an object of class <code>nifti</code> or <code>anlz</code> .
<code>mask</code>	is an object of class <code>nifti</code> or <code>anlz</code> .
<code>value</code>	is the value to assign to the <code>.Data</code> field.

## Examples

```

set.seed(2022)
img = nifti(array(rnorm(10^3), dim = rep(10, 3)))
mask = img > 1.5
mask_vals(img, mask)
testthat::expect_equal(sum(mask_vals(img, mask)), 117.628200302518)
mask_vals(img, mask) = rep(4, sum(mask))

```

```
mask_vals(img, as(mask, "array")) = rep(4, sum(mask))
mask_vals(as(img, "array"),
          as(mask, "array")) = rep(4, sum(mask))
```

---

mean.nifti	<i>Mean of Values in an Image</i>
------------	-----------------------------------

---

### Description

Computes the mean of values of an image with the option for a mask.

### Usage

```
## S3 method for class 'nifti'
mean(x, ..., mask)

## S3 method for class 'anlz'
mean(x, ..., mask)
```

### Arguments

x	Object of class nifti
...	Arguments passed to <a href="#">mean.default</a>
mask	object to subset the image. If missing, then all values of the image are plotted.

### Value

Output of [mean](#)

### Examples

```
img = nifti(array(rnorm(10^3), dim = rep(10, 3)))
mask = img > 0
mean(img, mask = mask)
```

---

minmax_img-methods	<i>Normalize Image using Range</i>
--------------------	------------------------------------

---

### Description

Calculates the range of values in an image, then scales the image minimum to 0 and maximum to 1

**Usage**

```

minmax_img(img)

## S4 method for signature 'nifti'
minmax_img(img)

## S4 method for signature 'array'
minmax_img(img)

## S4 method for signature 'ANY'
minmax_img(img)

## S4 method for signature 'character'
minmax_img(img)

## S4 method for signature 'factor'
minmax_img(img)

## S4 method for signature 'list'
minmax_img(img)

```

**Arguments**

`img` character path of image or an object of class `nifti`, or list of images

**Value**

A `nifti` object (or list of them) or class of object passed in if not specified

**Examples**

```

set.seed(5)
dims = rep(10, 4)
arr = array(rpois(prod(dims), lambda = 2), dim = dims)
nim = oro.nifti::nifti(arr)
mimg = minmax_img(nim)
marr = minmax_img(arr)
testthat::expect_equal(array(mimg, dim = dim(mimg)), marr)

set.seed(5)
dims = rep(10, 3)
arr = array(rnorm(prod(dims)), dim = dims)
arr[,,10] = 0
nim = oro.nifti::nifti(arr)
rnifti = RNifti::asNifti(nim)
timg = tempimg(nim)
limg = list(factor(timg), factor(timg))
func = minmax_img
func(arr)
func(nim)

```

```
func(rnifti)
func(timg)
func(limg)
```

---

```
multi_overlay
```

```
Create Multi-Image Plot with Overlays
```

---

## Description

Creates a multi-row or multi-column plot with image slices and the potential for overlays as well.

## Usage

```
multi_overlay(
  x,
  y = NULL,
  z = NULL,
  w = 1,
  mask = NULL,
  col.x = gray(0:64/64),
  col.y = hotmetal(),
  zlim.x = NULL,
  zlim.y = NULL,
  ybreaks = NULL,
  plane = c("axial", "coronal", "sagittal"),
  xlab = "",
  ylab = "",
  axes = FALSE,
  direction = c("horizontal", "vertical"),
  par.opts = list(oma = c(0, 0, 0, 0), mar = rep(0, 4), bg = "black"),
  text = NULL,
  text.x = 0.5,
  text.y = 1.4,
  text.cex = 2.5,
  text.col = "white",
  main = NULL,
  main.col = text.col,
  main.cex = text.cex,
  NA.x = TRUE,
  NA.y = TRUE,
  pdim = NULL,
  useRaster = TRUE,
  ...
)

multi_overlay_center(x, y = NULL, ...)
```



**Arguments**

x	List of images of class <code>nifti</code> or character vector of filenames
y	List of images of class <code>nifti</code> or character vector of filenames. Same length as x.
z	Slice to display.
w	3D volume to display if x has 4-D elements
mask	<code>nifti</code> image to drop empty image dimensions if wanted. Passed to <a href="#">dropEmptyImageDimensions</a>
col.x	Color to display x images
col.y	Color to display y images
zlim.x	Limits for x to plot
zlim.y	Limits for y to plot
ybreaks	(numeric) breaks for y to passed to <a href="#">image</a>
plane	the plane of acquisition to be displayed
xlab	Label for x-axis
ylab	Label for y-axis
axes	Should axes be displayed
direction	Should images be a row or column? Ignored if <code>mfrow</code> is in <code>par.opts</code>
par.opts	Options to pass to <a href="#">par</a>
text	Text to be displayed
text.x	Location of text in x-domain
text.y	Location of text in y-domain
text.cex	Multiplier for text font
text.col	Color for text and main.
main	Title for each plot
main.col	Color for main. Will default to <code>text.col</code>
main.cex	Multiplier for text font. Will default to <code>text.cex</code>
NA.x	Should 0's in x be set to NA?
NA.y	Should 0's in y be set to NA?
pdim	Pixel dimensions if passing in arrays. Will be overridden if x is a <code>nifti</code> object
useRaster	if TRUE, a bitmap raster is used to plot the image instead of polygons. Passed to <a href="#">image</a>
...	Additional arguments to pass to <a href="#">image</a>

**Examples**

```

set.seed(5)
dims = rep(10, 4)
arr = array(rnorm(prod(dims)), dim = dims)
arr[,,c(3, 5)] = rpois(1000*2, lambda = 2)
nim = oro.nifti::nifti(arr)
mask = nim > 2
add_mask = nim[,,1] > 0
imgs = img_ts_to_list(nim)
masks = img_ts_to_list(mask)
multi_overlay(imgs, masks)
multi_overlay(imgs, masks,
main = "hey", direction = "vertical", plane = "coronal")
multi_overlay(imgs, masks, mask = add_mask,
main = "hey")

```

## Not run:

```

if (requireNamespace("brainR", quietly = TRUE)) {
  visits = 1:3
  y = paste0("Visit_", visits, ".nii.gz")
  y = system.file(y, package = "brainR")
  y = lapply(y, readnii)

  y = lapply(y, function(r){
    pixdim(r) = c(0, rep(1, 3), rep(0, 4))
    dropImageDimension(r)
  })

  x = system.file("MNI152_T1_1mm_brain.nii.gz",
    package = "brainR")
  x = readnii(x)
  mask = x > 0
  x = lapply(visits, function(tmp){
    x
  })
  alpha = function(col, alpha = 1) {
    cols = t(col2rgb(col, alpha = FALSE)/255)
    rgb(cols, alpha = alpha)
  }
  multi_overlay(x, y,
    col.y = alpha(hotmetal(), 0.5),
    mask = mask,
    main = paste0("\n", "Visit ", visits),
    text = LETTERS[visits],
    text.x = 0.9,
    text.y = 0.1,
    text.cex = 3)
}

```

```
## End(Not run)
```

---

newnii	<i>Resets image parameters for a copied nifti object</i>
--------	--

---

### Description

Resets the slots of a nifti object, usually because an image was loaded, then copied and filled in with new data instead of making a nifti object from scratch. Just a wrapper for smaller functions

### Usage

```
newnii(img, ...)
```

### Arguments

img	nifti object (or character of filename)
...	arguments to be passed to <code>datatype</code>

### Value

object of type nifti

---

niftiarr	<i>Make new nifti from array</i>
----------	----------------------------------

---

### Description

Make new nifti object by passing in old nifti and array

### Usage

```
niftiarr(img, arr)
```

### Arguments

img	object of class nifti
arr	array to be passed in to .Data slot

### Value

object of class nifti

---

nii.stub	<i>Grab nii file stubname</i>
----------	-------------------------------

---

**Description**

Quick helper function to strip off .nii or .nii.gz from filename

**Usage**

```
nii.stub(x, bn = FALSE)
```

**Arguments**

x	character vector of filenames ending in .nii or .nii.gz
bn	Take <a href="#">basename</a> of file?

**Value**

A character vector with the same length as x

---

orient_rpi	<i>Reorient an Image to RPI orientation</i>
------------	---

---

**Description**

Reorient an Image to RPI orientation

**Usage**

```
orient_rpi(file, verbose = TRUE)
orient_rpi_file(file, verbose = TRUE)
is_rpi_oriented(file, verbose = FALSE)
```

**Arguments**

file	Object of class <code>nifti</code> or character path
verbose	print diagnostic messages

**Value**

List of 3 elements

- `img`: Reoriented image of class `nifti`
- `convention`: Convention (Neurological/Radiological) of original image
- `orientation`: Original image orientations

**Note**

'orient\_rpi' and 'orient\_rpi\_file' uses 'RNifti' to ensure the reading orientation

**Examples**

```
lr_fname = system.file( "nifti", "mniLR.nii.gz", package = "oro.nifti")
img = readnii(lr_fname)

rl_fname = system.file( "nifti", "mniRL.nii.gz", package = "oro.nifti")
rl_img = readnii(rl_fname)
stopifnot(all(rl_img[nrow(rl_img):1,,] == img))

reor = orient_rpi(rl_fname)
stopifnot(all(img == reor$img))

rev = reverse_orient_rpi(reor$img, convention = reor$convention,
orientation = reor$orientation)
stopifnot(all(rev == rl_img))
```

---

ortho2

*Orthographic Display, added options*


---

**Description**

Copy of oro.nifti's [orthographic](#) function with some tweaks such as adding L/R designations for left and right

**Usage**

```
ortho2(
  x,
  y = NULL,
  xyz = NULL,
  w = 1,
  col = gray(0:64/64),
  col.y = oro.nifti::hotmetal(),
  zlim = NULL,
  zlim.y = NULL,
  NA.x = FALSE,
  NA.y = TRUE,
  crosshairs = TRUE,
  col.crosshairs = "red",
  xlab = "",
  ylab = "",
  axes = FALSE,
  oma = c(0, 0, 0, ifelse(ycolorbar, 5, 0)),
  mar = rep(0, 4),
  bg = "black",
```

```

text = NULL,
text.color = "white",
text.cex = 2,
text.x = 32,
text.y = 32,
add.orient = FALSE,
mfrow = c(2, 2),
ybreaks = NULL,
breaks = NULL,
addlegend = FALSE,
leg.x = 32,
leg.y = 32,
legend,
leg.col,
leg.title = NULL,
leg.cex,
window = NULL,
ycolorbar = FALSE,
clabels = TRUE,
add = TRUE,
pdim = NULL,
useRaster = is.null(y),
mask = NULL,
...
)

```

### Arguments

<code>x</code>	is an object of class <code>nifti</code> or similar.
<code>y</code>	is an object of class <code>nifti</code> or similar for the overlay.
<code>xyz</code>	is the coordinate for the center of the crosshairs.
<code>w</code>	is the time point to be displayed (4D arrays only).
<code>col</code>	is grayscale (by default).
<code>col.y</code>	is hotmetal (by default).
<code>zlim</code>	is the minimum and maximum 'z' values passed into image.
<code>zlim.y</code>	is the minimum and maximum 'z' values passed into image for the overlay.
<code>NA.x</code>	Set any values of 0 in x to NA
<code>NA.y</code>	Set any values of 0 in y to NA
<code>crosshairs</code>	is a logical value for the presence of crosshairs in all three orthogonal planes (default = TRUE).
<code>col.crosshairs</code>	is the color of the crosshairs (default = red).
<code>xlab</code>	is set to "" since all margins are set to zero.
<code>ylab</code>	is set to "" since all margins are set to zero.
<code>axes</code>	is set to FALSE since all margins are set to zero.

oma	is the size of the outer margins in the par function.
mar	is the number of lines of margin in the par function.
bg	is the background color in the par function.
text	allows the user to specify text to appear in the fourth (unused) pane.
text.color	is the color of the user-specified text (default = "white").
text.cex	is the size of the user-specified text (default = 2).
text.x	x coordinate for text
text.y	y coordinate for text
add.orient	(logical) Add left/right, A/P, etc. orientation
mfrow	(numeric) layout of the 3 slices
ybreaks	(numeric) breaks for y to passed to <a href="#">image</a>
breaks	(numeric) breaks for x to passed to <a href="#">image</a>
addlegend	(logical) add legend?
leg.x	(numeric) x coordinate for legend
leg.y	(numeric) y coordinate for legend
legend	(character) legend text
leg.col	(character) Colors for legend
leg.title	(character) title for legend
leg.cex	(numeric) cex for <a href="#">legend</a>
window	(vector) Length-2 vector to limit image to certain range
ycolorbar	(logical) Should a colorbar for y be plotted
clabels	Label for colorbar (see <a href="#">colorbar</a> )
add	Should the y-plot be added or its own plot? Used in <code>double_ortho</code>
pdim	Pixel dimensions if passing in arrays. Will be overridden if x is a <code>nifti</code> object
useRaster	logical; if TRUE a bitmap raster is used to plot the image instead of polygons. Passed to <a href="#">image</a> .
mask	If a mask is passed, <code>drop_empty_dim</code> is applied to both x and y
...	other arguments to the image function may be provided here.

**See Also**

[orthographic](#)

**Examples**

```
set.seed(10)
x = oro.nifti::nifti(array(rnorm(1000), dim = rep(10, 3)))
ortho2(x)
y = x > 2
mask = x > 2.5
ortho2(x, y)
```

```

ortho2(x, y, mask = mask, add.orient = TRUE)
ortho2(x, y, mask = mask, add.orient = TRUE, add = FALSE)
nim = RNifti::asNifti(x, internal = FALSE)
ortho2(nim, y, mask = mask)
neurobase::ortho2(nim, x, mask = mask,
ybreaks = seq(min(x), max(x), length.out = 65), ycolorbar = TRUE)

ortho2(nim, y, mask = mask, add = FALSE)
arr_x = as.array(x)
arr_y = as.array(y)
ortho2( arr_x)
ortho2( arr_x, arr_y, useRaster = FALSE)

set.seed(10)
x = oro.nifti::nifti(array(rnorm(10000), dim = rep(10, 4)))
y = x > 2
mask = x > 2.5
ortho2(x, y)

set.seed(10)
x = oro.nifti::nifti(array(rnorm(100), dim = rep(10, 2)))
y = x > 2
mask = x > 2.5
ortho2(x, y)

```

---

ortho\_diff

*Plot differences for Prediction and Gold Standard*

---

### Description

Uses `ortho2` to plot differences between a predicted binary image and the assumed ground truth (roi).

### Usage

```

ortho_diff(
  img,
  pred,
  roi,
  xyz = NULL,
  cols = c("#56B4E9", "#D55E00", "#009E73"),
  levels = c("False Negative", "False Positive", "True Positive"),
  addlegend = TRUE,
  center = TRUE,
  leg.cex = 1.5,
  ...
)

multi_overlay_diff(

```



```

    x,
    pred,
    roi,
    z = NULL,
    cols = c("#56B4E9", "#D55E00", "#009E73"),
    ...
)

```

### Arguments

img	image to be underlaid
pred	binary segmentation (prediction)
roi	binary manual segmentation (ground truth)
xyz	coordinate for the center of the crosshairs.
cols	colors for false negatives/positives
levels	labels for false negatives/positives
addlegend	add legend, passed to <a href="#">ortho2</a>
center	run <a href="#">xyz</a> on <a href="#">roi</a> . Disregarded if <a href="#">xyz</a> is not NULL
leg.cex	multiplier for legend size
...	arguments to be passed to <a href="#">ortho2</a> or <a href="#">multi_overlay</a>
x	List of images of class <code>nifti</code> or character vector of filenames
z	slice to display

### See Also

[ortho2](#)

### Examples

```

set.seed(5)
dims = rep(10, 3)
arr = array(rpois(prod(dims), lambda = 2), dim = dims)
nim = oro.nifti::nifti(arr)
roi = nim > 2
pred = nim > 1.5
ortho_diff(nim, pred, roi)
set.seed(5)
dims = rep(10, 3)
arr = array(rnorm(prod(dims)), dim = dims)
nim = oro.nifti::nifti(arr)
mask = nim > 2
pred = nim > 1.5
multi_overlay_diff(nim, roi = mask, pred = pred)

```

```

if (requireNamespace("brainR", quietly = TRUE)) {
  visits = 1:3
  y = paste0("Visit_", visits, ".nii.gz")
  y = system.file(y, package = "brainR")
  y = lapply(y, readnii)

  y = lapply(y, function(r){
    pixdim(r) = c(0, rep(1, 3), rep(0, 4))
    dropImageDimension(r)
  })

  x = system.file("MNI152_T1_1mm_brain.nii.gz",
                 package = "brainR")
  x = readnii(x)
  mask = x > 0
  alpha = function(col, alpha = 1) {
    cols = t(col2rgb(col, alpha = FALSE)/255)
    rgb(cols, alpha = alpha)
  }
  roi = y[[2]]
  pred = y
  multi_overlay_diff(x, roi = roi, pred = pred)
  multi_overlay_diff(x, roi = roi, pred = pred,
                    mask = mask,
                    main = paste0("\n", "Visit ", visits),
                    text = LETTERS[visits],
                    text.x = 0.9,
                    text.y = 0.1,
                    text.cex = 3)
}

```

---

parse\_img\_ext

*Parse Image Extensions*

---

## Description

Get image extensions from a filename

## Usage

```
parse_img_ext(file)
```

## Arguments

file                    Character filename to parse

## Value

Extension of file

## Examples

```
parse_img_ext("blah.nii.gz")
parse_img_ext("blah.mnc")
parse_img_ext("blah.nii")
parse_img_ext("blah")
parse_img_ext("blah.img")
parse_img_ext("blah.hdr")
parse_img_ext("blah.hdr.gz")
```

---

quantile.nifti

*Sample Quantiles*

---

## Description

Computes sample quantiles for an image, with the option of a mask.

## Usage

```
## S3 method for class 'nifti'
quantile(x, ..., mask)

## S3 method for class 'anlz'
quantile(x, ..., mask)
```

## Arguments

x	Object of class nifti
...	Arguments passed to <a href="#">quantile</a>
mask	object to subset the image. If missing, then all values of the image are used

## Value

Output of [quantile](#)

## Examples

```
img = nifti(array(rnorm(10^3), dim = rep(10, 3)))
mask = img > 0
quantile(img, mask = mask)
```

---

**quantile\_img**      *Create Quantile Image*

---

**Description**

Creates output image of the quantiles that each voxel is in, after applying the mask

**Usage**

```
quantile_img(img, mask = NULL, ...)
```

**Arguments**

<code>img</code>	Character vector, or object of class <code>nifti</code>
<code>mask</code>	Mask to determine cumulative distribution function (cdf) from
<code>...</code>	Additional arguments to pass to <a href="#">ecdf</a>

**Value**

Object of class `nifti`

**Examples**

```
set.seed(5)
dims = rep(10, 3)
arr = array(rnorm(prod(dims)), dim = dims)
nim = oro.nifti::nifti(arr)
qimg = quantile_img(nim)
qarr = quantile_img(arr)
testthat::expect_equal(qarr, array(qimg, dim = dim(qarr)))
qimg = quantile_img(nim, mask = nim > 0)
```

---

**randomize\_mask**      *Randomize Image based on Mask*

---

**Description**

Randomize the values within a mask

**Usage**

```
randomize_mask(img, mask)
```

**Arguments**

img	Object of class nifti with values to be randomized
mask	Binary mask indicating which values should be randomized.

**Value**

Object of class nifti

**Examples**

```
set.seed(5)
dims = rep(10, 3)
arr = array(rnorm(prod(dims)), dim = dims)
nim = oro.nifti::nifti(arr)
mask = abs(nim) > 1
randomize_mask(nim, mask)
```

---

random_nifti	<i>Create Random 'nifti' object</i>
--------------	-------------------------------------

---

**Description**

Create Random 'nifti' object

**Usage**

```
random_nifti(dim, ...)
```

**Arguments**

dim	dimensions for the 'nifti' object
...	arguments to send to <a href="#">nifti</a>

**Value**

A 'nifti' object

**Examples**

```
random_nifti(c(10, 10, 2))
random_nifti(c(10, 10))
```

---

 readNIFTI2

*readNIFTI with default non-reorientation*


---

### Description

This function calls the [readNIFTI](#) function from the `oro.nifti` package, but sets the reorientation to FALSE by default

### Usage

```
readNIFTI2(..., reorient = FALSE)
```

```
readnii(
  ...,
  reorient = FALSE,
  dtype = TRUE,
  drop_dim = TRUE,
  reset_slope = FALSE,
  warn = FALSE,
  rm_extensions = TRUE
)
```

### Arguments

...	Arguments to pass to <a href="#">readNIFTI</a>
reorient	Reorientation argument to pass to <a href="#">readNIFTI</a>
dtype	Should <a href="#">datatyper</a> be run after reading?
drop_dim	Should <a href="#">drop_img_dim</a> be run after reading?
reset_slope	Reset slope/intercept of image
warn	Should warnings from <a href="#">readNIFTI</a> be printed? If not, <a href="#">suppressWarnings</a> is called. Also passed to <a href="#">datatyper</a>
rm_extensions	should <code>niftiExtensions</code> be converted to simple nifti objects?

### Value

nifti object

---

read_rpi	<i>Read NIfTI file reoriented to RPI</i>
----------	--

---

**Description**

This function calls the `readnii` function after calling `orient_rpi_file` to force RPI orientation.

**Usage**

```
read_rpi(file, ..., verbose = TRUE)
```

**Arguments**

file	file name of the NIfTI file.
...	Arguments to pass to <code>readnii</code>
verbose	print diagnostics, passed to <code>orient_rpi_file</code>

**Note**

'read\_rpi' uses 'RNifti' to ensure the reading orientation

---

remake_img	<i>Remake Image from Vector</i>
------------	---------------------------------

---

**Description**

Wrapper function to take a vector of values and result in a `nifti` object

**Usage**

```
remake_img(vec, img, mask = NULL, warn = FALSE, ...)
```

**Arguments**

vec	vector of values to be in resulting image
img	object of class <code>nifti</code> to put vector into
mask	binary array/ <code>nifti</code> object to denote where vector values are to be.
warn	Should a warning be issued if defaulting to FLOAT32?
...	additional arguments passed to <code>datatyper</code>

**Value**

Object of class `nifti`

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

```

set.seed(5)
dims = rep(10, 3)
arr = array(rnorm(prod(dims)), dim = dims)
arr[, , 10] = 0
nim = oro.nifti::nifti(arr)
remake_img(c(nim), nim)
mask = nim > 5
vals = nim[mask]
vals = sqrt(vals)
remake_img(vals, nim, mask = mask)

```

---

remap_filename	<i>Build Filename (usually for images)</i>
----------------	--

---

**Description**

This is a simple function that helps with the case where you want to construct a filename (usually for an image) with the same base of the filename, the same directory (default), but things added to the front or end of that base filename, with the same extension.

**Usage**

```
remap_filename(x, sub_dir = NULL, prefix = "", suffix = "")
```

**Arguments**

x	input filename/character vector
sub_dir	sub-directory for the new filename. If NULL, then the directory is the the same directory as x
prefix	string to put in front of base of filename
suffix	string to put at the end of base of filename

**Value**

Character vector

**Examples**

```

fname = file.path("/path/to/file", "original.nii.gz")
remap_filename(fname, prefix = "preproc_", "_with_gz")
fname = "original.nii"
remap_filename(fname, prefix = "note_", "_has_directory")
remap_filename(c(fname, "other.nii.gz"), prefix = "note_", "_has_directory")

```



---

`replaceEmptyImageDimensions-methods`*Replace Subsetting from Empty Image Dimensions*

---

**Description**

Simple wrapper for subsetting an image with indices, dropping empty dimensions.

**Usage**

```
replaceEmptyImageDimensions(  
  img,  
  inds,  
  target_dim,  
  value = 0,  
  reorient = FALSE,  
  ...  
)  
  
## S4 method for signature 'nifti'  
replaceEmptyImageDimensions(  
  img,  
  inds,  
  target_dim,  
  value = 0,  
  reorient = FALSE,  
  ...  
)  
  
## S4 method for signature 'character'  
replaceEmptyImageDimensions(  
  img,  
  inds,  
  target_dim,  
  value = 0,  
  reorient = FALSE,  
  ...  
)  
  
## S4 method for signature 'factor'  
replaceEmptyImageDimensions(  
  img,  
  inds,  
  target_dim,  
  value = 0,  
  reorient = FALSE,  
  ...  
)
```

```
)

## S4 method for signature 'list'
replaceEmptyImageDimensions(
  img,
  inds,
  target_dim,
  value = 0,
  reorient = FALSE,
  ...
)

## S4 method for signature 'array'
replaceEmptyImageDimensions(
  img,
  inds,
  target_dim,
  value = 0,
  reorient = FALSE,
  ...
)

## S4 method for signature 'anlz'
replaceEmptyImageDimensions(
  img,
  inds,
  target_dim,
  value = 0,
  reorient = FALSE,
  ...
)

## S4 method for signature 'ANY'
replaceEmptyImageDimensions(
  img,
  inds,
  target_dim,
  value = 0,
  reorient = FALSE,
  ...
)

replace_empty_dim(img, ...)
```

### Arguments

`img` image, nifti object, or array  
`inds` indices of subset from [getEmptyImageDimensions](#) or [dropEmptyImageDimensions](#).

target_dim	Original dimension from which the data was subset, the final dimension of the output
value	value to replace in the image where outside the indices
reorient	Should image be reoriented if a filename?
...	not used

**Value**

Object of class `nifti` or array if `nifti` is not supplied

**Note**

`replace_empty_dim` is a shorthand for `replaceEmptyImageDimensions` with all the same arguments.

**See Also**

[getEmptyImageDimensions](#), [dropEmptyImageDimensions](#)

**Examples**

```

dims = rep(10, 3)
arr = array(rnorm(prod(dims)), dim = dims)
arr[, , 10] = 0
nim = oro.nifti::nifti(arr)
rnifti = RNifti::asNifti(nim)
ting = tempimg(nim)
limg = list(factor(ting), ting)
inds = getEmptyImageDimensions(nim)
inds_arr = getEmptyImageDimensions(arr)
testthat::expect_equal(inds, inds_arr)

out = applyEmptyImageDimensions(nim, inds = inds)
result = replaceEmptyImageDimensions(out, inds = inds,
target_dim = dim(nim))
testthat::expect_equal(array(result, dim = dim(result)),
array(nim, dim = dim(nim)))
replace_empty_dim(out, inds = inds,
target_dim = dim(nim))

target_dim = dim(nim)

arr = array(out, dim = dim(out))
nim = oro.nifti::nifti(arr)
rnifti = RNifti::asNifti(nim)
ting = tempimg(nim)
limg = list(factor(ting), factor(ting))
func = function(...) replaceEmptyImageDimensions(...,
target_dim = target_dim, inds = inds)
func(arr)
func(nim)

```

```
func(rnifti)
func(timg)
func(limg)
```

---

```
replace_dropped_dimensions
```

*Remake Dropped Dimensions*

---

### Description

This function is the reverse of `dropEmptyImageDimensions`. If `dropEmptyImageDimensions` was run, and the output is a list, usually if `keep_ind = TRUE`, this function reverses that.

### Usage

```
replace_dropped_dimensions(img, inds, orig.dim)
```

### Arguments

<code>img</code>	Object of class <code>nifti</code> where image dimensions were dropped.
<code>inds</code>	List of length 3 of indices from <code>dropEmptyImageDimensions</code> or <code>getEmptyImageDimensions</code>
<code>orig.dim</code>	Original dimension of pre-dropped image. Output image will have dimensions same as this value

### Value

Object of class `nifti`

### Examples

```
## Not run:
# nim is an object of class nifti
dd = dropEmptyImageDimensions(nim, keep_ind = TRUE)
remake = replace_dropped_dimensions(img = dd$outimg,
inds = dd$inds,
orig.dim = dd$orig.dim)
all.equal(nim, remake)

## End(Not run)
```

---

 replace\_outside\_surface

*Replace Values Outside Surface of image*


---

### Description

Determines values outside the surface of an image and gives a mask back with those values set to a replacement.

### Usage

```
replace_outside_surface(
  img,
  value = 0,
  threshold = 0,
  replace_value = NA,
  reorient = FALSE
)
```

### Arguments

img	nifti object or array
value	Value to check against. If zero, then <code>replace_outside_surface</code> will include any dimension that has fewer than <code>threshold</code> zeroes. May be a vector of values, matched with <a href="#">match</a>
threshold	Include dimension if fewer than <code>threshold</code> voxels are in the slice
replace_value	Value to replace those outside the surface.
reorient	Should image be reoriented if a filename? Passed to <a href="#">check_nifti</a>

### Value

Creates an array of binary values. If `img` is a nifti object, then a nifti is returned

### Examples

```
set.seed(5)
dims = rep(10, 3)
arr = array(0, dim = dims)

arr[ 3:5, 4:6, c(2, 6:8, 5)] = 1
nim = oro.nifti::nifti(arr)
out = replace_outside_surface(nim, replace_value = 0)
out_arr = replace_outside_surface(arr, replace_value = 0)
testthat::expect_equal(out_arr, array(out, dim = dim(out)))
```

rescale\_img

*Image Rescaler***Description**

Rescales an image to be in certain value range. This was created as sometimes DICOM scale and slope parameters may be inconsistent across sites and the data need to be value restricted

**Usage**

```
rescale_img(
  filename,
  pngname = NULL,
  write.nii = FALSE,
  outfile = NULL,
  min.val = -1024,
  max.val = 3071,
  ROIformat = FALSE,
  drop_dim = TRUE,
  writer = "dcm2nii",
  ...
)
```

**Arguments**

filename	filename of image to be read into R or nifti object
pngname	filename of png of histogram of values of image to be made. For no png - set to NULL (default)
write.nii	logical - should the image be written.
outfile	if write.nii = TRUE, filename of output file
min.val	minimum value of image (default -1024 (for CT)). If no thresholding set to -Inf
max.val	maximum value of image (default 3071 (for CT)). If no thresholding set to Inf
ROIformat	if TRUE, any values $\leq 0$ will be set to 0
drop_dim	Should <code>drop_img_dim</code> be applied?
writer	character value to add to description slot of NIFTI header
...	extra methods to be passed to <code>writenii</code>

**Value**

Object of class nifti

**Examples**

```
img = nifti(array(rnorm(10^3, sd = 1000), dim = rep(10, 3)))
outfile = tempfile(fileext = ".nii.gz")
pngname = tempfile(fileext = ".png")
rescale_img(img, write.nii = TRUE, outfile = outfile,
pngname = pngname)
```

---

reverse\_orient\_rpi      *Reverse Reorientation an Image to RPI orientation*

---

**Description**

Reverse Reorientation an Image to RPI orientation

**Usage**

```
reverse_orient_rpi(
  file,
  convention = c("NEUROLOGICAL", "RADIOLOGICAL"),
  orientation,
  verbose = TRUE
)

reverse_orient_rpi_file(
  file,
  convention = c("NEUROLOGICAL", "RADIOLOGICAL"),
  orientation,
  verbose = TRUE
)
```

**Arguments**

file	Object of class <code>nifti</code> or character path
convention	Convention of original image (usually from <code>orient_rpi</code> )
orientation	Vector of length 3 from original image (usually from <code>orient_rpi</code> )
verbose	print diagnostic messages

**Value**

Object of class `nifti`

**Note**

'reverse\_orient\_rpi' and 'reverse\_orient\_rpi\_file' uses 'RNifti' to ensure the reading orientation

---

robust_window	<i>Window image based on quantiles of Image</i>
---------------	---

---

**Description**

Takes an image, finds the quantiles given by probs, then sets values outside these values to other values, as determined by replace argument passed to [window\\_img](#)

**Usage**

```
robust_window(img, non_zero = FALSE, probs = c(0, 0.999), ..., mask = NULL)
```

**Arguments**

img	object of class nifti
non_zero	Should zeroes be excluded from the calculation of quantiles?
probs	quantiles to constrain the image these define the window sent to <a href="#">window_img</a>
...	additional arguments sent to <a href="#">window_img</a>
mask	binary image to use to calculate quantiles

**Value**

Object of class nifti with values outside quantiles replaced by values depending on replace argument passed to [window\\_img](#)

---

same_dims	<i>Check if Objects have Same Dimensions</i>
-----------	--

---

**Description**

Wrapper to check if multiple objects all have the same dimensions

**Usage**

```
same_dims(...)
```

**Arguments**

...	Arguments (matrices or arrays) where the dimension will be checked against the first object's dimension
-----	---

**Value**

Logical indicating if all have the same dimensions or not



### Examples

```
mat1 = matrix(1:9, ncol = 3)
mat2 = matrix(rnorm(9), ncol = 3)
mat3 = matrix(rnorm(16), ncol = 4)
mat4 = matrix(rnorm(9), ncol = 3)
same_dims(mat1, mat2)
same_dims(mat1, mat3)
same_dims(mat1, mat2, mat4)
```

---

separate\_img-methods    *Separate Labeled Image into Multiple Binary Images*

---

### Description

Takes in an image, gets the unique values, then creates a list of binary images for each one of those values.

### Usage

```
separate_img(img, levels = NULL, drop_zero = TRUE)

## S4 method for signature 'nifti'
separate_img(img, levels = NULL, drop_zero = TRUE)

## S4 method for signature 'array'
separate_img(img, levels = NULL, drop_zero = TRUE)

## S4 method for signature 'ANY'
separate_img(img, levels = NULL, drop_zero = TRUE)

## S4 method for signature 'factor'
separate_img(img, levels = NULL, drop_zero = TRUE)

## S4 method for signature 'character'
separate_img(img, levels = NULL, drop_zero = TRUE)

## S4 method for signature 'list'
separate_img(img, levels = NULL, drop_zero = TRUE)
```

### Arguments

img	character path of image or an object of class <code>nifti</code> , or list of images
levels	if levels is given, then the separation is only done for those levels and not unique values of the image.
drop_zero	Should zeroes be dropped from the labels? Zero usually denotes background or non-interesting voxels

**Value**

A nifti object (or list of them) or class of object passed in if not specified

**Note**

Exact equalling is using ==

**Examples**

```
set.seed(5)
dims = rep(10, 3)
arr = array(rpois(prod(dims), lambda = 2), dim = dims)
nim = oro.nifti::nifti(arr)
simg = separate_img(nim)
simg_arr = separate_img(arr)
slist = lapply(simg, function(x) array(x, dim(x)))
testthat::expect_equal(slist, simg_arr)

rnifti = RNifti::asNifti(nim)
timg = tempimg(nim)
limg = list(factor(timg), factor(timg))
func = separate_img
func(arr)
func(nim)
func(rnifti)
func(timg)
func(limg)
```

---

slice\_colour\_df

*Slice a Image Color Data.frame*

---

**Description**

Slices a image color data.frame along the 3 planes (axial, coronal, sagittal) and returns it in a ggplot-ready format for faceting.

**Usage**

```
slice_colour_df(img_df, xyz = NULL)
```

**Arguments**

img_df	an image data.frame, usually from <code>img_colour_df</code> . Must have the columns: dim1, dim2, dim3, colour, and value.
xyz	coordinates to slice the data.frame in x, y, and z - domains

**Value**

A data.frame with x and y coordinates, colour, and intensity values, along with the associated planes that were sliced.

**Examples**

```
img = nifti(array(rnorm(10^3), dim = rep(10, 3)))
df = img_colour_df(img)
sliced = slice_colour_df(df, c(5, 5, 4))
```

---

subset\_dti-methods      *Subset DTI data based on b-values #'*

---

**Description**

Subset DTI data based on b-values #'

**Usage**

```
subset_dti(
  img,
  bvals,
  bvecs,
  b_step = 1,
  maximum = Inf,
  shells = NULL,
  verbose = TRUE,
  ...
)

## S4 method for signature 'nifti'
subset_dti(
  img,
  bvals,
  bvecs,
  b_step = 1,
  maximum = Inf,
  shells = NULL,
  verbose = TRUE,
  ...
)

## S4 method for signature 'ANY'
subset_dti(
  img,
  bvals,
  bvecs,
```

```

    b_step = 1,
    maximum = Inf,
    shells = NULL,
    verbose = TRUE,
    ...
)

## S4 method for signature 'character'
subset_dti(
  img,
  bvals,
  bvecs,
  b_step = 1,
  maximum = Inf,
  shells = NULL,
  verbose = TRUE,
  ...
)

## S4 method for signature 'list'
subset_dti(
  img,
  bvals,
  bvecs,
  b_step = 1,
  maximum = Inf,
  shells = NULL,
  verbose = TRUE,
  ...
)

```

### Arguments

<code>img</code>	character or nifti object
<code>bvals</code>	filename of b-values (assuming 1 row)
<code>bvecs</code>	filename of b-vectors (assuming 3 rows)
<code>b_step</code>	step of b-values to round to
<code>maximum</code>	maximum b-value threshold
<code>shells</code>	Shells to keep (after rounding)
<code>verbose</code>	print diagnostic messages
<code>...</code>	options passed to <a href="#">checking</a>

### Value

List of filenames of image, b-values, and b-vectors that were subsetted.

**Author(s)**

John Muschelli <muschellij2@gmail.com>

**Examples**

```
## Not run:
img = "~/Downloads/data.nii.gz"
bvals = "~/Downloads/bvals"
bvecs = "~/Downloads/bvals"
verbose = TRUE
b_step = 50
maximum = 1500
shells = NULL
sub = subset_dti(img = img, bvals = bvals, bvecs = bvecs,
maximum = 1500,
b_step = 50)

## End(Not run)
```

---

tempimg

---

*Create temporary nii.gz file*


---

**Description**

Takes in a object of class nifti, writes it to a temp file, appends .nii.gz as [writenii](#) adds it.

**Usage**

```
tempimg(
  nim,
  gzipped = TRUE,
  checknan = TRUE,
  check_type = FALSE,
  warn = FALSE,
  drop_dim = TRUE,
  dtype = TRUE,
  ...
)
```

**Arguments**

nim	object of class nifti
gzipped	Should file be gzipped? Passed to <a href="#">writenii</a>
checknan	Check for NAs or NaNs
check_type	Check the datatype for an image. Will run <a href="#">datatyper</a> .
warn	Should warnings be displayed if <a href="#">writenii</a> has any? Passed to <a href="#">writenii</a> .

drop_dim	Should <a href="#">drop_img_dim</a> be applied?
dtype	Should <a href="#">datatyper</a> be run before writing? Should override 'check_type'
...	Passed to <a href="#">writenii</a> .

**Value**

filename of output nii.gz

---

window_img	<i>nifti image windower</i>
------------	-----------------------------

---

**Description**

Windows an image to min and max values and also changes cal\_max and cal\_min parameters

**Usage**

```

window_img(
  x,
  window = c(0, 100),
  replace = c("window", "missing", "zero"),
  ...
)

```

**Arguments**

x	is either a character name for the image or an object of class nifti
window	numeric of length 2 that gives min and max for window
replace	either "window" if the any values outside of c(min, max) are set to the min or max or "missing" for these to be set to NA
...	not used

**Value**

Object of class nifti

**See Also**

[readnii](#)

**Examples**

```

set.seed(5)
dims = rep(10, 3)
arr = array(rnorm(prod(dims)), dim = dims)
nim = oro.nifti::nifti(arr)
window_img(nim, window = c(1, 5))
window_img(nim, window = c(1, 5), replace = "missing")
tfile = tempimg(nim)
window_img(tfile)
window_img(as.factor(tfile))
arr = window_img(img_data(nim))
rnim = RNifti::readNifti(tfile)
window_img(rnim, window = c(1, 5))
range(window_img(rnim, window = c(1, 5)))
window_img(rnim, window = c(1, 5), replace = "missing")
range(window_img(rnim, window = c(1, 5), replace = "missing"))

```

---

writeNIFTI2

*writeNIFTI with default non-reorientation*


---

**Description**

This function calls the [writeNIFTI](#) function from the `oro.nifti` package, but makes sure to remove `.nii` extension and warnings can be suppressed.

**Usage**

```
writeNIFTI2(nim, filename, dtype = FALSE, compression = 9, ...)
```

```

writenii(
  nim,
  filename,
  dtype = TRUE,
  drop_dim = TRUE,
  warn = FALSE,
  compression = 9,
  rm_extensions = TRUE,
  ...
)

```

**Arguments**

<code>nim</code>	object of class <code>nifti</code> , passed to <a href="#">writeNIFTI</a>
<code>filename</code>	path to save the NIFTI file. Suffix will be removed
<code>dtype</code>	Should <a href="#">datatyper</a> be run before writing?

compression	compression level for gzipped files.
...	Additional arguments passed to <code>writeNIFTI</code>
drop_dim	Should <code>drop_img_dim</code> be run before writing?
warn	Should warnings from <code>writeNIFTI</code> be printed? If not, <code>suppressWarnings</code> is called
rm_extensions	should <code>niftiExtensions</code> be converted to simple nifti objects before writing?

**Value**

Nothing

**Note**

While `writeNIFTI2` does not run `datatyper` as default, `writenii` does. Additional functionality will be added to `writenii` likely but will not to `writeNIFTI2`

**Examples**

```
set.seed(5)
dims = rep(10, 3)
arr = array(rnorm(prod(dims)), dim = dims)
nim = oro.nifti::nifti(arr)
rnifti = RNifti::asNifti(nim)
tfile = tempfile(fileext = ".nii.gz")
timg = writenii(nim, tfile, rm_extensions = TRUE, warn = TRUE)
timg = writeNIFTI2(nim, tfile, dtype = TRUE)
```

---

write_nifti	<i>General NIFTI Writer</i>
-------------	-----------------------------

---

**Description**

Writes out NIFTI files for multiple formats. Currently, for `nifti` objects and `niftiImage` objects from `RNifti`

**Usage**

```
write_nifti(nim, filename, ...)
```

**Arguments**

nim	Container for NIFTI Image
filename	Filename of image to be written out
...	additional arguments, to be passed to <code>writeNifti</code> or <code>writenii</code>

**Value**

Output from NIFTI writer



**Examples**

```

set.seed(5)
dims = rep(10, 4)
arr = array(rpois(prod(dims), lambda = 2), dim = dims)
nim = oro.nifti::nifti(arr)
tfile = tempfile(fileext = ".nii.gz")
write_nifti(nim, tfile)
ring = RNifti::readNifti(tfile)
write_nifti(ring, tfile)

```

---

xyz

*Image Center of Gravity Wrapper*


---

**Description**

Find Center of Gravity of Image, after thresholding and take ceiling (wrapper for [cog](#))

**Usage**

```
xyz(...)
```

**Arguments**

... Arguments passed to [cog](#)

**Value**

Vector of length 3

**Note**

Just a convenience wrapper for `cog(ceil=TRUE)`

---

zero\_pad

*Zero pads an image*


---

**Description**

This function zero pads an image by a certain number of dimensions, usually for convolution

**Usage**

```
zero_pad(img, kdim, invert = FALSE, pad_value = 0L, ...)
```

**Arguments**

img	Array or class nifti
kdim	Dimensions of kernel
invert	(logical) If FALSE, does zero padding. If TRUE, reverses the process.
pad_value	Value to pad the image with. May use other values, such as -1024 for CT data
...	Options to <a href="#">copyNIFTIHeader</a>

**Value**

Object of class nifti

**Examples**

```
kdim = c(3,3,5)
img = array(rnorm(30*30*36), dim = c(30, 30, 36))
pad = zero_pad(img, kdim)
back = zero_pad(pad, kdim, invert=TRUE)
all.equal(back, img)
```

---

zlimmer

*Find Image z-limits*


---

**Description**

Helper function for plotting - returns zlim for [image](#) plot function

**Usage**

```
zlimmer(x, zlim = NULL, computed_range = NULL)
```

**Arguments**

x	Object of class nifti
zlim	A user-specified zlim. If NULL, will calculate how <a href="#">ortho2</a> would calculate zlim
computed_range	If the range of the data was already computed, this can be passed in and will be used if relevant.

**Value**

If zlim = NULL, then vector of length 2, otherwise returns zlim

---

zscore\_img

*Get Z-score over a margin of an img*


---

**Description**

Standardizes an image either by the axial, sagittal, or coronal slice or whole image

**Usage**

```
zscore_img(
  img,
  mask = NULL,
  margin = NULL,
  centrality = c("mean", "median", "trimmed_mean"),
  variability = c("sd", "iqrdiff", "mad", "maddiff", "iqr", "trimmed_sd"),
  trim = 0.2,
  remove.na = TRUE,
  remove.nan = TRUE,
  remove.inf = TRUE,
  remove.val = 0,
  remask = TRUE
)
```

**Arguments**

img	character path of image or an object of class nifti
mask	character path of mask or an object of class nifti
margin	Margin of image to z-score over (NULL - whole brain, 3-Axial, 2-Sagittal, 1-Coronal)
centrality	(character) Measure to center the data, either mean or median
variability	(character) Measure to scale the data
trim	if centrality is trimmed_mean or variability is trimmed_sd, then the amount of trimming
remove.na	(logical) change NAs to remove.val
remove.nan	(logical) change NaN to remove.val
remove.inf	(logical) change Inf to remove.val
remove.val	(logical) value to put the NA/NaN/Inf
remask	(logical) Should the image be remasked after normalizing?

**Value**

Array of object of class nifti

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

```

dim = c(100, 30, 5)
img = array(rnorm(prod(dim), mean=4, sd=4),
dim=dim)

truth2 = img
for (i in 1:dim(img)[2]) {
truth2[,i,] = (truth2[,i,] - mean(truth2[,i,]))/sd(truth2[,i,])
}

truth1 = img
for (i in 1:dim(img)[1]) {
truth1[i,,] = (truth1[i,,] - mean(truth1[i,,]))/sd(truth1[i,,])
}

truth3 = img
for (i in 1:dim(img)[3]) {
truth3[,,i] = (truth3[,,i] - mean(truth3[,,i]))/sd(truth3[,,i])
}
try3 = zscore_img(img, margin=3)
stopifnot(all.equal(try3, truth3))
try2 = zscore_img(img, margin=2)
stopifnot(all.equal(try2, truth2))
try1 = zscore_img(img, margin=1)
stopifnot(all.equal(try1, truth1))

z = zscore_img(img, margin=NULL)
ztrim = zscore_img(img, margin=NULL,
centrality = "trimmed_mean", variability = "trimmed_sd")

z = zscore_img(img, centrality = "median", variability = "iqr")
z = zscore_img(img, centrality = "median", variability = "iqrdiff")
z = zscore_img(img, centrality = "median", variability = "maddiff")

```

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