

# Package ‘gifti’

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**Type** Package

**Title** Reads in 'Neuroimaging' 'GIFTI' Files with Geometry Information

**Version** 0.9.0

**Description** Functions to read in the geometry format under the 'Neuroimaging' 'Informatics' Technology Initiative (NIfTI), called 'GIFTI' <<https://www.nitrc.org/projects/gifti/>>. These files contain surfaces of brain imaging data.

**License** GPL-2

**Imports** xml2 (>= 1.1.1), base64enc, R.utils, tools, utils

**Suggests** rgl, grDevices, testthat, knitr, rmarkdown, covr

**BugReports** <https://github.com/muschellij2/gifti/issues>

**Encoding** UTF-8

**VignetteBuilder** knitr

**RoxygenNote** 7.3.3

**NeedsCompilation** no

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**Repository** CRAN

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convert\_binary\_datatype

*Convert Binary Data Type*

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## Description

Converts a data type to the size and what for [readBin](#), necessary for Base64Binary and GZipBase64Binary formats

## Usage

```
convert_binary_datatype(
  datatype = c("NIFTI_TYPE_UINT8", "NIFTI_TYPE_INT32", "NIFTI_TYPE_UINT32",
              "NIFTI_TYPE_FLOAT32")
)
```

## Arguments

datatype            data type from GIFTI image

## Value

List of length 2: with elements of size and what

## Examples

```
convert_binary_datatype()
convert_binary_datatype('NIFTI_TYPE_INT32')
testthat::expect_error(convert_binary_datatype('NIFTI_TYPE_BLAH'))
```

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convert_endian	<i>Convert Endian from GIFTI</i>
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**Description**

Converts Endian format from GIFTI

**Usage**

```
convert_endian(endian)
```

**Arguments**

endian	character passed from GIFTI XML
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**Value**

Character string

---

convert_intent	<i>Convert Intent</i>
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**Description**

Converts the intent field from a GIFTI image to a more standard naming

**Usage**

```
convert_intent(intent)
```

**Arguments**

intent	(character) string of intent type
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**Value**

A character string

`create_data_matrix`     *Create Data Matrix with ordering respected*

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**Description**

Create Data Matrix with ordering respected

**Usage**

```
create_data_matrix(  
  data,  
  dims,  
  ordering = c("RowMajorOrder", "ColumnMajorOrder")  
)
```

**Arguments**

<code>data</code>	Data output from <a href="#">data_decoder</a>
<code>dims</code>	Dimensions of output
<code>ordering</code>	Ordering of the data

**Value**

Matrix of Values

---

`data_array_attributes`     *Data Array Attributes*

---

**Description**

Parses a list of XML data to get the attributes

**Usage**

```
data_array_attributes(darray)
```

**Arguments**

<code>darray</code>	List of <code>xml_nodes</code> from GIFTI data array
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**Value**

`data.frame` of attributes

---

data_decoder	<i>Array Data Decoder</i>
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**Description**

Decodes values from a GIFTI image

**Usage**

```
data_decoder(
  values,
  encoding = c("ASCII", "Base64Binary", "GZipBase64Binary", "ExternalFileBinary"),
  datatype = NULL,
  endian = c("little", "big", "LittleEndian", "BigEndian"),
  ext_filename = NULL,
  n = NULL
)
```

**Arguments**

values	text from XML of GIFTI image
encoding	encoding of GIFTI values
datatype	Passed to <a href="#">convert_binary_datatype</a>
endian	Endian to pass in <a href="#">readBin</a>
ext_filename	if encoding = "ExternalFileBinary", then this is the external filename
n	number of values to read. Relevant if encoding = "ExternalFileBinary"

**Value**

Vector of values

**Examples**

```
if (have_gifti_test_data(outdir = NULL)) {
  gii_files = download_gifti_data(outdir = NULL)
  L = gifti_list(gii_files[1])
  orig = L$DataArray$Data[[1]]
  encoding = attributes(L$DataArray)$Encoding
  datatype = attributes(L$DataArray)$DataType
  endian = attributes(L$DataArray)$Endian
  vals = data_decoder(orig, encoding = encoding,
    datatype = datatype, endian = endian)
  enc = data_encoder(vals, encoding = encoding,
    datatype = datatype, endian = endian)
  enc == orig
}
```

---

 data\_encoder

*Array Data Encoder*


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### Description

Encodes values for a GIFTI image

### Usage

```
data_encoder(
  values,
  encoding = c("ASCII", "Base64Binary", "GZipBase64Binary"),
  datatype = NULL,
  endian = c("little", "big", "LittleEndian", "BigEndian")
)
```

### Arguments

values	values to be encoded
encoding	encoding of GIFTI values
datatype	Passed to <a href="#">convert_binary_datatype</a>
endian	Endian to pass in <a href="#">readBin</a>

### Value

Single character vector

### Examples

```
if (have_gifti_test_data(outdir = NULL)) {
  gii_files = download_gifti_data(outdir = NULL)
  L = gifti_list(gii_files[1])
  orig = L$DataArray$Data[[1]]
  encoding = attributes(L$DataArray)$Encoding
  datatype = attributes(L$DataArray)$DataType
  endian = attributes(L$DataArray)$Endian
  vals = data_decoder(orig, encoding = encoding,
    datatype = datatype, endian = endian)
  enc = data_encoder(vals, encoding = encoding,
    datatype = datatype, endian = endian)
  enc == orig
}
```

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decompress_gii	<i>Decompress Gzipped GIFTI (with extension .gz)</i>
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**Description**

If a GIFTI file is compressed, as in .gii.gz, this will decompress the file. This has nothing to do with the encoding WITHIN the file

**Usage**

```
decompress_gii(file)
```

**Arguments**

file	file name of GIFTI file
------	-------------------------

**Value**

Filename of decompressed GIFTI

**Examples**

```
if (have_gifti_test_data(outdir = NULL)) {  
  gii_files = download_gifti_data(outdir = NULL)  
  outfile = decompress_gii(gii_files[1])  
  print(outfile)  
}
```

---

download_gifti_data	<i>Download GIFTI Test Data</i>
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**Description**

Downloads GIFTI test data from [https://www.nitrc.org/frs/download.php/411/BV\\_GIFTI\\_1.3.tar.gz](https://www.nitrc.org/frs/download.php/411/BV_GIFTI_1.3.tar.gz)

**Usage**

```
download_gifti_data(  
  outdir = system.file(package = "gifti"),  
  overwrite = FALSE,  
  ...  
)
```

**Arguments**

outdir            Output directory for test file directory  
 overwrite        Should files be overwritten if already exist?  
 ...              additional arguments to [download.file](#)

**Value**

Vector of file names

---

gifti_list	<i>Convert GIFTI to List</i>
------------	------------------------------

---

**Description**

Reads in a GIFTI file and coerces it to a list

**Usage**

```
gifti_list(file)
```

**Arguments**

file            file name of GIFTI file

**Value**

List of elements

**Examples**

```
if (have_gifti_test_data(outdir = NULL)) {
  gii_files = download_gifti_data(outdir = NULL)
  L = gifti_list(gii_files[1])
  orig = L$DataArray$Data[[1]]
  encoding = attributes(L$DataArray)$Encoding
  datatype = attributes(L$DataArray)$DataType
  endian = attributes(L$DataArray)$Endian
  vals = data_decoder(orig, encoding = encoding,
    datatype = datatype, endian = endian)
  enc = data_encoder(vals, encoding = encoding,
    datatype = datatype, endian = endian)
  enc == orig
}
```

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gifti_map_value	<i>Map Values to Triangles from GIFTI</i>
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**Description**

Takes values and maps them to the correct triangles in space.

**Usage**

```
gifti_map_value(
  pointset,
  triangle,
  values,
  indices = seq(nrow(pointset)),
  add_one = TRUE
)
```

**Arguments**

pointset	pointset from GIFTI
triangle	triangles from GIFTI
values	Values to map to the triangles. Same length as indices
indices	indices to place the values, must be in the range of 1 and the number of rows of pointset
add_one	Should 1 be added to the indices for the triangle?

**Value**

A list of coordinates (in triangles) and the corresponding value mapped to those triangles

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have_gifti_test_data	<i>Check Presence of GIFTI Test Data</i>
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---

**Description**

Checks if GIFTI test data is downloaded

**Usage**

```
have_gifti_test_data(outdir = system.file(package = "gifti"))
```

**Arguments**

outdir	Output directory for test file directory
--------	--

**Value**

Logical indicator

**Examples**

```
have_gifti_test_data(outdir = NULL)
```

---

<code>is.gifti</code>	<i>Test if GIFTI</i>
-----------------------	----------------------

---

**Description**

Simple wrapper to determine if class is GIFTI

**Usage**

```
is.gifti(x)
```

```
is_gifti(x)
```

**Arguments**

x                    object to test

**Value**

Logical if x is GIFTI

---

<code>readgii</code>	<i>Read GIFTI File</i>
----------------------	------------------------

---

**Description**

Reads a GIFTI File and parses the output

**Usage**

```
readgii(file)
```

```
readGIFTI(file)
```

```
read_gifti(file)
```

**Arguments**

file                Name of file to read

**Value**

List of values

**Examples**

```

if (have_gifti_test_data(outdir = NULL)) {
  gii_files = download_gifti_data(outdir = NULL)
  gii_list = lapply(gii_files, readgii)
  surf_files = grep("white[.]surf[.]gii", gii_files, value = TRUE)
  surfs = lapply(surf_files, surf_triangles)

  col_file = grep("white[.]shape[.]gii", gii_files, value = TRUE)
  cdata = readgii(col_file)
  cdata = cdata$data$shape
  mypal = grDevices::colorRampPalette(colors = c("blue", "black", "red"))
  n = 4
  breaks = quantile(cdata)
  ints = cut(cdata, include.lowest = TRUE, breaks = breaks)
  ints = as.integer(ints)
  stopifnot(!any(is.na(ints)))
  cols = mypal(n)[ints]
  cols = cols[surfs[[1]]$triangle]
}
## Not run:
if (have_gifti_test_data(outdir = NULL)) {

  if (requireNamespace("rgl", quietly = TRUE)) {
    rgl::open3d()
    rgl::triangles3d(surfs[[1]]$pointset, color = cols)
    rgl::play3d(rgl::spin3d(), duration = 5)
  }
}

## End(Not run)

```

---

surf\_triangles

*Make Triangles from Gifti Image*

---

**Description**

Creates Triangles for plotting in RGL from a Gifti image

**Usage**

```
surf_triangles(file)
```

**Arguments**

file                    File name of Gifti image, usually surf.gii

**Value**

List of values corresponding to the data element from [readgii](#)

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writegii	<i>Write .gii xml from "gifti" object</i>
----------	---

---

**Description**

Writes a "gifti" object to a GIFTI file (ends in \*.gii).

**Usage**

```
writegii(gii, out_file, use_parsed_transformations = FALSE)
```

```
writeGIFTI(gii, out_file, use_parsed_transformations = FALSE)
```

```
write_gifti(gii, out_file, use_parsed_transformations = FALSE)
```

**Arguments**

gii                   The "gifti" object

out\_file             Where to write the new GIFTI file

use\_parsed\_transformations

Should the \$parsed\_transformations be written instead of the transformations?  
Use if the XML pointers in transformations might be invalid. Default: FALSE

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