

# Package ‘BHMSMAfMRI’

May 16, 2019

**Version** 1.3

**Date** 2019-05-15

**Title** Bayesian Hierarchical Multi-Subject Multiscale Analysis of Functional MRI Data

**Author** Nilotpal Sanyal [aut, cre],  
Marco A.R. Ferreira [aut]

**Maintainer** Nilotpal Sanyal <nsanyal@stanford.edu>

**Depends** oro.nifti, wavethresh, compiler

**Suggests** knitr

**VignetteBuilder** knitr

**Description** Performs Bayesian hierarchical multi-subject multiscale analysis of fMRI data as described in Sanyal & Ferreira (2012) <DOI:10.1016/j.neuroimage.2012.08.041> using wavelet based prior that borrows strength across subjects and returns posterior smoothed versions of the fMRI data and samples from the posterior distribution.

**License** GPL (>= 2)

**URL** <https://www.r-project.org>

**Repository** CRAN

**NeedsCompilation** no

**Date/Publication** 2019-05-16 07:50:03 UTC

## R topics documented:

|                              |    |
|------------------------------|----|
| BHMSMAfMRI-package . . . . . | 2  |
| BHMSMA . . . . .             | 2  |
| fmridata . . . . .           | 4  |
| glmcoeff . . . . .           | 5  |
| hyperparamest . . . . .      | 6  |
| pikljbar . . . . .           | 7  |
| postglmcoeff . . . . .       | 9  |
| postgroupcoeff . . . . .     | 10 |
| postsamples . . . . .        | 11 |

|                            |    |
|----------------------------|----|
| postwaveletcoeff . . . . . | 13 |
| read.fmridata . . . . .    | 14 |
| waveletcoeff . . . . .     | 16 |

|              |           |
|--------------|-----------|
| <b>Index</b> | <b>18</b> |
|--------------|-----------|

---

|                    |   |
|--------------------|---|
| BHMSMAfMRI-package | <i>Bayesian Hierarchical Multi-Subject Multiscale Analysis of Functional MRI Data</i> |
|--------------------|---|

---

### Description

Performs BHMSMA (Sanyal & Ferreira, 2012) of fMRI data using wavelet based prior that borrows strength across subjects and returns posterior smoothed versions of the fMRI data and samples from the posterior distribution. Currently considers 2D slices only.

### Details

Package: BHMSMAfMRI  
 Type: Package  
 Version: 1.3  
 Date: 2019-05-15  
 License: GPL (>= 2)

### Author(s)

Nilotpall Sanyal <nsanyal@stanford.edu>, Marco Ferreira <marf@vt.edu>

### References

Sanyal, Nilotpall, and Ferreira, Marco A.R. (2012). Bayesian hierarchical multi-subject multiscale analysis of functional MRI data. *Neuroimage*, 63, 3, 1519-1531.

---

|        |   |
|--------|---|
| BHMSMA | <i>Bayesian hierarchical multi-subject multiscale analysis of functional MRI data</i> |
|--------|---|

---

### Description

Performs BHMSMA (Sanyal & Ferreira, 2012) of fMRI data using wavelet based prior that borrows strength across subjects and returns posterior smoothed versions of the fMRI data

**Usage**

```
BHMSMA(nsubject, grid, Data, DesignMatrix, TrueCoeff=NULL, analysis,
       wave.family="DaubLeAsymm", filter.number=6, bc="periodic")
```

**Arguments**

|               |  |
|---------------|--|
| nsubject      | Number of subjects included in the analysis.   |
| grid          | The number of voxels in one row (or, one column) of the brain slice of interest. Must be a power of 2. The total number of voxels is $\text{grid}^2$ . The maximum grid value for this package is 512. |
| Data          | The data in form of an array with dimension (nsubject,grid,grid,ntime), where ntime is the size of the time series for each voxel.   |
| DesignMatrix  | The design matrix used to generate the data.   |
| TrueCoeff     | If available, the true GLM coefficients in form of an array with dimension (nsubject,grid,grid). By default, NULL.   |
| analysis      | "MSA" or "SSA", depending on whether performing multi-subject analysis or single subject analysis.   |
| wave.family   | The family of wavelets to use - "DaubExPhase" or "DaubLeAsymm". Default is "DaubLeAsymm".  |
| filter.number | The number of vanishing moments of the wavelet. By default 6.  |
| bc            | The boundary condition to use - "periodic" or "symmetric". Default is "periodic".  |

**Details**

The wavelet computations are performed by using R package 'wavethresh'. For details, check wavethresh package help.

**Value**

A list containing the following.

GLMCoeffStandardized

An array of dimension (nsubject, grid, grid), containing for each subject the standardized GLM coefficients obtained by fitting GLM to the time-series corresponding to the voxels.

GLMEstimatedSE

An array of dimension (nsubject, grid, grid), containing for each subject the estimated standard errors of the GLM coefficients.

WaveletCoefficientMatrix

A matrix of dimension (nsubject,  $\text{grid}^2-1$ ), containing for each subject the wavelet coefficients of all levels stacked together (by the increasing order of resolution level).

hyperparam

A vector containing the estimates of the six hyperparameters.

hyperparamVar

Estimated covariance matrix of the hyperparameters.

|                      |   |
|----------------------|---|
| pk1j.bar             | A matrix of dimension (nsubject, grid <sup>2</sup> -1), containing the piklj bar values (see Reference for details).  |
| PostMeanWaveletCoeff | A matrix of size (nsubject, grid <sup>2</sup> -1), containing for each subject the posterior mean of the wavelet coefficients of all levels stacked together (by the increasing order of resolution level). |
| GLMcoeffposterior    | An array of dimension (nsubject, grid, grid), containing for each subject the posterior means of the standardized GLM coefficients.   |
| MSE                  | MSE of the posterior estimates of the GLM coefficients, if the true values of the GLM coefficients are available.   |

**Author(s)**

Nilotpal Sanyal <nsanyal@stanford.edu>, Marco Ferreira <marf@vt.edu>

**References**

Sanyal, Nilotpal, and Ferreira, Marco A.R. (2012). Bayesian hierarchical multi-subject multiscale analysis of functional MRI data. *Neuroimage*, 63, 3, 1519-1531.

**Examples**

```
# Should take less than a minute to run
nsubject <- 3
grid <- 8
ntime <- 4
Data <- array(rnorm(3*8*8*4),dim=c(3,8,8,4))
DesignMatrix <- cbind(c(1,0,1,0), c(1,1,1,1))
analysis <- "multi"
BHMSMA.multi <- BHMSMA(nsubject, grid, Data, DesignMatrix, TrueCoeff=NULL, analysis)
```

---

fmridata

*A simulated fMRI data for 3 subjects*

---

**Description**

A simulated fMRI data containing true regression coefficients images for three subjects and design matrix

**Usage**

```
data(fmridata)
```

**Format**

A list containing the following.

- `grid = 32`. The image dimension is 32 by 32.
- `nsubject = 3`.
- `TrueCoeff` An array of dimension (3,32,32), containing the true regression coefficients for the 3 subjects.
- `DesignMatrix` A matrix with 9 columns and 2 rows. The first column is a column of ones.

**Details**

This dataset contains only the true coefficients. The noisy fMRI data, which are generated by adding Gaussian random noise to these true coefficients, are included in the `extdata` directory within the package directory. The function `read_fmridata` can be used to read those data files. The true coefficients and the noisy data both are generated using the R package `neuRosim`. The following specifications were used to generate the data: `totaltime=18`, `onsets=seq(1,18,by=8)`, `durations=1`, `TR=2`, `effectsize=1`, `hrf="double-gamma"`, `regions=3`, `radius=c(1,1,1)`, `form="sphere"`, `fading=1`, `SNR=1.5`, `noise="white"`. The centers of the activation regions were chosen manually. For information regarding the specifications, see `neuRosim` help.

---

|                       |  |
|-----------------------|--|
| <code>glmcoeff</code> | <i>Fit GLM to the data time-series and obtain GLM coefficients along with standard error estimates</i> |
|-----------------------|--|

---

**Description**

Fits General Linear Model to the time-series corresponding to each voxel in the data and returns the standardized GLM coefficients and their standard error estimates.

**Usage**

```
glmcoeff(nsubject, grid, Data, DesignMatrix)
```

**Arguments**

|                           |  |
|---------------------------|--|
| <code>nsubject</code>     | Number of subjects included in the analysis.   |
| <code>grid</code>         | The number of voxels in one row (or, one column) of the brain slice of interest. Must be a power of 2. The total number of voxels is <code>grid^2</code> . The maximum grid value for this package is 512. |
| <code>Data</code>         | The data in form of an array with dimension (nsubject,grid,grid,ntime), where <code>ntime</code> is the size of the time series for each voxel.  |
| <code>DesignMatrix</code> | The design matrix used to generate the data.   |

**Value**

A list containing the following.

**GLMCoeffStandardized**

An array of dimension (nsubject, grid, grid), containing for each subject the standardized GLM coefficients obtained by fitting GLM to the time-series corresponding to the voxels.

**GLMEstimatedSE**

An array of dimension (nsubject, grid, grid), containing for each subject the estimated standard errors of the GLM coefficients.

**Author(s)**

Nilotpal Sanyal <nsanyal@stanford.edu>, Marco Ferreira <marf@vt.edu>

**References**

Friston, K.J., Holmes, A.P., Worsley, K.J., Poline, J., Frith, C.D., Frackowiak, R.S.J., 1994. Statistical parametric maps in functional imaging: a general linear approach. *Hum. Brain Mapp.* 2 (4), 189-210.

**Examples**

```
nsubject <- 3
grid <- 8
Data <- array(dim=c(3,8,8,10),rnorm(3*8*8*10))
DesignMatrix <- cbind( c(rep(c(1,0),5)), rep(1,10) )
glm.fit <- glmcoeff(nsubject, grid, Data, DesignMatrix)
dim(glm.fit$GLMCoeffStandardized)
#[1] 3 8 8
```

---

hyperparamest

*Get the estimates of the hyperparameters of the BHMSME model along with the estimate of their covariance matrix.*

---

**Description**

Computes the MLEs of the hyperparameters of the BHMSME model following an empirical Bayes approach and the estimate of the covariance matrix of the hyperparameters.

**Usage**

```
hyperparamest(nsubject, grid, WaveletCoefficientMatrix, analysis)
```

**Arguments**

|                          |  |
|--------------------------|--|
| nsubject                 | Number of subjects included in the analysis.   |
| grid                     | The number of voxels in one row (or, one column) of the brain slice of interest. Must be a power of 2. The total number of voxels is grid <sup>2</sup> . The maximum grid value for this package is 512. |
| WaveletCoefficientMatrix | A matrix of dimension (nsubject, grid <sup>2</sup> -1), containing for each subject the wavelet coefficients of all levels stacked together (by the increasing order of resolution level).               |
| analysis                 | "multi" or "single", depending on whether performing multi-subject analysis or single subject analysis.  |

**Value**

A list containing the following.

|               |   |
|---------------|---|
| hyperparam    | A vector containing the estimates of the six hyperparameters of the BHMSME model. |
| hyperparamVar | Estimated covariance matrix of the hyperparameters.                               |

**Author(s)**

Nilotpai Sanyal <nsanyal@stanford.edu>, Marco Ferreira <marf@vt.edu>

**References**

Sanyal, Nilotpai, and Ferreira, Marco A.R. (2012). Bayesian hierarchical multi-subject multiscale analysis of functional MRI data. *Neuroimage*, 63, 3, 1519-1531.

**Examples**

```
nsubject <- 3
grid <- 8
WaveletCoefficientMatrix <- array(dim=c(3,63),rnorm(3*63))
analysis <- "multi"
hyper.est <- hyperparamest(nsubject, grid, WaveletCoefficientMatrix, analysis)
```

---

pikljbar

*Compute the piklj bar values of the BHMSMA model using Newton Cotes algorithm*

---

**Description**

Computes the values of piklj bar of the BHMSMA model using Newton Cotes algorithm. For details, check References.

**Usage**

```
pikljbar(nsubject, grid, WaveletCoefficientMatrix, hyperparam, analysis)
```

**Arguments**

|                          |  |
|--------------------------|--|
| nsubject                 | Number of subjects included in the analysis.   |
| grid                     | The number of voxels in one row (or, one column) of the brain slice of interest. Must be a power of 2. The total number of voxels is $\text{grid}^2$ . The maximum grid value for this package is 512. |
| WaveletCoefficientMatrix | A matrix of dimension $(\text{nsubject}, \text{grid}^2-1)$ , containing for each subject the wavelet coefficients of all levels stacked together (by the increasing order of resolution level).        |
| hyperparam               | A vector containing the estimates of the six hyperparameters.  |
| analysis                 | "MSA" or "SSA", depending on whether performing multi-subject analysis or single subject analysis.   |

**Value**

A list containing the following.

|          |   |
|----------|---|
| pklj.bar | A matrix of dimension $(\text{nsubject}, \text{grid}^2-1)$ , containing the piklj bar values. |
|----------|---|

**Author(s)**

Nilotpal Sanyal <nsanyal@stanford.edu>, Marco Ferreira <marf@vt.edu>

**References**

Sanyal, Nilotpal, and Ferreira, Marco A.R. (2012). Bayesian hierarchical multi-subject multiscale analysis of functional MRI data. *Neuroimage*, 63, 3, 1519-1531.

**Examples**

```
nsubject <- 3
grid <- 8
WaveletCoefficientMatrix <- matrix(nrow=3,ncol=63)
for(i in 1:3)
  WaveletCoefficientMatrix[i,] <- rnorm(63)
hyperparam <- rep(.1,6)
analysis <- "multi"
piklj.bar <- pikljbar(nsubject, grid, WaveletCoefficientMatrix, hyperparam, analysis)
dim(piklj.bar$pklj.bar)
#[1] 3 63
```



---

|              |  |
|--------------|--|
| postglmcoeff | <i>Obtain the posterior mean of the GLM coefficients using the posterior mean of the wavelet coefficients.</i> |
|--------------|--|

---

### Description

Computes the posterior mean of the GLM coefficients using the posterior mean of the wavelet coefficients by means of inverse discrete wavelet transform.

### Usage

```
postglmcoeff(nsubject, grid, GLMCoeffStandardized, PostMeanWaveletCoeff,
  wave.family="DaubLeAsymm", filter.number=6, bc="periodic")
```

### Arguments

|                      |   |
|----------------------|---|
| nsubject             | Number of subjects included in the analysis.  |
| grid                 | The number of voxels in one row (or, one column) of the brain slice of interest. Must be a power of 2. The total number of voxels is grid <sup>2</sup> . The maximum grid value for this package is 512.    |
| GLMCoeffStandardized | An array of dimension (nsubject, grid, grid), containing for each subject the standardized GLM coefficients obtained by fitting GLM to the time-series corresponding to the voxels.                         |
| PostMeanWaveletCoeff | A matrix of size (nsubject, grid <sup>2</sup> -1), containing for each subject the posterior mean of the wavelet coefficients of all levels stacked together (by the increasing order of resolution level). |
| wave.family          | The family of wavelets to use - "DaubExPhase" or "DaubLeAsymm". Default is "DaubLeAsymm".   |
| filter.number        | The number of vanishing moments of the wavelet. By default 6.   |
| bc                   | The boundary condition to use - "periodic" or "symmetric". Default is "periodic".   |

### Details

The wavelet reconstruction is performed by using R package 'wavethresh'. For details, check wavethresh package help.

### Value

A list containing the following.

|                   |   |
|-------------------|---|
| GLMcoeffposterior | An array of dimension (nsubject, grid, grid), containing for each subject the posterior means of the standardized GLM coefficients. |
|-------------------|---|

**Author(s)**

Nilotpal Sanyal <nsanyal@stanford.edu>, Marco Ferreira <marf@vt.edu>

**Examples**

```
nsubject <- 3
grid <- 8
GLMCoeffStandardized <- array(rnorm(3*8*8),dim=c(3,8,8))
PostMeanWaveletCoeff <- array(rnorm(3*63),dim=c(3,63))
postglm.coeff <- postglmcoeff(nsubject, grid, GLMCoeffStandardized, PostMeanWaveletCoeff)
dim(postglm.coeff$GLMcoeffposterior)
#[1] 3 8 8
```

---

|                |  |
|----------------|--|
| postgroupcoeff | <i>Obtain posterior group coefficients using the BHMSMA methodology.</i> |
|----------------|--|

---

**Description**

Computes posterior group coefficients using the BHMSMA methodology.

**Usage**

```
postgroupcoeff( nsubject, grid, GLMCoeffStandardized, PostMeanWaveletCoeff,
  wave.family="DaubLeAsymm", filter.number=6, bc="periodic" )
```

**Arguments**

|                      |   |
|----------------------|---|
| nsubject             | Number of subjects included in the analysis.  |
| grid                 | The number of voxels in one row (or, one column) of the brain slice of interest. Must be a power of 2. The total number of voxels is grid <sup>2</sup> . The maximum grid value for this package is 512.    |
| GLMCoeffStandardized | An array of dimension (nsubject, grid, grid), containing for each subject the standardized GLM coefficients obtained by fitting GLM to the time-series corresponding to the voxels.                         |
| PostMeanWaveletCoeff | A matrix of size (nsubject, grid <sup>2</sup> -1), containing for each subject the posterior mean of the wavelet coefficients of all levels stacked together (by the increasing order of resolution level). |
| wave.family          | The family of wavelets to use - "DaubExPhase" or "DaubLeAsymm". Default is "DaubLeAsymm".   |
| filter.number        | The number of vanishing moments of the wavelet. By default 6.   |
| bc                   | The boundary condition to use - "periodic" or "symmetric". Default is "periodic".   |

**Details**

The wavelet computations are performed by using R package 'wavethresh'. For details, check wavethresh package help.

**Value**

A list containing the following.

`groupcoeff` A matrix of dimension (grid, grid), containing the posterior group coefficients obtained by BHMSMA methodology.

**Author(s)**

Nilotpal Sanyal <nsanyal@stanford.edu>, Marco Ferreira <marf@vt.edu>

**References**

Sanyal, Nilotpal, and Ferreira, Marco A.R. (2012). Bayesian hierarchical multi-subject multiscale analysis of functional MRI data. *Neuroimage*, 63, 3, 1519-1531.

**Examples**

```

nsubject <- 3
grid <- 8
GLMCoeffStandardized <- array(rnorm(3*8*8),dim=c(3,8,8))
PostMeanWaveletCoeff <- array(rnorm(3*63),dim=c(3,63))
post.groupcoeff <- postgroupcoeff( nsubject, grid, GLMCoeffStandardized, PostMeanWaveletCoeff)
dim(post.groupcoeff$groupcoeff)
#[1] 8 8

```

---

|             |  |
|-------------|--|
| postsamples | <i>Generate samples from the posterior distribution of the GLM coefficients.</i> |
|-------------|--|

---

**Description**

Generates samples from the posterior distribution of the GLM coefficients.

**Usage**

```
postsamples(nsample, nsubject, grid, GLMCoeffStandardized, WaveletCoefficientMatrix,
hyperparam, pklj.bar, analysis, wave.family="DaubLeAsymm", filter.number=6, bc="periodic")
```

**Arguments**

|                                       |   |
|---------------------------------------|---|
| <code>nsample</code>                  | Number of samples to be generated.  |
| <code>nsubject</code>                 | Number of subjects included in the analysis.  |
| <code>grid</code>                     | The number of voxels in one row (or, one column) of the brain slice of interest. Must be a power of 2. The total number of voxels is $\text{grid}^2$ . The maximum grid value for this package is 512.      |
| <code>GLMCoeffStandardized</code>     | An array of dimension $(\text{nsubject}, \text{grid}, \text{grid})$ , containing for each subject the standardized GLM coefficients obtained by fitting GLM to the time-series corresponding to the voxels. |
| <code>WaveletCoefficientMatrix</code> | A matrix of dimension $(\text{nsubject}, \text{grid}^2-1)$ , containing for each subject the wavelet coefficients of all levels stacked together (by the increasing order of resolution level).             |
| <code>hyperparam</code>               | A vector containing the estimates of the six hyperparameters.   |
| <code>pklj.bar</code>                 | A matrix of dimension $(\text{nsubject}, \text{grid}^2-1)$ , containing the $p_{klj}$ bar values (see References for details).  |
| <code>analysis</code>                 | "MSA" or "SSA", depending on whether performing multi-subject analysis or single subject analysis.  |
| <code>wave.family</code>              | The family of wavelets to use - "DaubExPhase" or "DaubLeAsymm". Default is "DaubLeAsymm".   |
| <code>filter.number</code>            | The number of vanishing moments of the wavelet. By default 6.   |
| <code>bc</code>                       | The boundary condition to use - "periodic" or "symmetric". Default is "periodic".   |

**Details**

The wavelet computations are performed by using R package 'wavethresh'. For details, check wavethresh package help.

**Value**

A list containing the following.

|                            |  |
|----------------------------|--|
| <code>samples</code>       | An array of dimension $(\text{nsubject}, \text{grid}, \text{grid}, \text{nsample})$ , containing for each subject the posterior samples of the GLM coefficients.                               |
| <code>postdiscovery</code> | An array of dimension $(\text{nsubject}, \text{grid}, \text{grid})$ , containing for each subject the posterior discovery maps of the GLM coefficients (for details see Morris et al. (2011)). |

**Author(s)**

Nilotpal Sanyal <nsanyal@stanford.edu>, Marco Ferreira <marf@vt.edu>

## References

Sanyal, Nilotpal, and Ferreira, Marco A.R. (2012). Bayesian hierarchical multi-subject multiscale analysis of functional MRI data. *Neuroimage*, 63, 3, 1519-1531.

Morris, J.S. et al. (2011). Automated analysis of quantitative image data using isomorphic functional mixed models, with application to proteomic data. *Ann. Appl. Stat.* 5, 894-923.

## Examples

```
nsubject <- 3
grid <- 8
nsample <- 5
GLMCoeffStandardized <- array(rnorm(3*8*8),dim=c(3,8,8))
WaveletCoefficientMatrix <- array(rnorm(3*63),dim=c(3,63))
hyperparam <- rep(.2,6)
pklj.bar <- array(runif(3*63),dim=c(3,63))
analysis <- "multi"
post.samples <- postsamples(nsample, nsubject, grid, GLMCoeffStandardized,
WaveletCoefficientMatrix, hyperparam, pklj.bar, analysis)
dim(post.samples$samples)
#[1] 3 8 8 5
```

---

|                  |   |
|------------------|---|
| postwaveletcoeff | <i>Obtain posterior mean and posterior median of the wavelet coefficients using BHMSMA methodology.</i> |
|------------------|---|

---

## Description

Computes posterior mean and posterior median of the wavelet coefficients using BHMSMA methodology.

## Usage

```
postwaveletcoeff(nsubject, grid, WaveletCoefficientMatrix, hyperparam,
pklj.bar, analysis)
```

## Arguments

|                          |  |
|--------------------------|--|
| nsubject                 | Number of subjects included in the analysis.   |
| grid                     | The number of voxels in one row (or, one column) of the brain slice of interest. Must be a power of 2. The total number of voxels is grid <sup>2</sup> . The maximum grid value for this package is 512. |
| WaveletCoefficientMatrix | A matrix of dimension (nsubject, grid <sup>2</sup> -1), containing for each subject the wavelet coefficients of all levels stacked together (by the increasing order of resolution level).               |
| hyperparam               | A vector containing the estimates of the six hyperparameters.  |

|          |   |
|----------|---|
| pklj.bar | A matrix of dimension (nsubject, grid^2-1), containing the pklj bar values (see Reference for details). |
| analysis | "MSA" or "SSA", depending on whether performing multi-subject analysis or single subject analysis.      |

**Value**

A list containing the following.

PostMeanWaveletCoeff

A matrix of size (nsubject, grid^2-1), containing for each subject the posterior mean of the wavelet coefficients of all levels stacked together (by the increasing order of resolution level).

PostMedianWaveletCoeff

A matrix of size (nsubject, grid^2-1), containing for each subject the posterior median of the wavelet coefficients of all levels stacked together.

**Author(s)**

Nilotpal Sanyal <nsanyal@stanford.edu>, Marco Ferreira <marf@vt.edu>

**References**

Sanyal, Nilotpal, and Ferreira, Marco A.R. (2012). Bayesian hierarchical multi-subject multiscale analysis of functional MRI data. *Neuroimage*, 63, 3, 1519-1531.

**Examples**

```
nsubject <- 3
grid <- 8
nsample <- 5
GLMCoeffStandardized <- array(rnorm(3*8*8),dim=c(3,8,8))
WaveletCoefficientMatrix <- array(rnorm(3*63),dim=c(3,63))
hyperparam <- rep(.2,6)
pklj.bar <- array(runif(3*63),dim=c(3,63))
analysis <- "multi"
post.waveletcoeff <- postwaveletcoeff(nsubject, grid, WaveletCoefficientMatrix,
hyperparam, pklj.bar, analysis)
dim(post.waveletcoeff$PostMeanWaveletCoeff)
#[1] 3 63
```

---

read.fmridata

*Read fMRI data from fMRI image files.*

---

**Description**

Reads fMRI data from fMRI image files into a 4D array.

**Usage**

```
read.fmridata( directory, format, prefix, nimages, dim.image, nii=TRUE )
```

**Arguments**

|           |   |
|-----------|---|
| directory | Location of the directory where the fMRI image files are stored. Insert within quotations ("").   |
| format    | The format of the data file. One of "Analyze" (.img/.hdr files), "Nifti" (.img/.hdr files or .nii files) or "Afni" (.HEAD/.BRIK files).   |
| prefix    | If format is "Analyze" or "Nifti", then the part of the fMRI image file name appearing before the image number. The image number is assumed to have four digit representation, that is, the lowest number is 0001 and the highest possible number is 9999. If format is "Afni", then the file name. Insert within quotations(""). |
| nimages   | If format is "Analyze", number of images to be read beginning from the first image. If format is "Afni", not necessary.   |
| dim.image | Size of the 3D fMRI image. A vector with three elements.  |
| nii       | Necessary only for "Nifti" format. nii=TRUE (default) indicates the image files are in .nii files . nii=FALSE indicates the image files are .img/.hdr files.  |

**Details**

The function uses package 'oro.nifti' for reading from fMRI data files. For details, check oro.nifti package help.

**Value**

A list containing the following:

|          |   |
|----------|---|
| fmridata | An array of dimension (dim.image, nimages), containing the image data for all images/time-points. |
|----------|---|

**Author(s)**

Nilotpal Sanyal <nsanyal@stanford.edu>, Marco Ferreira <marf@vt.edu>

**See Also**

[readANALYZE](#), [readNIFTI](#), [readAFNI](#)

**Examples**

```
# See package vignette.
```

---

|              |  |
|--------------|--|
| waveletcoeff | <i>Apply discrete wavelet transform to the GLM coefficients and obtain the wavelet coefficients.</i> |
|--------------|--|

---

### Description

Applies 2D discrete wavelet transform to the standardized GLM coefficient maps and returns the wavelet coefficients of all resolution levels.

### Usage

```
waveletcoeff(nsubject, grid, GLMCoeffStandardized,
wave.family="DaubLeAsymm", filter.number=6, bc="periodic")
```

### Arguments

|                      |  |
|----------------------|--|
| nsubject             | Number of subjects included in the analysis.   |
| grid                 | The number of voxels in one row (or, one column) of the brain slice of interest. Must be a power of 2. The total number of voxels is grid <sup>2</sup> . The maximum grid value for this package is 512. |
| GLMCoeffStandardized | An array of dimension (nsubject, grid, grid), containing for each subject the standardized GLM coefficients obtained by fitting GLM to the time-series corresponding to the voxels.                      |
| wave.family          | The family of wavelets to use - "DaubExPhase" or "DaubLeAsymm". Default is "DaubLeAsymm".  |
| filter.number        | The number of vanishing moments of the wavelet. By default 6.  |
| bc                   | The boundary condition to use - "periodic" or "symmetric". Default is "periodic".  |

### Details

The wavelet decomposition is performed by using R package 'wavethresh'. For details, check wavethresh package help.

### Value

A list containing the following.

WaveletCoefficientMatrix

A matrix of dimension (nsubject, grid<sup>2</sup>-1), containing for each subject the wavelet coefficients of all levels stacked together (by the increasing order of resolution level).

### Author(s)

Nilotpal Sanyal <nsanyal@stanford.edu>, Marco Ferreira <marf@vt.edu>



**Examples**

```
nsubject <- 3
grid <- 8
Data <- array(dim=c(3,8,8,10),rnorm(3*8*8*10))
DesignMatrix <- cbind( c(rep(c(1,0),5)), rep(1,10) )
glm.fit <- glmcoeff(nsubject, grid, Data, DesignMatrix)
GLMCoeffStandardized <- glm.fit$GLMCoeffStandardized
wavelet.coeff <- waveletcoeff(nsubject, grid, GLMCoeffStandardized)
dim(wavelet.coeff$WaveletCoefficientMatrix)
#[1] 3 63
```

# Index

BHMSMA, [2](#)  
BHMSMAfMRI (BHMSMAfMRI-package), [2](#)  
BHMSMAfMRI-package, [2](#)  
  
fmridata, [4](#)  
  
glmcoeff, [5](#)  
  
hyperparamest, [6](#)  
  
pikljbar, [7](#)  
postglmcoeff, [9](#)  
postgroupcoeff, [10](#)  
postsamples, [11](#)  
postwaveletcoeff, [13](#)  
  
read.fmridata, [14](#)  
readAFNI, [15](#)  
readANALYZE, [15](#)  
readNIfTI, [15](#)  
  
waveletcoeff, [16](#)