

# Package ‘mcmcr’

June 27, 2019

**Title** Manipulate MCMC Samples

**Version** 0.2.0

**Description** Functions and classes to store, manipulate and summarise Monte Carlo Markov Chain (MCMC) samples. For more information see Brooks et al. (2011) <isbn:978-1-4200-7941-8>.

**License** MIT + file LICENSE

**Depends** R (>= 3.5)

**Imports** abind, err, checkr, coda, utils, stats

**Suggests** covr, graphics, testthat

**URL** <https://github.com/poissonconsulting/mcmcr>

**BugReports** <https://github.com/poissonconsulting/mcmcr/issues>

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 6.1.1

**Language** en-US

**NeedsCompilation** no

**Author** Joe Thorley [aut, cre] (<<https://orcid.org/0000-0002-7683-4592>>)

**Maintainer** Joe Thorley <[joe@poissonconsulting.ca](mailto:joe@poissonconsulting.ca)>

**Repository** CRAN

**Date/Publication** 2019-06-27 16:30:03 UTC

## R topics documented:

anyNA . . . . .	3
as.mccarray . . . . .	4
as.mcmc . . . . .	4
as.mcmc.list . . . . .	5
as.mcmcarray . . . . .	6
as.mcmcr . . . . .	7

as.mcmcrs . . . . .	8
as.term . . . . .	9
bind_chains . . . . .	10
bind_dimensions . . . . .	11
bind_dimensions_n . . . . .	12
bind_iterations . . . . .	12
bind_parameters . . . . .	13
check_mcmcarray . . . . .	14
check_mcmcr . . . . .	15
coef . . . . .	16
collapse_chains . . . . .	17
combine_dimensions . . . . .	18
combine_samples . . . . .	19
combine_samples_n . . . . .	20
converged . . . . .	20
dims . . . . .	22
esr . . . . .	22
ess . . . . .	24
estimates . . . . .	24
is.marray . . . . .	25
is.mcmcarray . . . . .	26
is.mcmcr . . . . .	26
is.mcmcrs . . . . .	27
is.term . . . . .	27
mcmcarray-object . . . . .	28
mcmcr-object . . . . .	28
mcmcrs . . . . .	29
mcmcrs-object . . . . .	29
mcmcr_example . . . . .	30
mcmc_aperm . . . . .	30
mcmc_map . . . . .	31
nchains . . . . .	32
ndims . . . . .	34
niters . . . . .	34
npars . . . . .	35
npdims . . . . .	36
npdims.default . . . . .	36
nsams . . . . .	37
nsims . . . . .	38
nterms . . . . .	39
parameters . . . . .	40
pvalue . . . . .	41
rhat . . . . .	42
sort . . . . .	43
split_chains . . . . .	44
subset . . . . .	45
tdims . . . . .	46
term . . . . .	46

<code>anyNA</code>	3
<code>terms</code> . . . . .	47
<code>thin</code> . . . . .	48
<code>zero</code> . . . . .	49

<b>Index</b>	<b>50</b>
--------------	-----------

---

<code>anyNA</code>	<i>Any Missing Values</i>
--------------------	---------------------------

---

### Description

Test whether there are any missing values.

### Usage

```
## S3 method for class 'marray'
anyNA(x, recursive = FALSE)
```

```
## S3 method for class 'mcmc'
anyNA(x, recursive = FALSE)
```

```
## S3 method for class 'mcmc.list'
anyNA(x, recursive = FALSE)
```

```
## S3 method for class 'mcmcarray'
anyNA(x, recursive = FALSE)
```

```
## S3 method for class 'mcmcr'
anyNA(x, recursive = FALSE)
```

```
## S3 method for class 'mcmcrs'
anyNA(x, recursive = FALSE)
```

### Arguments

<code>x</code>	The object to test.
<code>recursive</code>	Unused.

### Value

A flag indicating whether there are any missing values.

### See Also

[base::anyNA](#)

**Examples**

```
anyNA(mcmcr_example)
anyNA(mcmcr_example$beta)
```

---

as.marray	<i>Coerce to an marray object</i>
-----------	-----------------------------------

---

**Description**

Coerces MCMC objects to an marray object.

**Usage**

```
as.marray(x, ...)
```

```
## S3 method for class 'mcmarray'
as.marray(x, ...)
```

**Arguments**

x	object to coerce.
...	Unused.

**Methods (by class)**

- mcmarray: Coerces mcmarray object to an marray object

**Examples**

```
as.marray(mcmcr_example$beta)
```

---

as.mcmc	<i>Coerce to an mcmc object</i>
---------	---------------------------------

---

**Description**

Coerces MCMC objects to an [mcmc](#) object.

**Usage**

```
## S3 method for class 'mccarry'
as.mcmc(x, ...)

## S3 method for class 'mcmcarray'
as.mcmc(x, ...)

## S3 method for class 'mcmcr'
as.mcmc(x, ...)
```

**Arguments**

```
x          object to coerce.
...        Unused.
```

**Methods (by class)**

- `mccarry`: Coerces `mccarry` object (with 1 chain) to an `mcmc` object
- `mcmcarray`: Coerces `mcmcarray` object (with 1 chain) to an `mcmc` object
- `mcmcr`: Coerces `mcmcr` object (with 1 chain) to an `mcmc` object

**See Also**

`coda::mcmc`

**Examples**

```
as.mcmc(subset(mcmcr_example, chains = 1L))
```

---

<code>as.mcmc.list</code>	<i>Coerce to an <code>mcmc.list</code> object</i>
---------------------------	---

---

**Description**

Coerces MCMC objects to an `mcmc.list` object.

**Usage**

```
## S3 method for class 'mcmcarray'
as.mcmc.list(x, ...)

## S3 method for class 'mcmc'
as.mcmc.list(x, ...)

## S3 method for class 'mcmcr'
as.mcmc.list(x, ...)
```

**Arguments**

x                    object to coerce.  
 ...                  Unused.

**Methods (by class)**

- mcmcarray: Coerces mcmcarray object to an mcmc.list object
- mcmc: Coerces mcmc object to an mcmc.list object
- mcmcr: Coerces mcmcr object to an mcmc.list object

---

 as.mcmcarray

*Coerce to an mcmcarray object*


---

**Description**

Coerces MCMC objects to an [mcmcarray-object](#).

**Usage**

```
as.mcmcarray(x, ...)  
  
## Default S3 method:  
as.mcmcarray(x, ...)  
  
## S3 method for class 'marray'  
as.mcmcarray(x, ...)  
  
## S3 method for class 'mcmc'  
as.mcmcarray(x, ...)  
  
## S3 method for class 'mcmc.list'  
as.mcmcarray(x, ...)  
  
## S3 method for class 'mcmcr'  
as.mcmcarray(x, ...)
```

**Arguments**

x                    object to coerce.  
 ...                  Unused.

**Methods (by class)**

- `default`: Coerces vector, matrix or array to an `mcmcarray` object
- `mccarray`: Coerces `mccarray` object to an `mcmcarray` object
- `mcmc`: Coerces `mcmc` object (with one parameter) to an `mcmcarray` object
- `mcmc.list`: Coerces `mcmc.list` object (with one parameter) to an `mcmcarray` object
- `mcmcr`: Coerces `mcmcr` object (with one parameter) to an `mcmcarray` object

**Examples**

```
as.mcmcarray(as.mccarray(mcmcr_example$beta))
```

---

<code>as.mcmcr</code>	<i>Coerce to an mcmcr object</i>
-----------------------	----------------------------------

---

**Description**

Coerces MCMC objects to an [mcmcr-object](#).

**Usage**

```
as.mcmcr(x, ...)

## S3 method for class 'list'
as.mcmcr(x, ...)

## S3 method for class 'mccarray'
as.mcmcr(x, ...)

## S3 method for class 'mcmc'
as.mcmcr(x, ...)

## S3 method for class 'mcmc.list'
as.mcmcr(x, ...)

## S3 method for class 'mcmcarray'
as.mcmcr(x, ...)

## S3 method for class 'list'
as.mcmcrs(x, ...)
```

**Arguments**

<code>x</code>	object to coerce.
<code>...</code>	Unused.

**Methods (by class)**

- `list`: Coerces list (of `mcmcarray` objects) to an `mcmc` object
- `marray`: Coerces `marray` object to an `mcmc` object
- `mcmc`: Coerces `mcmc` object to an `mcmc` object
- `mcmc.list`: Coerces `mcmc.list` object to an `mcmc` object
- `mcmcarray`: Coerces `mcmcarray` object to an `mcmc` object
- `list`: Coerces list (of `mcmc` objects with the same parameters, chains and iterations) to an `mcmcs` object

**Examples**

```
as.mcmc(coda::as.mcmc.list(mcmc_example))
```

---

`as.mcmcs`

*Coerce to an mcmcs object*

---

**Description**

Coerces MCMC objects to an `mcmcs-object`.

**Usage**

```
as.mcmcs(x, ...)
```

**Arguments**

<code>x</code>	object to coerce.
<code>...</code>	Unused.

**Examples**

```
as.mcmcs(list(mcmc_example))
```



---

as.term	<i>Coerce to a term vector</i>
---------	--------------------------------

---

### Description

Coerces MCMC objects to a term vector.

### Usage

```
as.term(x, ...)  
  
## S3 method for class 'term'  
as.character(x, ...)  
  
## S3 method for class 'character'  
as.term(x, ...)  
  
## S3 method for class 'mcmc'  
as.term(x, ...)  
  
## S3 method for class 'mcmc.list'  
as.term(x, ...)  
  
## S3 method for class 'mcmcarray'  
as.term(x, ...)  
  
## S3 method for class 'mcmcr'  
as.term(x, ...)
```

### Arguments

x	The object to coerce
...	Unused.

### Methods (by class)

- term: Coerces term vector to a character vector
- character: Coerces character vector to a term vector
- mcmc: Coerces mcmc object to a term vector
- mcmc.list: Coerces mcmc.list object to a term vector
- mcmcarray: Coerces mcmcarray object to a term vector
- mcmcr: Coerces mcmcr object to a term vector

### Examples

```
as.term(mcmcr_example)
```

---

`bind_chains`*Combine MCMC objects by chains.*

---

### Description

Combines two MCMC objects (with the same parameters and iterations) by chains.

### Usage

```
bind_chains(x, x2, ...)

## S3 method for class 'mccarray'
bind_chains(x, x2, ...)

## S3 method for class 'mcmc'
bind_chains(x, x2, ...)

## S3 method for class 'mcmc.list'
bind_chains(x, x2, ...)

## S3 method for class 'mcmcarray'
bind_chains(x, x2, ...)

## S3 method for class 'mcmcr'
bind_chains(x, x2, ...)
```

### Arguments

<code>x</code>	an MCMC object.
<code>x2</code>	a second MCMC object
<code>...</code>	Unused.

### Methods (by class)

- `mccarray`: Binds two `mccarray` objects by their chains
- `mcmc`: Binds two `mcmc` objects by their chains
- `mcmc.list`: Binds two `mcmc.list` objects by their chains
- `mcmcarray`: Binds two `mcmcarray` objects by their chains
- `mcmcr`: Binds two `mcmcr` objects by their chains

### Examples

```
bind_chains(mcmcr_example, mcmcr_example)
```

---

bind_dimensions	<i>Combine two MCMC objects by dimensions</i>
-----------------	---

---

### Description

Combines multiple MCMC objects (with the same parameters, chains and iterations) by parameter dimensions.

### Usage

```
bind_dimensions(x, x2, along = NULL, ...)  
  
## S3 method for class 'mcmcarray'  
bind_dimensions(x, x2, along = NULL, ...)  
  
## S3 method for class 'mcmcr'  
bind_dimensions(x, x2, along = NULL, ...)
```

### Arguments

x	An MCMC object.
x2	a second MCMC object.
along	A count (or NULL) indicating the parameter dimension to bind along.
...	Unused.

### Methods (by class)

- mcmcarray: Binds two mcmcarray objects by their dimensions
- mcmcr: Binds two mcmcr objects by their dimensions

### See Also

[bind\\_dimensions\\_n](#)

### Examples

```
bind_dimensions(mcmcr_example, mcmcr_example)
```

---

bind\_dimensions\_n      *Combine multiple MCMC objects by parameter dimensions*

---

### Description

Combines multiple MCMC objects (with the same parameters, chains and iterations) by parameter dimensions.

### Usage

```
bind_dimensions_n(...)  
  
## S3 method for class 'mcmcarray'  
bind_dimensions_n(...)  
  
## S3 method for class 'mcmcr'  
bind_dimensions_n(...)
```

### Arguments

...                    one or more MCMC objects

### Methods (by class)

- mcmcarray: Binds multiple mcmcarray objects by their dimensions
- mcmcr: Binds multiple mcmcr objects by their dimensions

### See Also

[bind\\_dimensions](#)

### Examples

```
bind_dimensions_n(mcmcr_example, mcmcr_example, mcmcr_example)
```

---

bind\_iterations      *Combine two MCMC objects by iterations*

---

### Description

Combines two MCMC objects (with the same parameters and chains) by iterations.

**Usage**

```
bind_iterations(x, x2, ...)

## S3 method for class 'mccarray'
bind_iterations(x, x2, ...)

## S3 method for class 'mcmc'
bind_iterations(x, x2, ...)

## S3 method for class 'mcmc.list'
bind_iterations(x, x2, ...)

## S3 method for class 'mcmcarray'
bind_iterations(x, x2, ...)

## S3 method for class 'mcmcr'
bind_iterations(x, x2, ...)
```

**Arguments**

x	an MCMC object
x2	a second MCMC object
...	unused

**Methods (by class)**

- `mccarray`: Binds two `mccarray` objects by their iterations
- `mcmc`: Binds two `mcmc` objects by their iterations
- `mcmc.list`: Binds two `mcmc.list` objects by their iterations
- `mcmcarray`: Binds two `mcmcarray` objects by their iterations
- `mcmcr`: Binds two `mcmcr` objects by their iterations

**Examples**

```
bind_iterations(mcmcr_example, mcmcr_example)
```

---

bind_parameters	<i>Combine two MCMC object by parameters</i>
-----------------	--

---

**Description**

Combines two MCMC objects (with the same chains and iterations) by their parameters.

**Usage**

```

bind_parameters(x, ...)

## S3 method for class 'mcmc'
bind_parameters(x, x2, ...)

## S3 method for class 'mcmc.list'
bind_parameters(x, x2, ...)

## S3 method for class 'mcmcr'
bind_parameters(x, x2, ...)

```

**Arguments**

x	an MCMC object
...	unused
x2	a second MCMC object

**Methods (by class)**

- `mcmc`: Binds two `mcmc` objects by their parameters
- `mcmc.list`: Binds two `mcmc.list` objects by their parameters
- `mcmcr`: Binds two `mcmcr` objects by their parameters

**Examples**

```

bind_parameters(subset(mcmcr_example, parameters = "sigma"),
  subset(mcmcr_example, parameters = "beta"))

```

---

check_mcmcarray	<i>Check mcmcarray</i>
-----------------	------------------------

---

**Description**

Check mcmcarray

**Usage**

```

check_mcmcarray(x, x_name = substitute(x), error = TRUE)

```

**Arguments**

x	The object to check.
x_name	A string of the name of the object.
error	A flag indicating whether to throw an informative error or immediately generate an informative message if the check fails.

**Value**

An invisible copy of x (it if doesn't throw an error).

**Examples**

```
check_mcmcrarray(mcmcr::mcmcr_example$beta)
```

---

 check\_mcmcr

*Check mcmcr*


---

**Description**

Check mcmcr

**Usage**

```
check_mcmcr(x, sorted = FALSE, x_name = substitute(x), error = TRUE)
```

**Arguments**

x	The object to check.
sorted	A flag specifying whether the parameters must be sorted.
x_name	A string of the name of the object.
error	A flag indicating whether to throw an informative error or immediately generate an informative message if the check fails.

**Value**

An invisible copy of x (it if doesn't throw an error).

**Examples**

```
check_mcmcr(mcmcr::mcmcr_example)
```

---

coef *Term Coefficients*

---

**Description**

Gets coefficients for all the terms in an MCMC object.

**Usage**

```
## S3 method for class 'mccarray'
coef(object, conf_level = 0.95,
      estimate = stats::median, ...)

## S3 method for class 'mcmc'
coef(object, conf_level = 0.95,
      estimate = stats::median, ...)

## S3 method for class 'mcmc.list'
coef(object, conf_level = 0.95,
      estimate = stats::median, ...)

## S3 method for class 'mcmcarray'
coef(object, conf_level = 0.95,
      estimate = stats::median, ...)

## S3 method for class 'mcmcr'
coef(object, conf_level = 0.95,
      estimate = stats::median, ...)
```

**Arguments**

object	The MCMC object to get the coefficients for
conf_level	A number specifying the confidence level. By default 0.95.
estimate	The function to use to calculate the estimate.
...	Unused

**Value**

An data frame of the coefficients with the columns indicating the term, estimate, standard deviation (sd), zscore, lower and upper credible intervals and pvalue.

**Methods (by class)**

- `mccarray`: Get coefficients for terms in `mccarray` object
- `mcmc`: Get coefficients for terms in `mcmc` object
- `mcmc.list`: Get coefficients for terms in `mcmc.list` object



- `mcmcarray`: Get coefficients for terms in `mcmcarray` object
- `mcmcr`: Get coefficients for terms in `mcmcr` object

### See Also

`stats::coef`

### Examples

```
coef(mcmcr_example)
```

---

collapse_chains	<i>Collapse Chains</i>
-----------------	------------------------

---

### Description

Collapses an MCMC object's chains into a single chain.

### Usage

```
collapse_chains(x, ...)  
  
## Default S3 method:  
collapse_chains(x, ...)  
  
## S3 method for class 'mcmc.list'  
collapse_chains(x, ...)  
  
## S3 method for class 'mcmcr'  
collapse_chains(x, ...)
```

### Arguments

<code>x</code>	An MCMC object.
<code>...</code>	Unused.

### Methods (by class)

- `default`: Collapse an MCMC object's chains into a single chain
- `mcmc.list`: Collapse an `mcmc.list` object's chains into a single chain
- `mcmcr`: Collapse an `mcmcr` object's chains into a single chain

### Examples

```
collapse_chains(mcmcr_example)
```

---

combine\_dimensions      *Combine Samples by Dimensions*

---

### Description

Combines MCMC object samples by dimensions along using fun.

### Usage

```
combine_dimensions(x, fun = mean, along = NULL, ...)
```

```
## S3 method for class 'mcmcarray'  
combine_dimensions(x, fun = mean, along = NULL,  
  ...)
```

```
## S3 method for class 'mcmcr'  
combine_dimensions(x, fun = mean, along = NULL, ...)
```

### Arguments

x	An MCMC object
fun	The function to use when combining dimensions
along	A positive integer (or NULL) indicating the parameter dimension(s) to bind along.
...	Unused

### Value

The MCMC object with reduced dimensions.

### Methods (by class)

- mcmcarray: Combine an mcmcarray object's samples by dimensions
- mcmcr: Combine an mcmcr object's samples by dimensions

### Examples

```
combine_dimensions(mcmcr_example$alpha)
```

---

combine_samples	<i>Combine MCMC Samples of Two Objects</i>
-----------------	--

---

## Description

Combines samples of two MCMC objects (with the same parameters, chains and iterations) using a function.

## Usage

```
combine_samples(x, x2, fun = mean, ...)  
  
## S3 method for class 'mcmcarray'  
combine_samples(x, x2, fun = mean, ...)  
  
## S3 method for class 'mcmcr'  
combine_samples(x, x2, fun = mean, ...)
```

## Arguments

x	An MCMC object.
x2	A second MCMC object.
fun	The function to use to combine the samples. The function must return a scalar.
...	Unused.

## Value

The combined samples as an MCMC object with the same parameters, chains and iterations as the original objects.

## Methods (by class)

- mcmcarray: Combine samples of two mcmcarray objects
- mcmcr: Combine samples of two mcmcr objects

## Examples

```
combine_samples(mcmcr_example, mcmcr_example, fun = sum)
```

---

combine_samples_n	<i>Combine MCMC Samples of multiple objects</i>
-------------------	---

---

### Description

Combines samples of multiple MCMC objects (with the same parameters, chains and iterations) using a function.

### Usage

```
combine_samples_n(x, ..., fun = mean)

## Default S3 method:
combine_samples_n(x, ..., fun = mean)

## S3 method for class 'list'
combine_samples_n(x, ..., fun = mean)
```

### Arguments

x	An MCMC object (or a list of mcmc objects).
...	Additional MCMC objects.
fun	A function.

### Methods (by class)

- default: Combine samples of multiple MCMC objects
- list: Combine samples of a list of multiple MCMC objects

### Examples

```
combine_samples_n(mcmcr_example, mcmcr_example, mcmcr_example, fun = sum)
```

---

converged	<i>Object Converged</i>
-----------	-------------------------

---

### Description

Test whether an object has converged.

## Usage

```
converged(x, ...)  
  
## Default S3 method:  
converged(x, rhat = 1.1, esr = 0.33, by = "all",  
          as_df = FALSE, ...)  
  
## S3 method for class 'mcmcrg' :  
converged(x, rhat = 1.1, esr = 0.33, by = "all",  
          as_df = FALSE, bound = FALSE, ...)
```

## Arguments

x	An object.
...	Unused.
rhat	The maximum rhat value.
esr	The minimum effective sampling rate.
by	A string indicating whether to determine by "term", "parameter" or "all".
as_df	A flag indicating whether to return the values as a data frame versus a named list.
bound	flag specifying whether to bind mcmcrg objects by their chains before calculating rhat.

## Methods (by class)

- default: Test whether an object has converged
- mcmcrg: Test whether an mcmcrg object has converged

## See Also

[rhat](#) and [esr](#)

## Examples

```
converged(mcmcrg_example)  
converged(mcmcrg(mcmcrg_example, mcmcrg_example))  
converged(mcmcrg(mcmcrg_example, mcmcrg_example), bound = TRUE)
```

---

 dims

*Dimensions*


---

**Description**

Dimensions

**Usage**

```
dims(x)
```

**Arguments**

x                    A vector, matrix or array.

**Value**

An integer vector of the dimensions.

**Examples**

```
dims(character())
dims(2:3)
dims(matrix(1:6, nrow = 2))
```

---

 esr

*Effective Sampling Rate*


---

**Description**

Calculates the effective sampling rate (esr) based on the formula

$$\frac{1}{1 + 2 \sum_{k=1}^{\infty} \rho_k(\theta)}$$

in Brooks et al. (2011). The infinite sum is truncated at lag  $k$  when  $\rho_{k+1}(\theta) < 0$ .

**Usage**

```
esr(x, ...)
```

```
## S3 method for class 'marray'
esr(x, by = "all", ...)
```

```
## S3 method for class 'mcmc'
esr(x, by = "all", ...)
```

```

## S3 method for class 'mcmc.list'
esr(x, by = "all", ...)

## S3 method for class 'mcmcarray'
esr(x, by = "all", as_df = FALSE, ...)

## S3 method for class 'mcmcr'
esr(x, by = "all", as_df = FALSE, ...)

## S3 method for class 'mcmcrs'
esr(x, by = "all", as_df = FALSE, ...)

```

### Arguments

x	An MCMC object.
...	Unused.
by	A string indicating whether to determine by "term", "parameter" or "all".
as_df	A flag indicating whether to return the values as a data frame versus a named list.

### Value

The esr value(s) as a data frame or list

### Methods (by class)

- marray: Effective Sampling Rate for an marray object
- mcmc: Effective Sampling Rate for an mcmc object
- mcmc.list: Effective Sampling Rate for an mcmc.list object
- mcmcarray: Effective Sampling Rate for an mcmcarray object
- mcmcr: Effective Sampling Rate for an mcmcr object
- mcmcrs: Effective Sampling Rate for an mcmcrs object

### References

Brooks, S., Gelman, A., Jones, G.L., and Meng, X.-L. (Editors). 2011. Handbook for Markov Chain Monte Carlo. Taylor & Francis, Boca Raton.

### Examples

```

esr(mcmcr_example)
esr(mcmcrs(mcmcr_example, mcmcr_example))

```

---

ess *Effective Sample Size*

---

### Description

Calculates the effective sample size based on [esr](#).

### Usage

```
ess(x, by = "all", as_df = FALSE)
```

### Arguments

**x** An MCMC object.

**by** A string indicating whether to determine by "term", "parameter" or "all".

**as\_df** A flag indicating whether to return the results as a data frame or list.

### Examples

```
ess(mcmcr_example)
```

---

estimates *Estimates*

---

### Description

Get the estimates for an MCMC object.

### Usage

```
estimates(object, ...)

## S3 method for class 'mccarray'
estimates(object, fun = stats::median, as_df = FALSE,
  ...)

## S3 method for class 'mcmc'
estimates(object, fun = stats::median, as_df = FALSE,
  ...)

## S3 method for class 'mcmc.list'
estimates(object, fun = stats::median,
  as_df = FALSE, ...)

## S3 method for class 'mcmcarray'
```



```

estimates(object, fun = stats::median,
  as_df = FALSE, ...)

## S3 method for class 'mcmcr'
estimates(object, fun = stats::median, as_df = FALSE,
  ...)

```

### Arguments

object	An MCMC object.
...	Unused.
fun	The function to use.
as_df	A flag indicating whether to return the estimates as a data frame versus a named list.

### Value

A named list or data frame.

### Methods (by class)

- marray: Estimates for an marray object
- mcmc: Estimates for an mcmc object
- mcmc.list: Estimates for an mcmc.list object
- mcmarray: Estimates for an mcmarray object
- mcmcr: Estimates for an mcmcr object

### Examples

```

estimates(mcmcr_example)
estimates(mcmcr_example, as_df = TRUE)

```

---

is.marray	<i>Is marray Object</i>
-----------	-------------------------

---

### Description

Tests whether an object is an marray.

### Usage

```
is.marray(x)
```

### Arguments

x	The object to test.
---	---------------------

**Value**

A flag indicating whether the test was positive.

**Examples**

```
is.marray(mcmcr_example)
```

---

is.mcmarray	<i>Is mcmarray Object</i>
-------------	---------------------------

---

**Description**

Tests whether an object is an [mcmarray-object](#).

**Usage**

```
is.mcmarray(x)
```

**Arguments**

x                   The object to test.

**Value**

A flag indicating whether the test was positive.

**Examples**

```
is.mcmarray(mcmcr_example$beta)
```

---

is.mcmcr	<i>Is mcmcr Object</i>
----------	------------------------

---

**Description**

Tests whether an object is an [mcmcr-object](#).

**Usage**

```
is.mcmcr(x)
```

**Arguments**

x                   The object to test.

**Value**

A flag indicating whether the test was positive.

**Examples**

```
is.mcmcr(mcmcr_example)
```

---

is.mcmcrs

*Is mcmcrs Object*

---

**Description**

Tests whether an object is an [mcmcrs-object](#).

**Usage**

```
is.mcmcrs(x)
```

**Arguments**

x                    The object to test.

**Value**

A flag indicating whether the test was positive.

**Examples**

```
is.mcmcrs(mcmcrs(mcmcr_example))
```

---

is.term

*Is Term*

---

**Description**

Test whether an object is a term.

**Usage**

```
is.term(x)
```

**Arguments**

x                    The object to test.

**Value**

A flag indicating whether the test was positive.

**Examples**

```
is.term(terms(mcmcr_example))
```

---

mcmcrarray-object	<i>mcmcrarray</i>
-------------------	-------------------

---

**Description**

An *mcmcrarray* object is an array where the first dimension is the chains, the second dimension is the iterations and the subsequent dimensions represent the dimensionality of the parameter. The name *mcmcrarray* reflects the fact that the MCMC dimensions, ie the chains and iterations, precede the parameter dimensions.

**Examples**

```
mcmcr_example$beta
```

---

mcmcr-object	<i>mcmcr</i>
--------------	--------------

---

**Description**

An *mcmcr* object stores multiple uniquely named *mcmcrarray-object* objects with the same number of chains and iterations.

**Details**

*mcmcr* objects allow a set of dimensionality preserving parameters to be manipulated and queried as a whole.

**Examples**

```
mcmcr_example
```

---

mcmcrs	<i>Create mcmcrs</i>
--------	----------------------

---

**Description**

Creates an `mcmcrs-object` from multiple `link{mcmcr-object}`s.

**Usage**

```
mcmcrs(...)
```

**Arguments**

...                    Objects of class `mcmcr`.

**Value**

An object of class `mcmcrs`

**Examples**

```
mcmcrs(mcmcr_example, mcmcr_example)
```

---

mcmcrs-object	<i>mcmcrs</i>
---------------	---------------

---

**Description**

An `mcmcrs` object stores multiple `mcmcr-objects` with the same parameters and the same number of chains and iterations.

**Details**

`mcmcrs` objects allow the results of multiple analyses using the same model to be manipulated and queried as a whole.

**Examples**

```
mcmcrs(mcmcr_example, mcmcr_example)
```

---

mcmcr\_example

*An Example mcmcr Object*


---

### Description

An example `mcmcr-object` derived from on `coda::line`.

### Usage

```
mcmcr_example
```

### Format

An object of class `mcmcr` of length 3.

### Examples

```
mcmcr_example
```

---

mcmc\_aperm

*MCMC Object Transposition*


---

### Description

Transpose an MCMC object by permuting its parameter dimensions.

### Usage

```
mcmc_aperm(x, perm, ...)
```

```
## S3 method for class 'mcmcarray'
```

```
mcmc_aperm(x, perm = NULL, ...)
```

```
## S3 method for class 'mcmc'
```

```
mcmc_aperm(x, perm = NULL, ...)
```

```
## S3 method for class 'mcmc.list'
```

```
mcmc_aperm(x, perm = NULL, ...)
```

```
## S3 method for class 'mcmcr'
```

```
mcmc_aperm(x, perm = NULL, ...)
```

```
## S3 method for class 'mcmcrs'
```

```
mcmc_aperm(x, perm = NULL, ...)
```

**Arguments**

x	The MCMC object to transpose.
perm	A integer vector of the new order for the parameter dimensions. Missing parameter dimensions are added on the end. If perm = NULL (the default) the parameter dimensions are reversed.
...	Unused

**Value**

The modified MCMC object

**Methods (by class)**

- `mcmcarray`: Transpose an `mcmcarray` object
- `mcmc`: Transpose an `mcmc` object
- `mcmc.list`: Transpose an `mcmc.list` object
- `mcmcr`: Transpose an `mcmcr` object
- `mcmcrs`: Transpose an `mcmcrs` object

---

mcmc\_map

*MCMC Map*


---

**Description**

Adjust the sample values of an MCMC object using a function.

**Usage**

```
mcmc_map(.x, .f, .by = 1:npdims(.x), ...)
```

```
## S3 method for class 'mcmcarray'
mcmc_map(.x, .f, .by = 1:npdims(.x), ...)
```

```
## S3 method for class 'mcmc'
mcmc_map(.x, .f, .by = TRUE, ...)
```

```
## S3 method for class 'mcmc.list'
mcmc_map(.x, .f, .by = TRUE, ...)
```

```
## S3 method for class 'mcmcr'
mcmc_map(.x, .f, .by = TRUE, ...)
```

```
## S3 method for class 'mcmcrs'
mcmc_map(.x, .f, .by = TRUE, ...)
```

**Arguments**

<code>.x</code>	An MCMC object
<code>.f</code>	The function to use
<code>.by</code>	A positive integer vector of the dimensions to apply the function over.
<code>...</code>	Additional arguments passed to <code>.f</code> .

**Value**

The updated MCMC object.

**Methods (by class)**

- `mcmcarray`: Adjust the sample values of an MCMC object
- `mcmc`: Adjust the sample values of an `mcmc.list` object
- `mcmc.list`: Adjust the sample values of an `mcmc.list` object
- `mcmcr`: Adjust the sample values of an `mcmcr` object
- `mcmcrs`: Adjust the sample values of an `mcmcrs` object

**Examples**

```
mcmc_map(mcmcr_example$beta, exp)
```

---

nchains	<i>Number of MCMC chains</i>
---------	------------------------------

---

**Description**

Gets the number of MCMC chains.

**Usage**

```
nchains(x, ...)
```

```
## S3 method for class 'marray'
```

```
nchains(x, ...)
```

```
## S3 method for class 'mcmc'
```

```
nchains(x, ...)
```

```
## S3 method for class 'mcmc.list'
```

```
nchains(x, ...)
```

```
## S3 method for class 'mcmcarray'
```

```
nchains(x, ...)
```



```

## S3 method for class 'mcmcr'
nchains(x, ...)

## S3 method for class 'mcmcrs'
nchains(x, ...)

## S3 method for class 'mccarray'
niters(x, ...)

## S3 method for class 'mcmc'
niters(x, ...)

## S3 method for class 'mcmc.list'
niters(x, ...)

## S3 method for class 'mcmccarray'
niters(x, ...)

## S3 method for class 'mcmcr'
niters(x, ...)

## S3 method for class 'mcmcrs'
niters(x, ...)

```

### Arguments

x	An MCMC object
...	Unused

### Value

A count indicating the number of MCMC chains

### Methods (by class)

- `mccarray`: Number of MCMC chains for an `mccarray` object
- `mcmc`: Number of MCMC chains for an `mcmc` object
- `mcmc.list`: Number of MCMC chains for an `mcmc.list` object
- `mcmccarray`: Number of MCMC chains for an `mcmccarray` object
- `mcmcr`: Number of MCMC chains for an `mcmcr` object
- `mcmcrs`: Number of MCMC chains for an `mcmcrs` object
- `mccarray`: Number of MCMC iterations for an `mccarray` object
- `mcmc`: Number of MCMC iterations for an `mcmc` object
- `mcmc.list`: Number of MCMC iterations for an `mcmc.list` object
- `mcmccarray`: Number of MCMC iterations for an `mcmccarray` object
- `mcmcr`: Number of MCMC iterations for an `mcmcr` object
- `mcmcrs`: Number of MCMC iterations for an `mcmcrs` object

**Examples**

```
nchains(mcmcr_example)
```

---

ndims	<i>Number of dimensions</i>
-------	-----------------------------

---

**Description**

Number of dimensions

**Usage**

```
ndims(x)
```

**Arguments**

x	A vector, matrix or array.
---	----------------------------

**Value**

A count of the number of dimensions

---

niters	<i>Number of MCMC samples</i>
--------	-------------------------------

---

**Description**

Gets the number of MCMC iterations (in a chain).

**Usage**

```
niters(x, ...)
```

**Arguments**

x	The object
...	Unused.

**Value**

A count indicating the number of MCMC iterations.

**Examples**

```
niters(mcmcr_example)
```

---

npars	<i>Number of Parameters</i>
-------	-----------------------------

---

**Description**

Gets the number of parameters for an object.

**Usage**

```
npars(x, ...)  
  
## Default S3 method:  
npars(x, ...)  
  
## S3 method for class 'marray'  
npars(x, ...)  
  
## S3 method for class 'mcmarray'  
npars(x, ...)  
  
## S3 method for class 'mcmcr'  
npars(x, ...)  
  
## S3 method for class 'mcmcrs'  
npars(x, ...)
```

**Arguments**

x	The object.
...	Not used.

**Value**

A count of the number of parameters.

**Methods (by class)**

- default: Number of parameters for an object
- marray: Number of parameters for an marray object
- mcmarray: Number of parameters for an mcmarray object
- mcmcr: Number of parameters for an mcmcr object
- mcmcrs: Number of parameters for an mcmcrs object

**Examples**

```
npars(mcmcr_example)
```

---

npdims	<i>Number of Parameter Dimensions</i>
--------	---------------------------------------

---

**Description**

Gets the number parameter dimensions of an object.

**Usage**

```
npdims(x, ...)
```

**Arguments**

x	The object
...	Unused.

**Examples**

```
pdims(mcmcr_example)
```

---

npdims.default	<i>Parameter Dimensions</i>
----------------	-----------------------------

---

**Description**

Gets the parameter dimensions of an object.

**Usage**

```
## Default S3 method:
npdims(x, ...)

## S3 method for class 'mcmc.list'
npdims(x, ...)

## S3 method for class 'mcmcr'
npdims(x, ...)

pdims(x, ...)

## S3 method for class 'term'
pdims(x, ...)

## S3 method for class 'mccarray'
pdims(x, ...)
```

```
## S3 method for class 'mcmc'
pdims(x, ...)

## S3 method for class 'mcmc.list'
pdims(x, ...)

## S3 method for class 'mcmcarray'
pdims(x, ...)

## S3 method for class 'mcmcr'
pdims(x, ...)
```

### Arguments

x	The object
...	Unused.

### Methods (by class)

- default: Parameter dimensions for a term vector
- mcmc.list: Parameter dimensions for an mcmcr object
- mcmcr: Parameter dimensions for an mcmcr object
- term: Parameter dimensions for a term vector
- marray: Parameter dimensions for an marray object
- mcmc: Parameter dimensions for an mcmc object
- mcmc.list: Parameter dimensions for an mcmc.list object
- mcmcarray: Parameter dimensions for an mcmcarray object
- mcmcr: Parameter dimensions for an mcmcr object

### Examples

```
pdims(mcmcr_example)
```

---

nsams

*Number of MCMC Samples*

---

### Description

The product of the number of simulations and the number of terms.

### Usage

```
nsams(x)
```

**Arguments**

x                    The MCMC object.

**Value**

A count of the total number of samples.

**Examples**

```
nsims(mcmcr_example)
```

---

nsims	<i>Number of MCMC Simulations</i>
-------	-----------------------------------

---

**Description**

The product of the number of chains and number of iterations (in each chain).

**Usage**

```
nsims(x)
```

**Arguments**

x                    The MCMC object.

**Value**

A count of the total number of simulations.

**Examples**

```
nsims(mcmcr_example)
```

---

nterms	<i>Number of Terms</i>
--------	------------------------

---

### Description

Gets the number of terms for an object.

### Usage

```
nterms(x, ...)  
  
## Default S3 method:  
nterms(x, ...)  
  
## S3 method for class 'mcmcarray'  
nterms(x, ...)  
  
## S3 method for class 'mcmcr'  
nterms(x, ...)  
  
## S3 method for class 'mcmcrs'  
nterms(x, ...)
```

### Arguments

x	The object.
...	Not used.

### Value

A count of the number of terms.

### Methods (by class)

- default: Number of terms for an object
- mcmcarray: Number of terms for an mcmcarray object
- mcmcr: Number of terms for an mcmcr object
- mcmcrs: Number of terms for an mcmcrs object

### Examples

```
nterms(mcmcr_example)
```

---

parameters

*Parameter Names*

---

## Description

Gets or sets the parameter names for an object.

## Usage

```
parameters(x, ...)  
  
parameters(x) <- value  
  
set_parameters(x, parameters)  
  
## S3 method for class 'term'  
parameters(x, scalar_only = FALSE, terms = FALSE, ...)  
  
## S3 method for class 'mcmc'  
parameters(x, scalar_only = FALSE, terms = FALSE, ...)  
  
## S3 method for class 'mcmc.list'  
parameters(x, scalar_only = FALSE, terms = FALSE,  
  ...)  
  
## S3 method for class 'mcmcr'  
parameters(x, scalar_only = FALSE, terms = FALSE, ...)  
  
## S3 method for class 'mcmcrs'  
parameters(x, scalar_only = FALSE, terms = FALSE, ...)
```

## Arguments

x	An MCMC object.
...	Not used.
value	A character vector of the new parameter names.
parameters	A character vector of the new parameter names.
scalar_only	A flag indicating whether to only get the names of parameters with one term.
terms	A flag indicating whether to return the parameter name for each term.

## Value

A character vector of the parameter names.



**Methods (by class)**

- `term`: Parameter names for a term vector
- `mcmc`: Parameter names for an `mcmc` object
- `mcmc.list`: Parameter names for an `mcmc.list` object
- `mcmcr`: Parameter names for an `mcmcr` object
- `mcmcrs`: Parameter names for an `mcmcrs` object

**Examples**

```
parameters(mcmcr_example)
parameters(mcmcr_example) <- c("gamma", "theta", "tau")
parameters(mcmcr_example)
parameters(mcmcr_example, scalar_only = TRUE)
parameters(mcmcr_example, terms = TRUE)
```

---

pvalue

*P-Value*

---

**Description**

Calculates the p-value.

**Usage**

```
pvalue(x)
```

**Arguments**

`x` A vector of MCMC value

**Value**

A number indicating the p-value.

**Examples**

```
pvalue(as.numeric(0:10))
```

---

 rhat

*R-hat*


---

### Description

By default calculates the uncorrected, unfolded, univariate, split R-hat (potential scale reduction factor) values.

### Usage

```
rhat(x, ...)

## S3 method for class 'marray'
rhat(x, by = "all", as_df = FALSE, ...)

## S3 method for class 'mcmc'
rhat(x, by = "all", as_df = FALSE, ...)

## S3 method for class 'mcmc.list'
rhat(x, by = "all", as_df = FALSE, ...)

## S3 method for class 'mcmcarray'
rhat(x, by = "all", as_df = FALSE, ...)

## S3 method for class 'mcmcr'
rhat(x, by = "all", as_df = FALSE, ...)

## S3 method for class 'mcmcrs'
rhat(x, by = "all", as_df = FALSE, bound = FALSE,
     ...)
```

### Arguments

x	An MCMC object.
...	Unused.
by	A string indicating whether to determine by "term", "parameter" or "all".
as_df	A flag indicating whether to return the values as a data frame versus a named list.
bound	flag specifying whether to bind mcmcrs objects by their chains before calculating rhat.

### Value

The rhat value(s).

**Methods (by class)**

- `mccarray`: R-hat for an `mccarray` object
- `mcmc`: R-hat for an `mcmc` object
- `mcmc.list`: R-hat for an `mcmc.list` object
- `mcmcarray`: R-hat for an `mcmcarray` object
- `mcmcr`: R-hat for an `mcmcr` object
- `mcmcrs`: R-hat for an `mcmcrs` object

**References**

Gelman, A., and Rubin, D.B. 1992. Inference from Iterative Simulation Using Multiple Sequences. *Statistical Science* 7(4): 457–472.

**Examples**

```
rhat(mcmcr_example)
rhat(mcmcr_example, by = "parameter")
rhat(mcmcr_example, by = "term")
rhat(mcmcr_example, by = "term", as_df = TRUE)
rhat(mcmcrs(mcmcr_example, mcmcr_example))
rhat(mcmcrs(mcmcr_example, mcmcr_example), bound = TRUE)
```

---

 sort

*Sort an MCMC Object*


---

**Description**

Sorts an MCMC object by its parameter names.

**Usage**

```
## S3 method for class 'mcmc'
sort(x, ...)

## S3 method for class 'mcmc.list'
sort(x, ...)

## S3 method for class 'mcmcr'
sort(x, ...)

## S3 method for class 'mcmcrs'
sort(x, ...)
```

**Arguments**

<code>x</code>	The MCMC object to sort
<code>...</code>	Unused

**Methods (by class)**

- mcmc: Sort an mcmc object
- mcmc.list: Sort an mcmc.list object
- mcmcr: Sort an mcmcr object
- mcmcrs: Sort an mcmcrs object

**Examples**

```
parameters(mcmcr_example)
mcmcr_example <- subset(mcmcr_example, parameters = c("beta", "alpha"))
parameters(mcmcr_example)
mcmcr_example <- sort(mcmcr_example)
parameters(mcmcr_example)
```

---

split\_chains

*Split Chains*


---

**Description**

Splits each chain in half to double the number chains and halve the number of iterations.

**Usage**

```
split_chains(x, ...)
```

```
## S3 method for class 'mcmcarray'
split_chains(x, ...)
```

```
## S3 method for class 'mcmcr'
split_chains(x, ...)
```

**Arguments**

```
x          An MCMC object.
...        Unused.
```

**Methods (by class)**

- mcmcarray: Split chains for an mcmcarray object
- mcmcr: Split chains for an mcmcr object

**Examples**

```
split_chains(mcmcr_example)
```

---

subset	<i>Subset an MCMC Object</i>
--------	------------------------------

---

### Description

Subsets an MCMC object by its chains, iterations and/or parameters.

### Usage

```
## S3 method for class 'term'
subset(x, parameters = NULL, ...)

## S3 method for class 'mcmc'
subset(x, iterations = NULL, parameters = NULL, ...)

## S3 method for class 'mcmc.list'
subset(x, chains = NULL, iterations = NULL,
       parameters = NULL, ...)

## S3 method for class 'mcmcarray'
subset(x, chains = NULL, iterations = NULL, ...)

## S3 method for class 'mcmcr'
subset(x, chains = NULL, iterations = NULL,
       parameters = NULL, ...)

## S3 method for class 'mcmcrs'
subset(x, chains = NULL, iterations = NULL,
       parameters = NULL, ...)
```

### Arguments

x	The MCMC object to subset
parameters	A character vector (or NULL) of the parameters to subset by.
...	Unused
iterations	An integer vector (or NULL) of the iterations to subset by.
chains	An integer vector (or NULL) of the chains to subset by.

### Methods (by class)

- term: Subset a term vector
- mcmc: Subset an mcmc object
- mcmc.list: Subset an mcmc.list object
- mcmcarray: Subset an mcmcarray object
- mcmcr: Subset an mcmcr object
- mcmcrs: Subset an mcmcrs object

**Examples**

```
subset(mcmcr_example, chains = 2L, iterations = 1:100,
       parameters = c("beta", "alpha"))
```

---

tdims	<i>Term Dimensions</i>
-------	------------------------

---

**Description**

Gets the term dimensions of an object.

**Usage**

```
tdims(x, ...)

## Default S3 method:
tdims(x, ...)

## S3 method for class 'term'
tdims(x, ...)
```

**Arguments**

x	The object
...	Unused.

**Methods (by class)**

- default: Term dimensions for an object
- term: Term dimensions for a term vector

**Examples**

```
tdims(mcmcr_example)
```

---

term	<i>Term Vector</i>
------	--------------------

---

**Description**

A term vector is a sortable vector of the terms from an analysis.

**Examples**

```
terms <- terms(mcmcr_example)
terms <- rev(terms)
terms
sort(terms)
```

---

terms	<i>MCMC Object Terms</i>
-------	--------------------------

---

**Description**

Gets terms for an MCMC object.

**Usage**

```
## S3 method for class 'mcmc'  
terms(x, ...)  
  
## S3 method for class 'mcmc.list'  
terms(x, ...)  
  
## S3 method for class 'mcmcarray'  
terms(x, ...)  
  
## S3 method for class 'mcmcr'  
terms(x, ...)
```

**Arguments**

x	A MCMC object.
...	Unused

**Value**

A term vector.

**Methods (by class)**

- `mcmc`: Get terms for an `mcmc` object
- `mcmc.list`: Get terms for an `mcmc.list` object
- `mcmcarray`: Get terms for an `mcmcarray` object
- `mcmcr`: Get terms for an `mcmcr` object

**See Also**

`stats::terms`

**Examples**

```
terms(mcmcr_example)
```

---

thin *Thin MCMC Samples*

---

### Description

Thin an MCMC objects samples.

### Usage

```
## S3 method for class 'mcmcarray'  
thin(x, nthin = 1L, ...)  
  
## S3 method for class 'mcmcr'  
thin(x, nthin = 1L, ...)  
  
## S3 method for class 'mcmcrs'  
thin(x, nthin = 1L, ...)
```

### Arguments

x	An MCMC object
nthin	A count of the thinning rate.
...	Unused

### Value

The thinned object.

### Methods (by class)

- `mcmcarray`: Thin MCMC samples for an `mcmcarray` object
- `mcmcr`: Thin MCMC samples for an `mcmcr` object
- `mcmcrs`: Thin MCMC samples for an `mcmcrs` object

### See Also

`coda::thin`

### Examples

```
thin(mcmcr_example, nthin = 10L)
```



---

zero

*Zero MCMC Sample Values*

---

### Description

Zeros an MCMC object's sample values.

### Usage

```
zero(x, ...)  
  
## S3 method for class 'marray'  
zero(x, ...)  
  
## S3 method for class 'mcmarray'  
zero(x, ...)  
  
## S3 method for class 'mcmcr'  
zero(x, parameters = NULL, ...)
```

### Arguments

x	The MCMC object.
...	Unused
parameters	A character vector (or NULL) of the parameters to zero.

### Details

It is used for removing the effect of a random effect where the expected value is 0.

### Methods (by class)

- marray: Zero an marray object
- mcmarray: Zero an mcmarray object
- mcmcr: Zero an mcmcr object

### Examples

```
zero(mcmcr_example, parameters = "beta")
```

# Index

## \*Topic **datasets**

- mcmcr\_example, 30
- anyNA, 3, 3
- as.character.term (as.term), 9
- as.marray, 4
- as.mcmc, 4
- as.mcmc.list, 5
- as.mcmcarray, 6
- as.mcmcr, 7
- as.mcmcrs, 8
- as.mcmcrs.list (as.mcmcr), 7
- as.term, 9
  
- bind\_chains, 10
- bind\_dimensions, 11, 12
- bind\_dimensions\_n, 11, 12
- bind\_iterations, 12
- bind\_parameters, 13
  
- check\_mcmcarray, 14
- check\_mcmcr, 15
- coef, 16, 17
- collapse\_chains, 17
- combine\_dimensions, 18
- combine\_samples, 19
- combine\_samples\_n, 20
- converged, 20
  
- dims, 22
  
- esr, 21, 22, 24
- ess, 24
- estimates, 24
  
- is.marray, 25
- is.mcmcarray, 26
- is.mcmcr, 26
- is.mcmcrs, 27
- is.term, 27
  
- line, 30
  
- mcmc, 4, 5
- mcmc.list, 5
- mcmc\_aperm, 30
- mcmc\_map, 31
- mcmcarray-object, 28
- mcmcr-object, 28
- mcmcr\_example, 30
- mcmcrs, 29
- mcmcrs-object, 29
  
- nchains, 32
- ndims, 34
- niters, 34
- niters.marray (nchains), 32
- niters.mcmc (nchains), 32
- niters.mcmcarray (nchains), 32
- niters.mcmcr (nchains), 32
- niters.mcmcrs (nchains), 32
- npars, 35
- npdims, 36
- npdims.default, 36
- npdims.mcmc.list (npdims.default), 36
- npdims.mcmcr (npdims.default), 36
- nsams, 37
- nsims, 38
- nterms, 39
  
- parameters, 40
- parameters<- (parameters), 40
- pdims (npdims.default), 36
- pvalue, 41
  
- rhat, 21, 42
  
- set\_parameters (parameters), 40
- sort, 43
- split\_chains, 44
- subset, 45

tdims, 46  
term, 46  
terms, 47, 47  
thin, 48, 48  
zero, 49