

Package ‘waspr’

September 11, 2023

Type Package

Title Wasserstein Barycenters of Subset Posteriors

Version 1.0.1

Description Functions to compute Wasserstein barycenters of subset posteriors using the swapping algorithm developed by Puccetti, Rüschen Dorf and Vanduffel (2020) <[doi:10.1016/j.jmaa.2017.02.003](https://doi.org/10.1016/j.jmaa.2017.02.003)>. The Wasserstein barycenter is a geometric approach for combining subset posteriors. It allows for parallel and distributed computation of the posterior in case of complex models and/or big datasets, thereby increasing computational speed tremendously.

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URL <https://github.com/joliencremers/waspr>

BugReports <https://github.com/joliencremers/waspr/issues>

Encoding UTF-8

LazyData true

Imports Rcpp (>= 1.0.4.6)

LinkingTo BH, Rcpp, RcppArmadillo,

RoxygenNote 7.2.3

Suggests knitr, rmarkdown, testthat, spelling

VignetteBuilder knitr

Language en-US

Biarch true

Depends R (>= 3.5.0)

NeedsCompilation yes

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Repository CRAN

Date/Publication 2023-09-11 14:00:03 UTC

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combine	<i>Combine output of the swapping algorithm</i>
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Description

This (non-exported) function combines the output from the swapping algorithm (Puccetti, Rüschen-dorf and Vanduffel, 2020).

Usage

```
combine(x)
```

Arguments

x a three dimensional array (rows = subsets, columns = par, slices = samples) containing posterior samples for all subsets

Value

A wasp object, which can be further analyzed using the associated function [summary.wasp](#).

Source

Puccetti, G., Rüschen-dorf, L. & Vanduffel, S. (2020). On the computation of Wasserstein barycenters, *Journal of Multivariate Analysis*, 176.

hpd_est	<i>Compute the 95 percent Highest Posterior Density interval</i>
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Description

Compute the 95 percent Highest Posterior Density interval

Usage

```
hpd_est(x)
```

Arguments

x a numeric vector

Value

A vector containing the lower and upper bound of the 96 Posterior Density interval of a numeric vector as computed by the methods from Venter (1967).

Source

Venter, J.H. (1967). On estimation of the mode, *Annals of Mathematical Statistics*, 38(5), 1446-1455.

Examples

```
library(waspr)
hpd_est(pois_logistic[1,1,])
```

mode_est	<i>Compute the mode</i>
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Description

Compute the mode

Usage

```
mode_est(x)
```

Arguments

x a numeric vector

Value

The mode of a numeric vector as computed by the methods from Venter (1967).

Source

Venter, J.H. (1967). On estimation of the mode, *Annals of Mathematical Statistics*, 38(5), 1446-1455.

Examples

```
library(waspr)
mode_est(pois_logistic[1,1,])
```

pois_logistic	<i>pois_logistic</i>
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Description

A set of mcmc samples from 8 subposteriors from the analysis of a joint model with a logistic and poisson outcome variable.

Usage

```
pois_logistic
```

Format

An array with 3 dimensions of which the first represents the subposteriors (size = 8), the second represents the parameters (size = 8) and the third represents the amount of mcmc samples (size = 450).

print.wasp	<i>Print posterior summaries for the Wasserstein barycenter of subset posteriors</i>
------------	--

Description

Prints selected output from a Bayesian circular mixed-effects model.

Usage

```
## S3 method for class 'wasp'
print(x, ...)
```

Arguments

x a wasp object obtained from the function `wasp()`.
... further arguments passed to or from other methods.

Value

A print of posterior summaries for the Wasserstein barycenter of subset posteriors

Examples

```
library(waspr)
out <- wasp(pois_logistic,
            par.names = c("beta_s", "alpha_l", "beta_l",
                          "baseline_sigma", "baseline_mu",
                          "correlation", "sigma_s", "sigma_l"))
print(out)
```

summary	<i>Posterior summaries for the Wasserstein barycenter of subset posteriors</i>
---------	--

Description

summary gives a posterior summary (mean, mode, sd, HPD)

Usage

```
summary(x)
```

Arguments

x a wasp object.

Details

the method [summary.wasp](#) has its own help page.

Examples

```
library(waspr)
```

summary.wasp	<i>Posterior summaries for the Wasserstein barycenter of subset posteriors</i>
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Description

Outputs and prints posterior summary statistics (mean, mode, sd, 95 Posterior Density interval)

Usage

```
## S3 method for class 'wasp'
summary(x)
```

Arguments

x a wasp object obtained from the function wasp().

Value

Posterior summary statistics (mean, mode, sd, 95 all the Wasserstein barycenter of subset posteriors of all parameters in the model).

Examples

```
library(waspr)
out <- wasp(pois_logistic,
  par.names = c("beta_s", "alpha_l", "beta_l",
    "baseline_sigma", "baseline_mu",
    "correlation", "sigma_s", "sigma_l"))
summary(out)
```

swap_rcpp	<i>The swapping algorithm for computing Wasserstein barycenters</i>
-----------	---

Description

The swapping algorithm for computing Wasserstein barycenters

Usage

```
swap_rcpp(samples, acc = 0.001, iter = 10L, out = FALSE)
```

Arguments

samples	A cube containing samples for all subset posteriors (rows = subsets, columns = par, slices = samples)
acc	accuracy
iter	maximum number of iterations of the algorithm
out	boolean indicating whether output for each iteration should be displayed (default = false)

Value

a three dimensional array (rows = subsets, columns = par, slices = samples) containing output from the swapping algorithm.

wasp	<i>Compute Wasserstein barycenters of subset posteriors</i>
------	---

Description

This function computes Wasserstein Barycenters of subset posteriors and gives posterior summaries for the full posterior.

Usage

```
wasp(mcmc, par.names = NULL, acc = 0.001, iter = 10, out = FALSE)
```

Arguments

mcmc	a three dimensional array (rows = number of subset posteriors, columns = number of parameters of the posterior distribution, slices = samples number of samples for each subset posterior) containing posterior samples for all subsets
par.names	optional character vector with parameter names
acc	accuracy of the swapping algorithm (default = 0.001)
iter	maximum number of iterations of the swapping algorithm (default = 10)
out	boolean indicating whether output for each iteration of the swapping algorithm should be displayed (default = false)

Details

The swapping algorithm developed by Puccetti, Rüschemdorf and Vanduffel (2020) is used to compute Wasserstein barycenters of subset posteriors.

Value

A wasp object, which can be further analyzed using the associated function `summary.wasp`.

A wasp object contains the following elements (some elements are not returned if not applicable)

`barycenter` A matrix of posterior samples (rows) for all parameters (columns) of the full posterior obtained by the swapping algorithm.

`raw` An array (`dim = c(subsets, parameters, samples)`) containing the raw output from the swapping algorithm.

`call` The call to the `wasp()` function.

`subsets` The amount of subset posteriors in mcmc.

`parameters` The amount of parameters in mcmc.

`samples` The amount of posterior samples for each subset posterior in mcmc.

`acc` Accuracy of the swapping algorithm, default = 0.001.

`iter` Maximum amount of iterations for the swapping algorithm, default = 10.

Source

Puccetti, G., Rüschemdorf, L. & Vanduffel, S. (2020). On the computation of Wasserstein barycenters, *Journal of Multivariate Analysis*, 176.

Examples

```
library(waspr)
out <- wasp(pois_logistic,
            par.names = c("beta_s", "alpha_1", "beta_1",
                          "baseline_sigma", "baseline_mu",
                          "correlation", "sigma_s", "sigma_1"))
summary(out)
```


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