# Package 'familial'

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Type Package

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Title Familial Inference

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<b>Description</b> Provides functionality for testing familial hypotheses. Currently supports tests of the Huber family of centers. Testing is carried out using the Bayesian bootstrap. One- and two-sample tests are supported, as are directional tests. Methods for visualizing output are provided.
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BugReports https://github.com/ryan-thompson/familial/issues
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R topics documented:
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## Description

Performs a Bayesian bootstrap for a statistic defined via a suitable function.

## Usage

```
bayes.boot(x, fun, nboot = 1000, cluster = NULL, ...)
```

## Arguments

X	a numeric vector to be passed as the first argument to fun
fun	the function to bootstrap; must accept data $\boldsymbol{x}$ and weights $\boldsymbol{w}$ (in that order), and return a data frame
nboot	the number of bootstraps to perform
cluster	an optional cluster for running bootstraps in parallel; must be set up using parallel::makeCluster
	any other arguments for fun

#### Value

An object of class bayes.boot; a data frame with the following columns:

```
boot.id the bootstrap iteration index
... any columns returned by fun
```

## Author(s)

Ryan Thompson <ryan.thompson@monash.edu>

## **Examples**

```
set.seed(1)
# Bootstrap
boot <- bayes.boot(MASS::galaxies, fun = fit.family, nboot = 100)
head(boot)</pre>
```

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center.test Center test

## Description

Performs a one- or two-sample test for a family of centers.

## Usage

```
center.test(
    x,
    y = NULL,
    family = "huber",
    alternative = c("two.sided", "less", "greater"),
    mu = 0,
    paired = FALSE,
    nboot = 1000,
    loss = NULL,
    cluster = NULL,
    ...
)
```

#### **Arguments**

X	a numeric vector of data
У	an optional numeric vector of data
family	the family of centers; currently only allows 'huber' for Huber family
alternative	the form of the alternative hypothesis; must be one of 'two.sided' (default), 'greater', or 'less'
mu	the null value of the center for a one-sample test, or the null value of the center of differences for a paired two-sample test, or the null value of the difference in centers for an independent two-sample test; can be an interval
paired	a logical indicating whether to treat x and y as paired
nboot	the number of bootstraps to perform
loss	an optional $c\times 2$ matrix of losses incurred from an incorrect decision, where c is the number of candidate choices (typically c=3: H0, H1, or indeterminate)
cluster	an optional cluster for running bootstraps in parallel; must be set up using parallel::makeCluster
	any other arguments

#### **Details**

Uses the Bayesian bootstrap to compute posterior probabilities for the hypotheses  $H_0: \mu(\lambda) = \mu_0$  for some  $\lambda \in \Lambda$  vs.  $H_1: \mu(\lambda) \neq \mu_0$  for all  $\lambda \in \Lambda$ , where  $\{\mu(\lambda): \lambda \in \Lambda\}$  is a family of centers. The default loss matrix results in a decision whenever the posterior probability for one of the hypotheses is greater than 0.95 and otherwise is indeterminate.

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#### Value

An object of class center.test; a list with the following components:

expected.loss the expected loss, calculated by post-multiplying loss with prob

decision the optimal decision given the expected loss

loss the loss matrix

prob the posterior probabilities of the null and alternative

boot the bootstrap output from bayes.boot

x the x that was supplied
y the y that was supplied
mu the mu that was supplied
family the family that was supplied

#### Author(s)

Ryan Thompson < ryan.thompson@monash.edu>

#### **Examples**

```
set.seed(1)
# Familial test using Huber family with point null
test <- center.test(MASS::galaxies, mu = 21000, nboot = 100)
print(test)
plot(test)
# Familial test using Huber family with interval null
test <- center.test(MASS::galaxies, mu = c(20500, 21500), nboot = 100)
print(test)
# Familial test in parallel
cl <- parallel::makeCluster(2)
test <- center.test(MASS::galaxies, mu = c(20500, 21500), nboot = 100, cluster = cl)
parallel::stopCluster(cl)
print(test)</pre>
```

fit.family

Fit family

#### Description

Fits a family of centers.

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#### Usage

```
fit.family(
   x,
   w = rep(1, length(x)),
   family = "huber",
   spread.fun = weighted.mad,
   eps = .Machine$double.eps
)
```

#### **Arguments**

x a numeric vector of dataw a numeric vector of weights

family the location family; currently only allows 'huber' for Huber family

spread. fun a function used for the spread of x in the Huber family; ensures that the results

are comparable across variables

eps a numerical tolerance parameter

#### Value

An object of class fit.family; a data frame with the following columns:

mu.hat the fitted values

lambda the thresholding parameter

#### Author(s)

Ryan Thompson <ryan.thompson@monash.edu>

#### **Examples**

```
fit <- fit.family(MASS::galaxies)
plot(fit)</pre>
```

plot.center.test

Plot function for center.test object

## Description

Plot the posterior distribution for the family of centers using a functional box plot.

#### Usage

```
## S3 method for class 'center.test' plot(x, band = c(0.5, 0.75, 0.95), ninterp = 25, ...)
```

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#### **Arguments**

x an object of class center.test

band a vector of band limits for the functional box plot

ninterp the number of interpolation points for the functional box plot; more points lead

to finer resolution of the plot at the expense of additional computation

... any other arguments

#### Value

A plot of the posterior distribution.

#### Author(s)

Ryan Thompson <ryan.thompson@monash.edu>

plot.fit.family

Plot function for fit.family object

#### **Description**

Plot a fitted family.

## Usage

```
## S3 method for class 'fit.family'
plot(x, y = NULL, ...)
```

#### Arguments

x an object of class fit.family y an object of class fit.family

... any other arguments

#### Value

A plot of the fitted family.

## Author(s)

Ryan Thompson <ryan.thompson@monash.edu>

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print.center.test

Print function for center.test object

## Description

Print objects of class center.test.

#### Usage

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

## **Arguments**

x an object of class center.test

... any other arguments

#### Value

The argument x.

#### Author(s)

Ryan Thompson <ryan.thompson@monash.edu>

rudirichlet

Uniform Dirichlet distribution

## Description

Random number generation for the uniform Dirichlet distribution (having all concentration parameters set to one).

#### Usage

```
rudirichlet(n, d)
```

#### **Arguments**

n the number of observations
d the number of dimensions

#### Value

A matrix; each row is a random draw and each column is a dimension.

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#### Author(s)

Ryan Thompson < ryan.thompson@monash.edu>

weighted

Weighted statistics

## Description

Assorted weighted statistics unavailable in base R

## Usage

```
weighted.median(x, w)
weighted.mad(x, w)
```

## **Arguments**

x a numeric vector of dataw a numeric vector of weights

#### Value

A length-one numeric vector.

## Author(s)

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