

# Package ‘BBcor’

October 12, 2022

**Type** Package

**Title** Bayesian Bootstrapping Correlations

**Version** 1.0.3

**Description** Efficiently draw samples from the posterior distribution of various correlation coefficients with the Bayesian bootstrap described in Rubin (1981) <[doi:10.1214/aos/1176345338](https://doi.org/10.1214/aos/1176345338)>. There are six correlation coefficients, including Pearson, Kendall, Spearman, Gaussian Rank, Blomqvist, and polychoric.

**Depends** R (>= 4.0.0)

**License** GPL-2

**Imports** parallel, pbapply (>= 1.4-2), psych (>= 1.9.12.31), wdm (>= 0.2.1), ggplot2 (>= 3.3.4), stats, utils, methods, bayeslincom (>= 1.2.0), Rdpack

**Suggests** BGGM

**RdMacros** Rdpack

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 7.1.1

**NeedsCompilation** no

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**Repository** CRAN

**Date/Publication** 2021-10-14 11:20:01 UTC

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bbcor	<i>Bayesian Bootstrapping Correlations</i>
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## Description

Efficiently draws samples from the posterior distribution of various correlation coefficients

## Usage

```
bbcor(x, method = "pearson", iter = 5000, cores = 2)
```

## Arguments

<code>x</code>	A matrix of dimensions $n$ by $p$
<code>method</code>	Character string. Which correlation coefficient should be computed. One of "pearson" (default), "kendall", "spearman", "polychoric", "gaussian_rank", or "blomqvist" (i.e., median correlation).
<code>iter</code>	Numeric. How many posterior samples (defaults to 5000) ?
<code>cores</code>	Numeric. How many cores for parallel computing (defaults to 2)?

## Value

- `cor_mean`: A matrix including the posterior mean
- `samps`: An array of dimensions  $p$  by  $b$  by `iter` that includes the sampled correlation matrices.

## Note

NAs are removed.

## Examples

```
Y <- mtcars[,1:2]
bb_samps <- bbcor(Y, method = "spearman")
```

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compare	<i>Compare Bayesian bootstrapped correlations</i>
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**Description**

See [lin\\_comb](#)

**Usage**

```
compare(lin_comb, obj, cred = 0.9, rope = NULL, contrast = NULL)
```

**Arguments**

lin_comb	A string specifying a linear combination of variables, or a list of variable names if using contrast.
obj	An object of class BGGM, bbcor, or a data.frame of posterior samples.
cred	The level for which a credible interval should be computed.
rope	Specify a ROPE. Optional.
contrast	A contrast matrix specifying which combinations to test. Optional.

**Value**

An object of class bayeslincom

**Examples**

```
Y <- mtcars[, 1:3]
bb <- bbcor(Y)
bb_compare <- compare("mpg--cyl > mpg--disp",
                      obj = bb,
                      cred = 0.90,
                      rope = c(-0.1, 0.1))

bb_compare
```

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cor_2_pcor	<i>Correlation to Partial Correlation</i>
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**Description**

Convert correlations into the corresponding partial correlations.

**Usage**

```
cor_2_pcor(x, ...)
```

**Arguments**

x	An object of class bbcor
...	Currently ignored
	<ul style="list-style-type: none"> <li>• pcor_mean: A matrix including the posterior mean.</li> <li>• sampls: An array of dimensions p by b by iter that includes the sampled partial correlation matrices.</li> </ul>

**Examples**

```
Y <- mtcars[,1:3]

fit <- bbcor(Y, method = "spearman")

cor_2_pcor(fit)
```

---

plot.bayeslincom      *Plot comparisons from compare*

---

**Description**

See [plot.bayeslincom](#)

**Usage**

```
## S3 method for class 'bayeslincom'
plot(
  x,
  point_col = "black",
  hist_col = "black",
  hist_fill = "gray",
  bar_col = "steelblue",
  bins = 30,
  display_comb_strings = TRUE,
  ...
)
```

**Arguments**

x	An object of class bayeslincom
point_col	Color for point indicating mean of posterior
hist_col	Color for histogram edges
hist_fill	Color for histogram bars
bar_col	Color of bar for credible interval
bins	Number of bins

```

display_comb_strings      If TRUE, displays full strings for combinations in ggplot facets when there is
                          more than one combination in x
...                       Currently ignored

```

**Value**

An object of class `ggplot`

**Examples**

```

Y <- mtcars[, 1:3]
bb <- bbcor(Y)
bb_compare <- compare("mpg--cyl > mpg--disp",
                      obj = bb,
                      cred = 0.90,
                      rope = c(-0.1, 0.1))

plot(bb_compare)

```

---

plot.bbcor

*Plot bbcor point estimates and intervals*


---

**Description**

Plot `bbcor` point estimates and intervals

**Usage**

```

## S3 method for class 'bbcor'
plot(x, ci = 0.9, point_col = "red", bar_col = "black", ...)

```

**Arguments**

```

x           An object of class bbcor
ci          Width of credible interval. Defaults to 0.9.
point_col   Color for point indicating mean of posterior
bar_col     Color of bar for credible interval
...        Currently ignored

```

**Value**

An object of class `ggplot`

**Examples**

```

Y <- mtcars[, 1:5]
bb <- bbcor(Y)
plot(bb)

```

posterior\_samples      *Extract Posterior Samples*

---

**Description**

Extract Posterior Samples

**Usage**

```
posterior_samples(object, ...)
```

**Arguments**

object	An object of class bbcor
...	Currently ignored

**Value**

A data frame including the posterior samples

**Examples**

```
Y <- mtcars[,1:5]

bb_samps <- bbcor(Y, method = "spearman")

# correlations
posterior_samples(bb_samps)

# partial correlations
posterior_samples(cor_2_pcor(bb_samps))
```

---

print.bayeslincom      *Print formatted summary of a bayeslincom object*

---

**Description**

Print formatted summary of a bayeslincom object

**Usage**

```
## S3 method for class 'bayeslincom'
print(x, decimals = 2, ...)
```

**Arguments**

x                    An object of class bayeslincom  
decimals            The number of decimals points to which estimates should be rounded  
...                   Other arguments to be passed to print

**Value**

A formatted summary of posterior samples

---

print.bbcor                    *Print bbcor Objects*

---

**Description**

Print the correlation or partial correlation matrix

**Usage**

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

**Arguments**

x                    An object of class bbcor  
...                   Currently ignored

---

srol2021                    *Data on the social consequences of COVID-19 conspiracy beliefs*

---

**Description**

Data from Study 1 in (Årol et al. 2021) examining effects of prejudice and descrimination on COVID-19 conspiracy beliefs

**Usage**

```
data("srol2021")
```

## Format

A data frame with 501 rows and 24 variables

- id: participant id
- gender: participants' indicated gender (1 = "male", 2 = "female")
- age: participants' indicated age
- education: participants' indicated highest attained education level(1 = "elementary education", 2 = "high school without diploma", 3 = "high school with diploma", 4 = "undergraduate college degree", 5 = "graduate college degree", 6 = "doctoral degree")
- combined\_covid\_conspiracy: average rating on 12 items of both generic and China-specific COVID-19 conspiracy beliefs
- china\_covid\_conspiracy: average rating on 4 items of China-specific COVID-19 conspiracy beliefs
- generic\_covid\_conspiracy: average rating on 8 items of generic COVID-19 conspiracy beliefs
- generic\_covid\_conspiracy\_wo\_hoax: average rating on 7 items of generic COVID-19 conspiracy beliefs (without the hoax theory item)
- combined\_covid\_conspiracy\_wo\_hoax: average rating on 11 items of both generic and China-specific COVID-19 conspiracy beliefs (without the hoax theory item)
- neg\_feelings\_italy: score on a feeling thermometer (higher score = more negative feelings) toward Italian people/ 0-100
- neg\_feelings\_china: score on a feeling thermometer (higher score = more negative feelings) toward Chinese people/ 0-100
- neg\_feelings\_roma: score on a feeling thermometer (higher score = more negative feelings) toward Roma people/ 0-100
- social\_distance\_italy: average rating on three items of social distance toward Italian people
- social\_distance\_china: average rating on three items of social distance toward Chinese people
- social\_distance\_roma: average rating on three items of social distance toward Roma people
- discrimination\_italy: rating on one discrimination item for Italian people
- discrimination\_china: rating on one discrimination item for Chinese people
- discrimination\_roma: rating on one discrimination item for Roma people
- italy\_composite: composite average of 5 z-scores (feeling thermometer, 3 social distance items, and discrimination) for Italian people
- china\_composite: composite average of 5 z-scores (feeling thermometer, 3 social distance items, and discrimination) for Chinese people
- roma\_composite: composite average of 5 z-scores (feeling thermometer, 3 social distance items, and discrimination) for Roma people
- information\_exposure: average rating on the 3 items of exposure to information about COVID-19 pandemic
- anxiety: average rating on the 4 items related to feelings of anxiety]
- lack\_of\_control: average rating on the 4 items related to the feeling of lack of control



## Details

Further details can be found at <https://osf.io/jkab7/>

## References

Årol J, Cavojava V, MikuÅkovÃi EB (2021). "Social consequences of COVID-19 conspiracy beliefs: Evidence from two studies in Slovakia." *PsyArXiv*.

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summary.bbcor

*Summarize posterior samples from bbcor object*

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## Description

Summarize posterior samples from bbcor object

## Usage

```
## S3 method for class 'bbc'or'  
summary(object, ci = 0.9, decimals = 2, ...)
```

## Arguments

object	An object of class bbcor
ci	The desired credible interval
decimals	The number of decimals points to which estimates should be rounded
...	Currently ignored

## Value

A data.frame summarizing the relations

## Examples

```
Y <- mtcars[, 1:5]  
bb <- bbcor(Y, method = "spearman")  
  
summary(bb)
```

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