# Package 'lancor'

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tle Statistical Inference via Lancaster Correlation		
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Maintainer Bernhard Klar <bernhard.klar@kit.edu></bernhard.klar@kit.edu>		
Description Implementation of the methods described in Holzmann, Klar (2024) <doi:10.1111 sjos.12733="">. Lancaster correlation is a correlation coefficient which equals the absolute value of the Pearson correlation for the bivariate normal distribution, and is equal to or slightly less than the maximum correlation coefficient</doi:10.1111>		
for a variety of bivariate distributions. Rank and moment-based estimators and corresponding confidence intervals are implemented, as well as independence tests based on these statistics.		
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Author Bernhard Klar [aut, cre] (ORCID:		
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Lancaster correlation

#### **Description**

Computes the Lancaster correlation coefficient.

#### Usage

#### **Arguments**

x a numeric vector, or a matrix or data frame with two columns.

y NULL (default) or a vector with same length as x.

type a character string indicating which lancaster correlation is to be computed. One of "rank" (default), or "linear": can be abbreviated.

#### **Details**

Let  $F_X$  and  $F_Y$  be the distribution functions of X and Y, and define

$$X^* = \Phi^{-1}(F_X(X)), \quad Y^* = \Phi^{-1}(F_Y(Y)),$$

where  $\Phi^{-1}$  is the standard normal quantile function. Furthermore for X and Y with finite fourth moment, let

$$\tilde{X} = (X - \mathbb{E}(X))/\operatorname{sd}(X), \quad \tilde{Y} = (Y - \mathbb{E}(Y))/\operatorname{sd}(Y).$$

Then

$$\rho_L(X,Y) = \max\{|\operatorname{Cor}_{\operatorname{Pearson}}(X^*,Y^*)|, \ |\operatorname{Cor}_{\operatorname{Pearson}}((X^*)^2,(Y^*)^2)|\}$$

and

$$\rho_{L,1}(X,Y) = \max\{|\operatorname{Cor}_{\operatorname{Pearson}}(X,Y)|, |\operatorname{Cor}_{\operatorname{Pearson}}((\tilde{X})^2, (\tilde{Y})^2)|\}$$

are called the Lancaster correlation coefficient and the linear Lancaster correlation coefficient, respectively. Two estimation methods are supported:

- Linear estimator for  $\rho_{L,1}$  (type = "linear"): Consider  $\rho_{L1} = \operatorname{Cor}_{\operatorname{Pearson}}(X,Y)$  and  $\rho_{L2} = \operatorname{Cor}_{\operatorname{Pearson}}((\tilde{X})^2,(\tilde{Y})^2)$ . Let  $\hat{\rho}_{L1}$  be the sample Pearson correlation and  $\hat{\rho}_{L2}$  the empirical correlation of the squares of the empirically standardized observations, and set  $\hat{\rho}_{L,1} = \max\{|\hat{\rho}_{L1}|, |\hat{\rho}_{L2}|\}$ .
- Rank-based estimator for  $\rho_L$  (type = "rank"): Consider  $\rho_{R1} = \operatorname{Cor}_{\operatorname{Pearson}}(X^*, Y^*)$  and  $\rho_{R2} = \operatorname{Cor}_{\operatorname{Pearson}}((X^*)^2, (Y^*)^2)$ . Let  $Q_i$  and  $R_i$  be the ranks of  $X_i$  and  $Y_i$ , within  $X_1, ..., X_n$  or  $Y_1, ..., Y_n$  respectively. Define

$$\hat{\rho}_{R1} = \frac{1}{n \, s_a^2} \sum_{j=1}^n a(Q_j) \, a(R_j),$$

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$$\hat{\rho}_{R2} = \frac{1}{n s_b^2} \sum_{j=1}^{n} (b(Q_j) - \bar{b}) (b(R_j) - \bar{b}),$$

where the scores are, for j = 1, ..., n,

$$a(j) = \Phi^{-1}\left(\frac{j}{n+1}\right), \quad b(j) = a(j)^2,$$

$$\bar{b} = \frac{1}{n} \sum_{j=1}^{n} b(j), \quad s_a^2 = \frac{1}{n} \sum_{j=1}^{n} (a(j) - \bar{a})^2, \quad s_b^2 = \frac{1}{n} \sum_{j=1}^{n} (b(j) - \bar{b})^2.$$

Finally, the rank-based Lancaster correlation is

$$\hat{\rho}_L = \max\{|\hat{\rho}_{R1}|, |\hat{\rho}_{R2}|\}.$$

#### Value

the sample Lancaster correlation.

#### Author(s)

Hajo Holzmann, Bernhard Klar

#### References

Holzmann, Klar (2024). "Lancester correlation - a new dependence measure linked to maximum correlation". doi:10.1111/sjos.12733

#### See Also

```
lcor.comp, lcor.ci, lcor.test
```

## **Examples**

```
Sigma <- matrix(c(1,0.1,0.1,1), ncol=2)
R <- chol(Sigma)
n <- 1000
x <- matrix(rnorm(n*2), n)
lcor(x, type = "rank")
lcor(x, type = "linear")

x <- matrix(rnorm(n*2), n)
nu <- 2
y <- x / sqrt(rchisq(n, nu)/nu)
cor(y[,1], y[,2], method = "spearman")
lcor(y, type = "rank")</pre>
```

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lcor.ci

Confidence intervals for the Lancaster correlation coefficient

#### **Description**

Computes confidence intervals for the Lancaster correlation coefficient. Lancaster correlation is a bivariate measures of dependence.

#### Usage

```
lcor.ci(
   x,
   y = NULL,
   conf.level = 0.95,
   type = c("rank", "linear"),
   con = TRUE,
   R = 1000,
   method = c("plugin", "boot", "pretest")
)
```

#### Arguments

x a numeric vector, or a matrix or data frame with two columns.

y NULL (default) or a vector with same length as x.

conf. level confidence level of the interval.

type a character string indicating which lancaster correlation is to be computed. One

of "rank" (default), or "linear": can be abbreviated.

con logical; if TRUE (default), conservative asymptotic confidence intervals are

computed.

R number of bootstrap replications.

method a character string indicating how the asymptotic covariance matrix is computed

if type ="linear". One of "plugin" (default), "boot" or "symmetric": can be

abbreviated.

## **Details**

Computes asymptotic and bootstrap-based confidence intervals for the (linear) Lancaster correlation coefficient  $\rho_L$  ( $\rho_{L,1}$ ). For more details see lcor.

Asymptotic confidence intervals are derived under two cases (analogously for  $\rho_L$ ; see Holzmann and Klar (2024)):

Case 1: If  $|\rho_{L1}| \neq |\rho_{L2}|$ , the  $1 - \alpha$  asymptotic interval is

$$\left[\max\{\hat{\rho}_{L,1} - z_{1-\alpha/2} s/\sqrt{n}, 0\}, \min\{\hat{\rho}_{L,1} + z_{1-\alpha/2} s/\sqrt{n}, 1\}\right],$$

where  $z_{1-\alpha/2}$  is the standard normal quantile and s is an estimator of the corresponding standard deviation.

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Case 2: If  $|\rho_{L1}| = |\rho_{L2}| = a > 0$ , let  $\tau$  denote the correlation between the two components and let  $q_{1-\alpha/2}$  be the  $1-\alpha/2$  quantile of the asymptotic distribution of  $\sqrt{n}(\hat{\rho}_{L,1}-a)$ . A conservative asymptotic interval is

$$\left[\max\{\hat{\rho}_{L,1} - q_{1-\alpha/2}/\sqrt{n}, 0\}, \min\{\hat{\rho}_{L,1} + z_{1-\alpha/2} s/\sqrt{n}, 1\}\right].$$

Additionally, bootstrap-based intervals can be obtained by resampling and estimating the covariance matrix of the rank or linear correlation components.

## Value

a vector containing the lower and upper limits of the confidence interval.

#### Author(s)

Hajo Holzmann, Bernhard Klar

#### References

Holzmann, Klar (2024). "Lancester correlation - a new dependence measure linked to maximum correlation". doi:10.1111/sjos.12733

#### See Also

```
lcor, lcor.comp, lcor.test
```

## **Examples**

```
n <- 1000
x <- matrix(rnorm(n*2), n)
nu <- 2
y <- x / sqrt(rchisq(n, nu)/nu) # multivariate t
lcor(y, type = "rank")
lcor.ci(y, type = "rank")</pre>
```

lcor.comp

Lancaster correlation and its components

## **Description**

Computes the Lancaster correlation coefficient and its components.

## Usage

```
lcor.comp(x, y = NULL, type = c("rank", "linear"), plot = FALSE)
```

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## **Arguments**

X	a numeric vector, or a matrix or data frame with two columns.
У	NULL (default) or a vector with same length as x.
type	a character string indicating which lancaster correlation is to be computed. One of "rank" (default), or "linear": can be abbreviated.
plot	logical; if TRUE, scatterplots of the transformed x and y values and of their squares are drawn.

## **Details**

For more details see lcor.

#### Value

a vector containing the two components rho1 and rho2 and the sample Lancaster correlation.

#### Author(s)

Hajo Holzmann, Bernhard Klar

## References

Holzmann, Klar (2024). "Lancester correlation - a new dependence measure linked to maximum correlation". doi:10.1111/sjos.12733

#### See Also

```
lcor, lcor.comp, lcor.test
```

## **Examples**

```
Sigma <- matrix(c(1,0.1,0.1,1), ncol=2)
R <- chol(Sigma)
n <- 1000
x <- matrix(rnorm(n*2), n)
nu <- 8
y <- x / sqrt(rchisq(n, nu)/nu) #multivariate t
cor(y[,1], y[,2])
lcor.comp(y, type = "linear")

x <- matrix(rnorm(n*2), n)
nu <- 2
y <- x / sqrt(rchisq(n, nu)/nu) #multivariate t
cor(y[,1], y[,2], method = "spearman")
lcor.comp(y, type = "rank", plot = TRUE)</pre>
```

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lcor.test

Lancaster correlation test

## Description

Lancaster correlation test of bivariate independence. Lancaster correlation is a bivariate measures of dependence.

## Usage

```
lcor.test(
    x,
    y = NULL,
    type = c("rank", "linear"),
    nperm = 999,
    method = c("permutation", "asymptotic", "symmetric")
)
```

#### **Arguments**

x a numeric vector, or a matrix or data frame with two columns.

y NULL (default) or a vector with same length as x

type a character string indicating which lancaster correlation is to be computed. One

of "rank" (default), or "linear": can be abbreviated.

nperm number of permutations.

method a character string indicating how the p-value is computed if type ="linear". One

of "permutation" (default), "asymptotic" or "symmetric": can be abbreviated.

#### Details

For more details on the testing procedure see  $Remark\ 2$  in Holzmann, Klar (2024).

#### Value

A list containing the following components:

lcor the value of the test statistic pval the p-value of the test

#### Author(s)

Hajo Holzmann, Bernhard Klar

#### References

Holzmann, Klar (2024). "Lancester correlation - a new dependence measure linked to maximum correlation". doi:10.1111/sjos.12733

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#### See Also

lcor, lcor.comp, lcor.ci and for for performing an ACE permutation test of independence see acepack (https://cran.r-project.org/package=acepack).

## **Examples**

```
n <- 200
x <- matrix(rnorm(n*2), n)
nu <- 2
y <- x / sqrt(rchisq(n, nu)/nu)
cor.test(y[,1], y[,2], method = "spearman")
lcor.test(y, type = "rank")</pre>
```

Sigma.est

Covariance matrix of components of Lancaster correlation coefficient

## **Description**

Estimate of covariance matrix of the two components of Lancaster correlation. Lancaster correlation is a bivariate measures of dependence.

## Usage

```
Sigma.est(xx)
```

## Arguments

XX

a matrix or data frame with two columns.

#### **Details**

For more details see the Appendix in Holzmann, Klar (2024).

## Value

the estimated covariance matrix.

#### Author(s)

Hajo Holzmann, Bernhard Klar

#### References

Holzmann, Klar (2024). "Lancester correlation - a new dependence measure linked to maximum correlation". doi:10.1111/sjos.12733

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## See Also

lcor.ci

## Examples

```
Sigma <- matrix(c(1,0.1,0.1,1), ncol=2)
R <- chol(Sigma)
n <- 1000
x <- matrix(rnorm(n*2), n)
nu <- 8
y <- x / sqrt(rchisq(n, nu)/nu) #multivariate t
Sigma.est(y)</pre>
```

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