

# Package ‘regcorr’

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**Type** Package

**Title** Regression Models of Pearson Correlation Coefficient

**Version** 0.1.0

**Description** Provides statistical tools for evaluating how covariates influence the strength of Pearson correlation coefficients between two response variables. Supports bivariate normal and bivariate binary responses, with likelihood-based inference and bootstrap-based significance testing. The methodology is based on Dufera, Liu and Xu (2023) “Regression models of Pearson correlation coefficient” <[doi:10.1080/24754269.2023.2164970](https://doi.org/10.1080/24754269.2023.2164970)>.

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**Imports** stats

**Suggests** testthat (>= 3.0.0)

**Config/testthat/edition** 3

**URL** <https://github.com/lonze-nb/regcorr>

**BugReports** <https://github.com/lonze-nb/regcorr/issues>

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**NeedsCompilation** no

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genDataBB	<i>Generate data from bivariate Bernoulli</i>
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### Description

Generate data from bivariate Bernoulli

### Usage

```
genDataBB(numSample, p, betaTrue, eta1True, eta2True, link)
```

```
genDataBB(numSample, p, betaTrue, eta1True, eta2True, link)
```

### Arguments

numSample	Sample size.
p	Number of covariates.
betaTrue	True beta vector.
eta1True	True eta1 vector.
eta2True	True eta2 vector.
link	Link function indicator ("1" = logistic; "2" = tanh).

### Value

A list containing X, Y, and rho.

### Examples

```
set.seed(123)
dat <- genDataBB(
  numSample = 20,
  p = 2,
  betaTrue = c(0.2, 0, 0),
  eta1True = c(0, 0, 0),
  eta2True = c(0, 0, 0),
  link = "1"
```

```
)  
names(dat)
```

---

genDataBN	<i>Generate data from bivariate normal</i>
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---

### Description

Simulate bivariate normal responses with a covariate-dependent Pearson correlation coefficient.

### Usage

```
genDataBN(numSample, p, betaTrue, eta1True, eta2True, link)  
genDataBN(numSample, p, betaTrue, eta1True, eta2True, link)
```

### Arguments

numSample	Number of observations.
p	Number of covariates.
betaTrue	True regression coefficients for the correlation model.
eta1True	Included for interface consistency.
eta2True	Included for interface consistency.
link	Correlation link function: "1" for logistic, "2" for tanh.

### Value

A list containing X, Y, and rho.

A list containing:

**X** Covariate matrix.

**Y** Response matrix with two columns.

**rho** True correlation coefficients.

### Examples

```
set.seed(123)  
dat <- genDataBN(  
  numSample = 20,  
  p = 2,  
  betaTrue = c(0.2, 0, 0),  
  eta1True = c(0, 0, 0),  
  eta2True = c(0, 0, 0),  
  link = "1"  
)  
names(dat)
```

```
dim(dat$X)
dim(dat$Y)
```

---

logistic

*Logistic Function*

---

### Description

Compute the logistic transformation of a numeric input.

### Usage

```
logistic(x)
```

```
logistic(x)
```

### Arguments

x                    A numeric vector.

### Value

A numeric value or vector with elements in (0,1).

The calculated logistic probability.

### Examples

```
logistic(0)
logistic(c(-1, 0, 1))
```

---

NRfitBivBernoulli

*Fit a Regression Model for Pearson Correlation Coefficient*

---

### Description

Estimate regression coefficients in a covariate-dependent Pearson correlation model for bivariate binary responses using the Newton-Raphson algorithm.

### Usage

```
NRfitBivBernoulli(Y, X, beta0, link)
```

```
NRfitBivBernoulli(Y, X, beta0, link)
```

**Arguments**

Y	n by 2 matrix, paired responses.
X	n by p matrix, covariate matrix including first column of ones.
beta0	Initial estimate of beta.
link	Indicator of link function ("1" = logistic, "2" = tanh).

**Value**

A list containing:

**betaCurrent** Estimated regression coefficients.

**numIter** Number of iterations.

**restart** Number of restarts.

A list containing betaCurrent, numIter, and restart.

**Examples**

```
set.seed(123)

dat <- genDataBB(
  numSample = 30,
  p = 1,
  betaTrue = c(0.3, 0.1),
  eta1True = c(0, 0),
  eta2True = c(0, 0),
  link = "1"
)

fit <- NRfitBivBernoulli(
  Y = dat$Y,
  X = dat$X,
  beta0 = c(0, 0),
  link = "1"
)

fit$numIter
```

**Description**

Estimate regression coefficients in a covariate-dependent Pearson correlation model for bivariate normal responses using the Newton-Raphson algorithm.

**Usage**

```
NRfitBivNormal(Y, X, betaIni, link)
```

```
NRfitBivNormal(Y, X, betaIni, link)
```

**Arguments**

Y	n by 2 matrix, paired responses.
X	n by p matrix, covariate matrix including first column of ones.
betaIni	Initial estimate of beta.
link	Indicator of link function ("1" = logistic, "2" = tanh).

**Value**

A list containing:

**betaCurrent** Estimated regression coefficients.

**numIter** Number of Newton-Raphson iterations.

**restart** Number of restarts.

A list containing betaCurrent, numIter, and restart.

**Examples**

```
set.seed(123)

dat <- genDataBN(
  numSample = 30,
  p = 1,
  betaTrue = c(0.5, 0.2),
  eta1True = c(0, 0),
  eta2True = c(0, 0),
  link = "1"
)

fit <- NRfitBivNormal(
  Y = dat$Y,
  X = dat$X,
  betaIni = c(0, 0),
  link = "1"
)

fit$numIter
```

---

`rbinary`*Generate Correlated Binary Data*

---

**Description**

Generate bivariate binary random variables with specified marginal probabilities and correlation.

**Usage**

```
rbinary(n, p, rho)
```

```
rbinary(n, p, rho)
```

**Arguments**

<code>n</code>	Number of rows.
<code>p</code>	1 by 2 mean vector of bivariate variables.
<code>rho</code>	Correlation of bivariate variables.

**Details**

The implementation follows Qaqish (2003).

**Value**

An  $n \times 2$  matrix of binary random variables.  
 $n$  by 2 matrix of generated binary variables.

**References**

Qaqish, B. F. (2003). A family of multivariate binary distributions for simulating correlated binary variables with specified marginal means and correlations. *Biometrika*, 90(2), 455–463.

**Examples**

```
set.seed(123)
y <- rbinary(
  n = 10,
  p = c(0.4, 0.5),
  rho = 0.2
)

dim(y)
```

---

subRoutineTest

*Subroutine to Test the Significance of Individual Parameters*


---

### Description

Conduct simulation studies for testing regression coefficients in correlation regression models.

### Usage

```
subRoutineTest(
  numSample,
  p,
  link,
  model,
  betaTrue,
  betaIni,
  eta1True,
  eta2True,
  numSimu,
  numBoot
)
```

### Arguments

numSample	Sample size.
p	Number of covariates.
link	Link function. "1" for logistic and "2" for tanh.
model	Model type. "1" for bivariate normal and "2" for bivariate Bernoulli.
betaTrue	True beta coefficients.
betaIni	Initial beta coefficients.
eta1True	True eta1 coefficients.
eta2True	True eta2 coefficients.
numSimu	Number of simulation replications.
numBoot	Number of bootstrap iterations.

### Value

A list containing:

**RMSE** Root mean squared error of the estimated coefficients.

**ConsistRate** Consistency rate of estimated correlations.

**power** Estimated testing power.

**Examples**

```
set.seed(123)

res <- subRoutineTest(
  numSample = 20,
  p = 1,
  link = "1",
  model = "1",
  betaTrue = c(0.2, 0.1),
  betaIni = c(0, 0),
  eta1True = c(0, 0),
  eta2True = c(0, 0),
  numSimu = 1,
  numBoot = 5
)

names(res)
```

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