

# Package: logisticRR (via r-universe)

May 14, 2026

**Type** Package

**Title** Adjusted Relative Risk from Logistic Regression

**Version** 0.3.0

**Author** Youjin Lee

**Maintainer** Youjin Lee <youjin.lee@penmedicine.upenn.edu>

**Imports** stats, nnet

**Suggests** testthat, rmarkdown, knitr

**Description** Adjusted odds ratio conditional on potential confounders can be directly obtained from logistic regression. However, those adjusted odds ratios have been widely incorrectly interpreted as a relative risk. As relative risk is often of interest in public health, we provide a simple code to return adjusted relative risks from logistic regression model under potential confounders.

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**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 7.0.0

**VignetteBuilder** knitr

**URL** <https://github.com/youjin1207/logisticRR>

**NeedsCompilation** no

**Repository** <https://cranhaven.r-universe.dev>

**Date/Publication** 2026-05-14 14:02:00 UTC

**RemoteUrl** <https://github.com/cranhaven/cranhaven.r-universe.dev>

**RemoteRef** package/logisticRR

**RemoteSha** 2c27bc84da8fecf33073272aeea9500def4c5bbc

**RemoteSubdir** logisticRR

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logisticRR	<i>Calculate adjusted relative risks</i>
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### Description

When response variable is binary and exposure variable is binary or continuous, this function derives adjusted relative risks conditional on fixed other confounders' value from logistic regression.

### Usage

```
logisticRR(
  formula,
  basecov = 0,
  fixcov = NULL,
  data,
  boot = FALSE,
  n.boot = 100
)
```

### Arguments

formula	a formula term that is passed into <code>glm()</code> having a form of response ~ terms where response is binary response vector and terms is a collection of terms connected by '+'. The first term of predictors will be used as a predictor of interest to calculate relative risks with respect to response variable.
basecov	a baseline value of exposure variable. Defaults to 0.
fixcov	a data frame of fixed value for each of adjusted confounders. If there is no confounder other than an exposure variable of interest, <code>fixcov = NULL</code> ; if <code>fixcov</code> is missing for covariates, they are all set to 0 (for numerical covariates) or first levels (for factor covariates).
data	a data frame containing response variable and all the terms used in formula.
boot	a logical value whether bootstrap samples are generated or not. Defaults to FALSE.
n.boot	if <code>boot = TRUE</code> , the number of bootstrap samples. Defaults to 100.

**Value**

fit	an object of class glm.
RR	(conditional) relative risk in response under exposure at baseline (basecov) and basecov + 1.
delta.var	estimated variance of relative risk (RR) using Delta method.
boot.rr	if boot = TRUE, a vector of RR's using bootstrap samples.
boot.var	estimated sampled variance using bootstraps if boot = TRUE.
fix.cov	a data frame of fixed value for each of adjusted confounders.

**Author(s)**

Youjin Lee

**Examples**

```
n <- 500
set.seed(1234)
X <- rbinom(n, 1, 0.3)
W <- rbinom(n, 1, 0.3);
W[sample(1:n, n/3)] = 2
Y <- rbinom(n, 1, plogis(X - W))
dat <- as.data.frame(cbind(Y, X, W))
result <- logisticRR(Y ~ X + W, basecov = 0, data = dat,
                    boot = TRUE, n.boot = 200)
```

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multinRR

*Inference on relative risk under multinomial logistic regression*


---

**Description**

Inference on relative risk under multinomial logistic regression

**Usage**

```
multinRR(
  formula,
  basecov,
  comparecov,
  fixcov = NULL,
  data,
  boot = FALSE,
  n.boot = 100
)
```

**Arguments**

formula	a formula term that is passed into <code>multinom()</code> where response should be a factor having K different levels. Every term appearing in the formula should be well-defined as a column of data. Reference response should be specified as the first level.
basecov	a baseline value of exposure variable. Defaults to 0.
comparecov	a value of exposure variable for comparison. Defaults to the first level.
fixcov	a data frame of fixed value for each of adjusted confounders. If there is no confounder other than the exposure variable of interest, <code>fixcov = NULL</code> ; if <code>fixcov</code> is missing for existing covariates, they are all set to 0 (for numerical covariates) or to the first level (for factor covariates).
data	a data frame containing response variable and all the terms used in formula.
boot	a logical value whether bootstrap samples are generated or not. Defaults to FALSE.
n.boot	if <code>boot = TRUE</code> , the number of bootstrap samples. Defaults to 100.

**Value**

fit	an object of class <code>multinom</code> .
RRR	(adjusted) relative risk ratio of K different responses compared to reference response under exposure at baseline ( <code>basecov</code> ) and <code>basecov + 1</code> .
RR	(adjusted) relative risk of K different responses under exposure at baseline ( <code>basecov</code> ) and <code>basecov + 1</code> .
delta.var	estimated variance of relative risk (RR) using Delta method.
boot.rr	if <code>boot = TRUE</code> , a vector of RR's using bootstrap samples.
boot.rrr	if <code>boot = TRUE</code> , a vector of relative risk ratio (RRR)'s using bootstrap samples.
boot.var	estimated sampled variance using bootstraps if <code>boot = TRUE</code> .
fix.cov	a data frame of fixed value for each of adjusted confounders.

**Author(s)**

Youjin Lee

**Examples**

```
n <- 500
set.seed(1234)
X <- rbinom(n, 1, 0.3)
W <- rbinom(n, 1, 0.3)
W[sample(1:n, n/3)] = 2
Y <- rbinom(n, 1, plogis(X - W))
multiY <- ifelse(X == 1, rbinom(n, 1, 0.7) + Y, rbinom(n, 1, 0.2) + Y)
print(table(multiY))
dat <- as.data.frame(cbind(multiY, X, W))
dat$W <- as.factor(dat$W)
```

```

result <- multinRR(multiY ~ W + X, basecov = 0, comparecov = 1,
data = dat, boot = TRUE)
print(apply(result$boot.rr, 2, sd)) # estimated standard errors using Delta method
print(sqrt(result$delta.var)) # estimated standard errors using bootstrap

```

multiRR

*Inference on relative risk under multinomial logistic regression***Description**

Inference on relative risk under multinomial logistic regression

**Usage**

```
multiRR(formula, basecov = 0, fixcov = NULL, data, boot = FALSE, n.boot = 100)
```

**Arguments**

formula	a formula term that is passed into <code>multinom()</code> where response should be a factor having K different levels. Every term appearing in the formula should be well-defined as a column of data. Reference response should be specified as the first level.
basecov	a baseline value of exposure variable. Defaults to 0.
fxcov	a data frame of fixed value for each of adjusted confounders. If there is no confounder other than the exposure variable of interest, <code>fxcov = NULL</code> ; if <code>fxcov</code> is missing for existing covariates, they are all set to 0 (for numerical covariates) or to the first level (for factor covariates).
data	a data frame containing response variable and all the terms used in formula.
boot	a logical value whether bootstrap samples are generated or not. Defaults to FALSE.
n.boot	if <code>boot = TRUE</code> , the number of bootstrap samples. Defaults to 100.

**Value**

fit	an object of class <code>multinom</code> .
RRR	(adjusted) relative risk ratio of K different responses compared to reference response under exposure at baseline ( <code>basecov</code> ) and <code>basecov + 1</code> .
RR	(adjusted) relative risk of K different responses under exposure at baseline ( <code>basecov</code> ) and <code>basecov + 1</code> .
delta.var	estimated variance of relative risk (RR) using Delta method.
boot.rr	if <code>boot = TRUE</code> , a vector of RR's using bootstrap samples.
boot.rrr	if <code>boot = TRUE</code> , a vector of relative risk ratio (RRR)'s using bootstrap samples.
boot.var	estimated sampled variance using bootstraps if <code>boot = TRUE</code> .
fix.cov	a data frame of fixed value for each of adjusted confounders.

**Author(s)**

Youjin Lee

**Examples**

```

n <- 500
set.seed(1234)
X <- rbinom(n, 1, 0.3)
W <- rbinom(n, 1, 0.3)
W[sample(1:n, n/3)] = 2
Y <- rbinom(n, 1, plogis(X - W))
dat <- as.data.frame(cbind(Y, X, W))
result <- multiRR(W ~ X + Y, basecov = 0, data = dat, boot = TRUE, n.boot = 100)
print(apply(result$boot.rr, 2, sd)) # estimated standard errors using Delta method
print(sqrt(result$delta.var)) # estimated standard errors using bootstrap

```

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nominalRR

---

*Calculate adjusted relative risks under nominal exposure variable*


---

**Description**

When response variable is binary and exposure variable is categorical this function derives adjusted relative risks conditional on fixed other confounders' value from logistic regression.

**Usage**

```

nominalRR(
  formula,
  basecov = NULL,
  comparecov = NULL,
  fixcov = NULL,
  data,
  boot = FALSE,
  n.boot = 100
)

```

**Arguments**

formula	a formula term that is passed into <code>glm()</code> having a form of response ~ terms where response is binary response vector and terms is a collection of terms connected by '+'. The first term of predictors will be used as a predictor of interest to calculate relative risks with respect to response variable.
basecov	a baseline value of exposure variable. Defaults to the first level.
comparecov	a value of exposure variable for comparison. Defaults to the first level.

fixcov	a data frame of fixed value for each of adjusted confounders. If there is no confounder other than an exposure variable of interest, fixcov = NULL; if fixcov is missing for covariates, they are all set to 0 (for numerical covariates) or first levels (for factor covariates).
data	a data frame containing response variable and all the terms used in formula.
boot	a logical value whether bootstrap samples are generated or not. Defaults to FALSE.
n.boot	if boot = TRUE, the number of bootstrap samples. Defaults to 100.

**Value**

fit	an object of class glm.
RR	(conditional) relative risk in response under exposure at baseline (basecov) and comparecov.
delta.var	estimated variance of relative risk (RR) using Delta method.
boot.rr	if boot = TRUE, a vector of RR's using bootstrap samples.
boot.var	estimated sampled variance using bootstraps if boot = TRUE.
fix.cov	a data frame of fixed value for each of adjusted confounders.

**Author(s)**

Youjin Lee

**Examples**

```
n <- 500
set.seed(1234)
W <- rbinom(n, 1, 0.3); W[sample(1:n, n/3)] = 2
dat <- as.data.frame(W)
dat$X <- sample( c("low", "medium", "high"), size = n, replace = TRUE)
dat$Y <- ifelse(dat$X == "low", rbinom(n, 1, plogis(W + 0.5)),
               ifelse(dat$X == "medium", rbinom(n, 1, plogis(W + 0.2)),
                      rbinom(n, 1, plogis(W - 0.4)) ))
dat$X <- as.factor(dat$X)
result <- nominalRR(Y ~ X + W, basecov = "low", comparecov = "high", data = dat,
boot = TRUE, n.boot = 200)
```

---

printmnRR

*Print adjusted relative risk using multinomial logistic regression under nominal exposure variable.*

---

**Description**

Print adjusted relative risk using multinomial logistic regression under nominal exposure variable.

**Usage**

```
printmnRR(formula, basecov, comparecov, fixcov = NULL, data)
```

**Arguments**

formula	a formula term that is passed into <code>multinom()</code> where response should be a factor having K different levels. Every term appearing in the formula should be well-defined as a column of data. Reference response should be specified as the first level.
basecov	a baseline value of exposure variable. Defaults to $\emptyset$ .
comparecov	a value of exposure variable for comparison. Defaults to the first level.
fixcov	a data frame of fixed value for each of adjusted confounders. If there is no confounder other than the exposure variable of interest, <code>fixcov = NULL</code> ; if <code>fixcov</code> is missing for existing covariates, they are all set to $\emptyset$ (for numerical covariates) or to the first level (for factor covariates).
data	a data frame containing response variable and all the terms used in formula.

**Value**

fit	an object of class <code>multinom</code> .
RRR	(adjusted) relative risk ratio of K different responses compared to reference response under exposure at baseline ( <code>basecov</code> ) and <code>basecov + 1</code> .
RR	(adjusted) relative risk of K different responses under exposure at baseline ( <code>basecov</code> ) and <code>basecov + 1</code> .
delta.var	estimated variance of relative risk (RR) using Delta method.
fix.cov	a data frame of fixed value for each of adjusted confounders.

**Author(s)**

Youjin Lee

**Examples**

```
n <- 500
set.seed(1234)
X <- rbinom(n, 1, 0.3)
W <- rbinom(n, 1, 0.3)
W[sample(1:n, n/3)] = 2
Y <- rbinom(n, 1, plogis(X - W))
multiY <- ifelse(X == 1, rbinom(n, 1, 0.7) + Y, rbinom(n, 1, 0.2) + Y)
print(table(multiY))
dat <- as.data.frame(cbind(multiY, X, W))
dat$W <- as.factor(dat$W)
result <- printmnRR(multiY ~ W + X, basecov = 0, comparecov = 1, data = dat)
```

---

printmRR	<i>Print adjusted relative risk using multinomial logistic regression under binary or ordinal exposure variable.</i>
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### Description

Print adjusted relative risk using multinomial logistic regression under binary or ordinal exposure variable.

### Usage

```
printmRR(formula, basecov = 0, fixcov = NULL, data)
```

### Arguments

formula	a formula term that is passed into <code>multinom()</code> where response should be a factor having K different levels. Every term appearing in the formula should be well-defined as a column of data. Reference response should be specified as the first level.
basecov	a baseline value of exposure variable. Defaults to 0.
fixcov	a data frame of fixed value for each of adjusted confounders. If there is no confounder other than the exposure variable of interest, <code>fixcov = NULL</code> ; if <code>fixcov</code> is missing for existing covariates, they are all set to 0 (for numerical covariates) or to the first level (for factor covariates).
data	a data frame containing response variable and all the terms used in formula.

### Value

fit	an object of class <code>multinom</code> .
RRR	(adjusted) relative risk ratio of K different responses compared to reference response under exposure at baseline ( <code>basecov</code> ) and <code>basecov + 1</code> .
RR	(adjusted) relative risk of K different responses under exposure at baseline ( <code>basecov</code> ) and <code>basecov + 1</code> .
delta.var	estimated variance of relative risk (RR) using Delta method.
fix.cov	a data frame of fixed value for each of adjusted confounders.

### Author(s)

Youjin Lee

**Examples**

```
n <- 500
set.seed(1234)
X <- rbinom(n, 1, 0.3)
W <- rbinom(n, 1, 0.3)
W[sample(1:n, n/3)] = 2
Y <- rbinom(n, 1, plogis(X - W))
dat <- as.data.frame(cbind(Y, X, W))
result <- printmRR(W ~ X + Y, basecov = 0, data = dat)
```

---

printnRR

---

*Print adjusted relative risk under nominal exposure variable.*


---

**Description**

Print adjusted relative risk under nominal exposure variable.

**Usage**

```
printnRR(formula, basecov, comparecov, fixcov = NULL, data)
```

**Arguments**

formula	a formula term that is passed into <code>glm()</code> having a form of response ~ terms where response is binary response vector and terms is a collection of terms connected by '+'. The first term of predictors will be used as a predictor of interest to calculate relative risks with respect to response variable.
basecov	a baseline value of exposure variable. Defaults to the first level.
comparecov	a value of exposure variable for comparison. Defaults to the first level.
fixcov	a data frame of fixed value for each of adjusted confounders. If there is no confounder other than an exposure variable of interest, <code>fixcov = NULL</code> ; if <code>fixcov</code> is missing for covariates, they are all set to 0 (for numerical covariates) or first levels (for factor covariates).
data	a data frame containing response variable and all the terms used in formula.

**Value**

fit	an object of class <code>glm</code> .
RR	(adjusted) relative risk in response under exposure at baseline ( <code>basecov</code> ) and <code>comparecov</code> .
delta.var	estimated variance of relative risk (RR) using Delta method.
fix.cov	a data frame of fixed value for each of adjusted confounders.

**Author(s)**

Youjin Lee

**Examples**

```
n <- 500
set.seed(1234)
W <- rbinom(n, 1, 0.3); W[sample(1:n, n/3)] = 2
dat <- as.data.frame(W)
dat$X <- sample( c("low", "medium", "high"), size = n, replace = TRUE)
dat$Y <- ifelse(dat$X == "low", rbinom(n, 1, plogis(W + 0.5)),
               ifelse(dat$X == "medium", rbinom(n, 1, plogis(W + 0.2)),
                       rbinom(n, 1, plogis(W - 0.4)) ))
dat$X <- as.factor(dat$X)
result <- printRR(Y ~ X + W, basecov = "high", comparecov = "low", data = dat)
```

printRR

*Print adjusted relative risk under binary or ordinal exposure variable.***Description**

Print adjusted relative risk under binary or ordinal exposure variable.

**Usage**

```
printRR(formula, basecov = 0, fixcov = NULL, data)
```

**Arguments**

formula	a formula term that is passed into <code>glm()</code> having a form of response ~ terms where response is binary response vector and terms is a collection of terms connected by '+'. The first term of predictors will be used as a predictor of interest to calculate relative risks with respect to response variable.
basecov	a baseline value of exposure variable. Defaults to 0.
fixcov	a data frame of fixed value for each of adjusted confounders. If there is no confounder other than an exposure variable of interest, <code>fixcov = NULL</code> ; if <code>fixcov</code> is missing for covariates, they are all set to 0 (for numerical covariates) or first levels (for factor covariates).
data	a data frame containing response variable and all the terms used in formula.

**Value**

fit	an object of class <code>glm</code> .
RR	(adjusted) relative risk in response under exposure at baseline ( <code>basecov</code> ) and <code>basecov + 1</code> .
delta.var	estimated variance of relative risk (RR) using Delta method.
fix.cov	a data frame of fixed value for each of adjusted confounders.

**Examples**

```
n <- 500
set.seed(1234)
X <- rbinom(n, 1, 0.3)
W <- rbinom(n, 1, 0.3)
W[sample(1:n, n/3)] = 2
Y <- rbinom(n, 1, plogis(X - W))
dat <- as.data.frame(cbind(Y, X, W))
result <- printRR(Y ~ X + W, basecov = 0, data = dat)
```

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