

# Package: stddiff (via r-universe)

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**Type** Package

**Title** Calculate the Standardized Difference for Numeric, Binary and Category Variables

**Version** 3.1

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**Description** Contains three main functions including `stddiff.numeric()`, `stddiff.binary()` and `stddiff.category()`. These are used to calculate the standardized difference between two groups. It is especially used to evaluate the balance between two groups before and after propensity score matching.

**License** GPL-3

**NeedsCompilation** no

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**Repository** <https://marsdu1989.r-universe.dev>

**RemoteUrl** <https://github.com/cran/stddiff>

**RemoteRef** HEAD

**RemoteSha** a3886a6b89516d719c336b118d06023f1c6e5de4

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### Description

Contains three main functions including `stddiff.numeric()`, `stddiff.binary()` and `stddiff.category()`. These are used to calculate the standardized difference between two groups. It is especially used to evaluate the balance between two groups before and after propensity score matching.

### Usage

```
stddiff.numeric(data, gcol, vcol)
stddiff.binary(data, gcol, vcol)
stddiff.category(data, gcol, vcol)
```

### Arguments

<code>data</code>	a dataframe
<code>gcol</code>	a column number of group variable in data, 0 for control group, 1 for treatment group
<code>vcol</code>	one or more column numbers of different types variables in data

### Details

`stddiff.numeric()` is used for the numeric variables. For the skewed variables, you should change to the rank using the `rank()` function before computing the "stddiff".

`stddiff.binary()` is used for the binomial variables.

`stddiff.category()` is used for the categorical variables.

Imbalance was usually defined as "stddiff" greater than 0.1 or 0.2 (which means the small effect size).

### Value

for `stddiff.numeric` function:

<code>mean.c</code>	the mean of control group
<code>sd.c</code>	the standard deviation of control group
<code>mean.t</code>	the mean of treatment group
<code>sd.t</code>	the standard deviation of treatment group
<code>missing.c</code>	the counts of missing value of control group
<code>missing.t</code>	the counts of missing value of treatment group
<code>stddiff.l</code>	the standardized difference between two groups
<code>stddiff.l</code>	the lower limit of the 95 percentage confidence interval of standardized difference

stddiff.u        the upper limit of the 95 percentage confidence interval of standardized difference

for stddiff.binary function:

p.c             the proportion of last level in the control group

p.t             the proportion of last level in the treatment group

missing.c      the counts of missing value of control group

missing.t      the counts of missing value of treatment group

stddiff        the standardized difference between two groups

stddiff.l      the lower limit of the 95 percentage confidence interval of standardized difference

stddiff.u      the upper limit of the 95 percentage confidence interval of standardized difference

for stddiff.category function:

p.c             the proportion of each level in the control group

p.t             the proportion of each level in the treatment group

missing.c      the counts of missing value of control group

missing.t      the counts of missing value of treatment group

stddiff        the standardized difference between two groups

stddiff.l      the lower limit of the 95 percentage confidence interval of standardized difference

stddiff.u      the upper limit of the 95 percentage confidence interval of standardized difference

## Note

Update:

version 2.0: Avoiding the negative number for the 'stddiff' of stddiff.numeric() and stddiff.binary()

version 3.0: Fixing the incorrect format in the results of stddiff.category()

version 3.1: Fixing the incorrect counts of missing values of stddiff.numeric(), stddiff.binary(), stddiff.category()

## Author(s)

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## References

Yang DS, Dalton JE. A Unified Approach to Measuring the Effect Size Between Two Groups Using SAS. SAS Global Forum 2012. paper 335

## See Also

nothing

**Examples**

```
set.seed(2016)
treat<-round(abs(rnorm(100)+1)*10,0)
numeric<-round(abs(rnorm(100)+1)*10,0)
binary<-round(abs(rnorm(100)+1)*10,0)
category<-round(abs(rnorm(100)+1)*10,0)
data<-data.frame(treat,numeric,binary,category)
stddiff.numeric(data=data,gcol=1,vcol=c(2,2))
#stddiff.binary(data=data,gcol=1,vcol=c(3,3))
#stddiff.category(data=data,gcol=1,vcol=c(4,4))
```

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