

# Package: `smd` (via `r-universe`)

September 3, 2024

**Type** Package

**Title** Compute Standardized Mean Differences

**Version** 0.7.0

**Description** Computes standardized mean differences and confidence intervals for multiple data types based on Yang, D., & Dalton, J. E. (2012) <<https://support.sas.com/resources/papers/proceedings12/335-2012.pdf>>.

**Imports** MASS (>= 7.3-50), methods (>= 3.5.1)

**Suggests** testthat, stddiff, tableone, knitr, dplyr, purrr, markdown, rmarkdown

**License** MIT + file LICENSE

**URL** <https://bsaul.github.io/smd/>

**BugReports** <https://github.com/bsaul/smd/issues>

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 7.3.1

**VignetteBuilder** knitr

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

**RemoteUrl** <https://github.com/bsaul/smd>

**RemoteRef** HEAD

**RemoteSha** e7758588096c0fd517724dccb07560bf60213c5

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smd

*Compute Standardized Mean Difference***Description**

Computes the standardized mean difference (SMD) between two groups.

$$d = \sqrt{D'S^{-1}D}$$

where  $D$  is a vector of differences between group 1 and 2 and  $S$  is the covariance matrix of these differences. If  $D$  is length 1, the result is multiplied by  $sign(D)$ .

In the case of a numeric or integer variable, this is equivalent to:

$$d = \frac{\bar{x}_1 - \bar{x}_2}{\sqrt{(s_1^2 + s_2^2)/2}}$$

where  $\bar{x}_g$  is the sample mean for group  $g$  and  $s_g^2$  is the sample variance.

For a logical or factor with only two levels, the equation above is  $\bar{x}_g = \hat{p}_g$ , i.e. the sample proportion and  $s_g^2 = \hat{p}_g(1 - \hat{p}_g)$ .

When using the SMD to evaluate the effectiveness of weighting in achieving covariate balance, it is important to isolate the change in SMD before and after weighting to the change in mean difference, so the denominator (covariance matrix) must be held constant (Stuart 2008, doi:10.1002/sim.3207). By default, the unweighted covariance matrix is used to compute SMD in both the unweighted and weighted case. If the weights are not being used to adjust for covariate imbalance (e.g. case weights), the `unwgt.var` argument can be set to `FALSE` to use the weighted covariance matrix as the denominator.

**Usage**

```
smd(x, g, w, std.error = FALSE, na.rm = FALSE, gref = 1L, unwgt.var = TRUE)

## S4 method for signature 'character,ANY,missing'
smd(x, g, w, std.error = FALSE, na.rm = FALSE, gref = 1L, unwgt.var = TRUE)

## S4 method for signature 'character,ANY,numeric'
smd(x, g, w, std.error = FALSE, na.rm = FALSE, gref = 1L, unwgt.var = TRUE)

## S4 method for signature 'logical,ANY,missing'
smd(x, g, w, std.error = FALSE, na.rm = FALSE, gref = 1L, unwgt.var = TRUE)

## S4 method for signature 'logical,ANY,numeric'
smd(x, g, w, std.error = FALSE, na.rm = FALSE, gref = 1L, unwgt.var = TRUE)

## S4 method for signature 'matrix,ANY,missing'
smd(x, g, w, std.error = FALSE, na.rm = FALSE, gref = 1L, unwgt.var = TRUE)
```

```
## S4 method for signature 'matrix,ANY,numeric'
smd(x, g, w, std.error = FALSE, na.rm = FALSE, gref = 1L, unwtg.var = TRUE)

## S4 method for signature 'list,ANY,missing'
smd(x, g, w, std.error = FALSE, na.rm = FALSE, gref = 1L, unwtg.var = TRUE)

## S4 method for signature 'list,ANY,numeric'
smd(x, g, w, std.error = FALSE, na.rm = FALSE, gref = 1L, unwtg.var = TRUE)

## S4 method for signature 'data.frame,ANY,missing'
smd(x, g, w, std.error = FALSE, na.rm = FALSE, gref = 1L, unwtg.var = TRUE)

## S4 method for signature 'data.frame,ANY,numeric'
smd(x, g, w, std.error = FALSE, na.rm = FALSE, gref = 1L, unwtg.var = TRUE)
```

### Arguments

x	a vector or matrix of values
g	a vector of at least 2 groups to compare. This should be coercible to a factor.
w	a vector of numeric weights (optional)
std.error	Logical indicator for computing standard errors using <code>compute_smd_var</code> . Defaults to FALSE.
na.rm	Remove NA values from x? Defaults to FALSE.
gref	an integer indicating which level of g to use as the reference group. Defaults to 1.
unwtg.var	Use unweighted or weighted covariance matrix. Defaults to TRUE

### Value

a `data.frame` containing standardized mean differences between levels of g for values of x. The `data.frame` contains the columns:

- term: the level being compared to the reference level
- estimate: SMD estimates
- std.error: (if `std.error = TRUE`) SMD standard error estimates

### Examples

```
x <- rnorm(100)
g <- rep(1:2, each = 50)
smd(x, g)
```

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