

# Package ‘tidyrstats’

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**Type** Package

**Depends** R (>= 4.1.0)

**Title** Tidy Common R Statistical Functions

**Version** 0.1.0

**Maintainer** Brendan Ansell <ansell.b@wehi.edu.au>

**Description** Provides functions to scale, log-transform and fit linear models within a 'tidyverse'-style R code framework.

Intended to smooth over inconsistencies in output of base R statistical functions, allowing ease of teaching, learning and daily use. Inspired by the tidy principles used in 'broom' Robinson (2017) <[doi:10.21105/joss.00341](https://doi.org/10.21105/joss.00341)>.

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**Encoding** UTF-8

**RoxygenNote** 7.3.2

**Imports** broom, glue, purrr, dplyr, rlang, stringr, stats

**Suggests** ggplot2, knitr, magrittr, rmarkdown

**NeedsCompilation** no

**Author** Brendan Ansell [aut, cre] (ORCID:  
<<https://orcid.org/0000-0003-0297-897X>>)

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`lm_test`*Linear Model Testing for Grouped, Nested, or Ungrouped Data*

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### Description

Applies a linear model to a data frame and returns tidy model summaries. Supports ungrouped, grouped (`dplyr::group_by()`), and nested (`tidyr::nest_by()`) input data.

### Usage

```
lm_test(input_data, formula)
```

### Arguments

`input_data` A data frame or tibble. Can be ungrouped, grouped, or nested.  
`formula` A model formula, either quoted or unquoted (e.g., `y ~ x * z`, or `"y ~ x * z"`).

### Details

Designed to allow seamless 'in-line' chaining to fit linear models to columns of a tibble. Compatible with ungrouped, grouped or nested input. Compatible with native and magrittr pipe. Uses `broom::tidy()` to extract model summaries.

### Value

A tibble with tidy model output sorted by p value, including:

**term** Model term (e.g., intercept, predictors, interactions)

**estimate** Estimated coefficient / beta

**std.error** Standard error of the estimate

**statistic** t-statistic

**p.value** p-value for the hypothesis test

If the input is grouped or nested, group identifiers are retained in the output. In the nested case, nested terms are relocated to the left-most column of the tibble.

### Examples

```
library(ggplot2)
library(dplyr)

# Ungrouped
mpg |> lm_test(cty ~ hwy * cyl)

# Grouped
mpg |> group_by(class) |> lm_test(cty ~ hwy * cyl)

# Nested
```

```
mpg |> nest_by(class) |> lm_test(cty ~ hwy * cyl)
```

---

**neg\_log***Negative Logarithm (Base 10 by Default)*

---

**Description**

Computes the negative logarithm of a numeric input using base 10 by default.

**Usage**

```
neg_log(x, base = 10)
```

```
neglog(x, base = 10)
```

**Arguments**

**x** A numeric vector. Values must be positive.

**base** A numeric value specifying the base of the logarithm. Default is 10.

**Details**

This function returns the negative logarithm of 'x'. By default, it uses base 10, but you can specify a different base using the 'base' argument. Designed for quickly transforming p values for statistical analysis.

**Value**

A numeric vector of negative logarithmic values.

**Examples**

```
pvals <- 10^runif(10, -15, -1)
neg_log(pvals)
```

---

`scale_this`*Scale a numeric vector without converting to a matrix*

---

**Description**

This function scales and centres a numeric vector by subtracting the mean and dividing by the standard deviation. Unlike `scale()`, it returns a numeric vector, not a matrix. Note this function does not allow control over centering or scaling.

**Usage**

```
scale_this(x)
```

**Arguments**

`x` A numeric vector.

**Value**

A numeric vector of scaled values.

**Examples**

```
iris_dat <- head(iris$Sepal.Length)
scale_this(iris_dat)
scale_this(c(iris_dat, NA))
```

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