

Package ‘ParamANOVA’

May 7, 2026

Title Fully Parameterizable ANOVA Tests

Version 0.1.4

Description Allows the user to perform ANOVA tests (in a strict sense: continuous and normally-distributed Y variable and 1 or more factorial/categorical X variable(s)), with the possibility to specify the type of sum of squares (1, 2 or 3), the types of variables (Fixed or Random) and their relationships (crossed or nested) with the sole function of the package (FullyParamANOVA()). The resulting outputs are the same as in 'SAS' software. A dataset (Butterfly) to test the function is also joined.

License MIT + file LICENSE

Encoding UTF-8

RoxygenNote 7.2.3

Imports dplyr (>= 1.1.2), magrittr (>= 2.0.3), mlr3misc (>= 0.16.0),
rlang (>= 1.1.1), rstatix (>= 0.7.2), stats (>= 4.2.1), tibble
(>= 3.2.1)

Depends R (>= 3.5)

LazyData true

NeedsCompilation no

Author Victor Brans [aut, cre] (ORCID:
<<https://orcid.org/0000-0002-8960-100X>>)

Maintainer Victor Brans <vicbrans003@gmail.com>

Repository CRAN

Date/Publication 2026-04-23 11:20:08 UTC

Contents

Butterfly	2
FullyParamANOVA	2

Index	5
--------------	----------

Butterfly

Wingspans of butterflies in different sites

Description

Imaginary data set of wingspans (in cm) of butterflies in different patches of different sites

Usage

```
data("Butterfly")
```

Format

A data frame with 300 observations of the following 4 variables.

Sex a character vector: M=males; F=Females

Site a numeric vector: 3 different sites

Patch a numeric vector: 5 patches per site

Wingspan a numeric vector

Source

Data created by Estelle Laurent

References

Data created by Estelle Laurent

FullyParamANOVA

Fully Parameterizable ANOVA test

Description

Function allowing the user to perform ANOVA tests (in a strict sense: one continuous and normally-distributed response variable and one or more factorial/categorical explaining variable(s)), with the possibility to specify the type of sum of squares (1, 2 or 3), the types of factors (Fixed or Random) and their relationships (crossed or nested). This function is built on the `'?rstatix::anova_test()'`. The resulting outputs are the same as in 'SAS' software.

Usage

```
FullyParamANOVA(
  InputData,
  form,
  variables_type = NULL,
  variables_relationships = c(NA),
  sig_threshold = 0.05,
  sum_of_squares = 3,
  print_ges = FALSE,
  output_table_only = FALSE
)
```

Arguments

InputData	The name of the input dataset.
form	To be written as <i>Response variable ~ Explaining variable(s)</i> . If there are multiple <i>Explaining variables</i> , they should be separated by a <i>*</i> .
variables_type	Type of factors in a vector. One argument per <i>Explaining variable</i> , in the same order as in the formula: 'Fxd' for Fixed factors, or 'Rndm' for Random factors
variables_relationships	Relationships between factors in a vector. One argument per <i>Explaining variable</i> , in the same order as in the formula: <i>NA</i> if the factor is not nested in any other factor, or the name of the factor it is nested into.
sig_threshold	The threshold below which a p-value is considered significant. By default, it is set to the usual value of 0.05.
sum_of_squares	Type of sum of squares used in the test: 1, 2 or 3. These numbers correspond to those in 'SAS' software. By default, the type 3 is used as it is the most often recommended one.
print_ges	If <i>TRUE</i> , prints effect sizes (ges = generalized eta squared) in the result table. See '?rstatix::anova_test()' for more details. It is <i>FALSE</i> by default.
output_table_only	If <i>TRUE</i> , reports only the result table and not the text below specifying the parameters used for the test, allowing for the output table to be directly usable or stored in an object for example. It is <i>FALSE</i> by default.

Value

ANOVA result table.

There is one row per effect tested and a last row for the residuals.

The first column ("Effect") specifies the effects tested.

The second column ("SS") gives the Sum of Squares associated with the effects tested.

The third column ("DF") gives the Degrees of Freedom associated with the effects tested.

The fourth column ("MSS") gives the Mean Sum of Squares (=SS/DF) associated with the effects tested.

The fifth column ("Fval") gives the F statistic value associated with the effects tested.

The sixth column ("pval") gives the p-value associated with the effects tested.

The seventh column ("Sig") shows a star if the associated p-value is below the statistical threshold of 0.05 by default or the one indicated during the call of the function for each effects tested.

The optional eighth column ("ges") gives the effect sizes (ges = generalized eta squared) associated with the effects tested.

If the argument *output_table_only* is *FALSE* (which is the case by default), the function also specifies below this result table which types of Sum of Squares were used, what is the response variable, which factors were fixed or random, and if there are at least two factors, it specifies the relationships between these factors.

You will find warnings below if the setup is not appropriate.

Examples

```
data("Butterfly")
# Example of ANOVA I
FullyParamANOVA(InputData=Butterfly, form=Wingspan~Site, variables_type = c('Fxd'))

# Example of ANOVA II
FullyParamANOVA(InputData=Butterfly, form=Wingspan~Patch*Site,
variables_type = c('Rndm','Fxd'), variables_relationships = c("Site",NA))

# Example of ANOVA III
FullyParamANOVA(InputData=Butterfly, form=Wingspan~Sex*Patch*Site,
variables_type = c('Fxd','Rndm','Fxd'), variables_relationships = c(NA,"Site",NA))

# Example of ANOVA III, using arguments different than those by default
# (changing the threshold below which a p-value is considered significant to 0.1,
# the types of sum of squares to 2, adding the effect sizes (ges = generalized eta squared)
# in the output table, and outputting only the table to be able to use the information within
# or store it in an object)
FullyParamANOVA(InputData=Butterfly, form=Wingspan~Sex*Patch*Site, sig_threshold=0.1,
sum_of_squares = 2, variables_type = c('Fxd','Rndm','Fxd'),
variables_relationships = c(NA,"Site",NA), print_ges = TRUE, output_table_only = TRUE)
```

Index

* **datasets**

Butterfly, [2](#)

Butterfly, [2](#)

FullyParamANOVA, [2](#)