

statsExpressions: R Package for Tidy Dataframes and Expressions with Statistical Details

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Summary

The statsExpressions package has two key aims: to provide a consistent syntax to do statistical analysis with tidy data, and to provide statistical expressions (i.e., a preformatted in-text statistical result) for plotting functions. Currently, it supports common types of statistical approaches and tests: parametric, nonparametric, robust, and Bayesian t-test, one-way ANOVA, correlation analyses, contingency table analyses, and meta-analyses. The functions are pipe-friendly and compatible with tidy data.

Statement of need

Statistical packages exhibit substantial diversity in terms of their syntax and expected input type. This can make it difficult to switch from one statistical approach to another. For example, some functions expect vectors as inputs, while others expect dataframes. Depending on whether it is a repeated measures design or not, different functions might expect data to be in wide or long format. Some functions can internally omit missing values, while other functions error in their presence. Furthermore, if someone wishes to utilize the objects returned by these packages downstream in their workflow, this is not straightforward either because even functions from the same package can return a list, a matrix, an array, a dataframe, etc., depending on the function.

This is where statsExpressions comes in: It can be thought of as a unified portal through which most of the functionality in these underlying packages can be accessed, with a simpler interface and no requirement to change data format.

Comparison to Other Packages

Unlike broom (Robinson, Hayes, & Couch, 2021) or parameters (Lüdecke, Ben-Shachar, Patil, & Makowski, 2020), the goal of statsExpressions is not to convert model objects into tidy dataframes, but to provide a consistent and easy syntax to carry out statistical tests. Additionally, none of these packages return statistical expressions.

Tidy Dataframes from Statistical Analysis

The package offers six primary functions that let users choose a statistical approach without changing the syntax (i.e., by only specifying a single argument). The users are always expected to provide a dataframe in tidy format (Wickham et al., 2019), and all functions work with missing data. Moreover, they always return a dataframe that can be further utilized downstream in the workflow (for a visualization, e.g.).

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Table 1: A summary table listing the primary functions in the package and the statistical approaches they support. For a more detailed description of the tests and outputs from these functions, the readers are encouraged to read vignettes on the package website: https://indrajeetpatil.github.io/statsExpressions/articles/.

Function	Parametric	Non-parametric	Robust	Bayesian
one_sample_test	√	√	√	√
two_sample_test	\checkmark	\checkmark	\checkmark	\checkmark
oneway_anova	\checkmark	\checkmark	\checkmark	\checkmark
corr_test	\checkmark	\checkmark	\checkmark	\checkmark
contingency_table	\checkmark	\checkmark	-	\checkmark
meta_analysis	\checkmark	-	\checkmark	\checkmark

statsExpressions internally relies on stats package for parametric and non-parametric (R Core Team, 2021), WRS2 package for robust (Mair & Wilcox, 2020), and BayesFactor package for Bayesian statistics (Morey & Rouder, 2020). The random-effects meta-analysis is carried out using metafor (parametric) (Viechtbauer, 2010), metaplus (robust) (Beath, 2016), and metaBMA (Bayesian) (Heck et al., 2019) packages. Additionally, it relies on easystats packages (Ben-Shachar, Lüdecke, & Makowski, 2020; Lüdecke, Ben-Shachar, Patil, & Makowski, 2020; Lüdecke, Ben-Shachar, Patil, Waggoner, & Makowski, 2021; Lüdecke, Waggoner, & Makowski, 2019; Makowski, Ben-Shachar, & Lüdecke, 2019; Makowski, Ben-Shachar, Patil, & Lüdecke, 2020) to compute appropriate effect size/posterior estimates and their confidence/credible intervals.

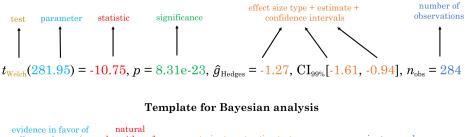
To illustrate the simplicity of this syntax, let's say we want to run a two-sample *t*-test. If we first run a parametric *t*-test and then decide to run a robust *t*-test instead, the syntax remains the same and the statistical approach can be modified by changing a single argument:

```
mtcars %>% two_sample_test(am, wt, type = "parametric") # Welch's t-test
#> # A tibble: 1 x 14
#>
     term group mean.group1 mean.group2 statistic df.error
                                                                  p.value
#>
     <chr> <chr>
                        <\!db\,l>
                                    <db1>
                                               <dbl>
                                                        <db1>
                                                                    <db1>
                         3.77
#> 1 wt
           a.m
                                     2.41
                                                5.49
                                                         29.2 0.00000627
#>
                              estimate conf.level conf.low conf.high effectsize
     met.h.od.
#>
     <chr>
                                 <dbl>
                                             <dbl>
                                                      <dbl>
                                                                 <dbl> <chr>
#> 1 Welch Two Sample t-test
                                  1.84
                                              0.95
                                                       1.00
                                                                  2.66 Hedges' q
     expression
#>
     t>
#> 1 < language>
mtcars %>% two_sample_test(am, wt, type = "robust") # Yuen's t-test
#> # A tibble: 1 x 10
#>
     statistic df.error
                           p.value
#>
         <dbl>
                  <dbl>
                             <d.b1.>
#> 1
          5.84
                   13.6 0.0000485
#>
     method.
                                                            estimate conf.low
#>
                                                                <db1>
                                                                         <db1>
#> 1 Yuen's test on trimmed means for independent samples
                                                                0.915
                                                                         0.754
     conf.high conf.level effectsize
#>
                                                                expression
#>
         <db1>
                     <dbl> <chr>
                                                                t>
                     0.95 Explanatory measure of effect size <language>
```

These functions are also compatible with other popular data manipulation packages (see Appendix for an example).



Template for Frequentist analysis



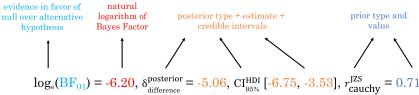


Figure 1: The templates used in 'statsExpressions' to display statistical details in a plot.

Expressions for Plots

In addition to other details contained in the dataframe, there is also a column titled expression, which contains a pre-formatted text with statistical details. These expressions (Figure 1) attempt to follow the gold standard in statistical reporting for both Bayesian (Doorn et al., 2020) and Frequentist (Association & others, 2019) frameworks.

This expression be easily displayed in a plot (Figure 2). Displaying statistical results in the context of a visualization is indeed a philosophy adopted by the ggstatsplot package (Patil, 2021), and statsExpressions functions as its statistical processing backend.

Licensing and Availability

statsExpressions is licensed under the GNU General Public License (v3.0), with all source code stored at GitHub. In the spirit of honest and open science, requests and suggestions for fixes, feature updates, as well as general questions and concerns are encouraged via direct interaction with contributors and developers by filing an issue while respecting *Contribution Guidelines*.

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Kruskal-Wallis Rank Sum Test

 $\chi^2_{\text{Kruskal-Wallis}}(2) = 217.60, p = 5.61e-48, \hat{\epsilon}^2_{\text{ordinal}} = 0.64, \text{Cl}_{95\%} [0.61, 0.66], n_{\text{obs}} = 342$

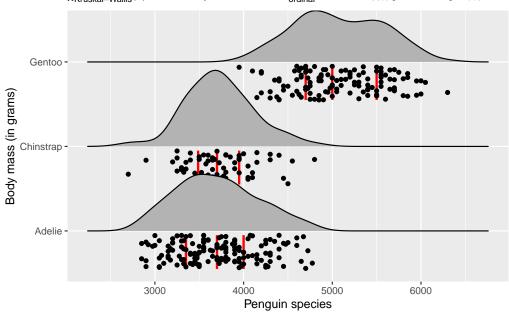


Figure 2: Example illustrating how 'statsExpressions' functions can be used to display results from a statistical test in a plot. Code to create this figure is reported in Appendix.

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Appendix

Example with dplyr

We can use combination of dplyr and statsExpressions to repeat the same statistical analysis across grouping variables.

```
# running one-sample proportion test for all levels of `cyl`
mtcars %>%
 group_by(cyl) %>%
 group_modify(~ contingency_table(.x, am), .keep = TRUE) %>%
 ungroup()
#> # A tibble: 3 x 11
      cyl statistic
                       df p.value method
#>
     <dbl>
              <dbl> <dbl> <dbl> <chr>
              2.27
#> 1
                      1 0.132 Chi-squared test for given probabilities
        4
                        1 0.705 Chi-squared test for given probabilities
#> 2
         6
              0.143
                        1 0.00753 Chi-squared test for given probabilities
#> 3
        8
              7.14
#>
    estimate conf.level conf.low conf.high effectsize
                                                             expression
#>
        <dbl>
                           <dbl>
                                     <dbl> <chr>
                                                             t>
                  <db1>
#> 1
        0.344
                   0.95
                           0
                                     0.917 Cramer's V (adj.) <language>
                   0.95
                                           Cramer's V (adj.) <language>
#> 2
        0
#> 3
        0.685
                   0.95
                           0.127
                                     1.18 Cramer's V (adj.) <language>
```

Code to reproduce for Figure 2

```
# needed libraries
library(statsExpressions)
library(ggplot2)
library(ggridges)
library(palmerpenguins) # `penguins` dataset is from this package
```



```
# creating a dataframe
res <- oneway_anova(penguins, species, body_mass_g, type = "nonparametric")
# create a ridgeplot using `ggridges` package
ggplot(penguins, aes(x = body_mass_g, y = species)) +
  geom_density_ridges(
   jittered_points = TRUE,
    quantile_lines = TRUE,
   scale = 0.9,
   vline_size = 1,
   vline_color = "red",
   position = position_raincloud(adjust_vlines = TRUE)
  ) + # use 'expression' column to display results in the subtitle
  labs(
   x = "Penguin species",
   y = "Body mass (in grams)",
   title = "Kruskal-Wallis Rank Sum Test",
    subtitle = res$expression[[1]]
```