

Package ‘tidystats’

June 15, 2020

Type Package

Title Save Output of Statistical Tests

Version 0.4.1

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Description Produce a file containing the output of statistical tests
that can be shared or used to report statistics in scientific papers.

URL <https://willemsleegers.github.io/tidystats/>

BugReports <https://github.com/WillemSleegers/tidystats/issues>

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Encoding UTF-8

RoxygenNote 7.1.0

LazyData true

Depends R (>= 2.10)

Imports dplyr, tidyr, purrr, stringr, readr, jsonlite, tibble

Suggests knitr, rmarkdown, testthat, BayesFactor

VignetteBuilder knitr

NeedsCompilation no

Repository CRAN

Date/Publication 2020-06-15 06:00:08 UTC

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Index**13****add_stats***Add statistical output to a tidystats list***Description**

`add_stats` is used to add the output of a statistical test to a tidystats list. While adding the output, additional information about the test can be added, including the type of test (primary, secondary, or exploratory), whether the test was preregistered, and additional notes. Please note that not all statistical tests are supported. See 'Details' below for a list of supported statistical tests.

Usage

```
add_stats(
  results,
  output,
  identifier = NULL,
  type = NULL,
  preregistered = NULL,
  notes = NULL
)
```

Arguments

<code>results</code>	A tidystats list.
<code>output</code>	Output of a statistical test.
<code>identifier</code>	A character string identifying the model. Automatically created if not provided.
<code>type</code>	A character string specifying the type of analysis: primary, secondary, or exploratory.
<code>preregistered</code>	A boolean specifying whether the analysis was preregistered or not.
<code>notes</code>	A character string specifying additional information.

Details

Currently supported functions:

- `t.test()`
- `cor.test()`
- `chisq.test()`
- `wilcox.test()`
- `fisher.test()`

- oneway.test()
- aov()
- lm()

Examples

```
# Load dplyr for access to the piping operator
library(dplyr)

# Conduct statistical tests
# t-test:
sleep_test <- t.test(extra ~ group, data = sleep, paired = TRUE)

# lm:
ctl <- c(4.17,5.58,5.18,6.11,4.50,4.61,5.17,4.53,5.33,5.14)
trt <- c(4.81,4.17,4.41,3.59,5.87,3.83,6.03,4.89,4.32,4.69)
group <- gl(2, 10, 20, labels = c("Ctl","Trt"))
weight <- c(ctl, trt)
lm_D9 <- lm(weight ~ group)

# ANOVA:
npk_aov <- aov(yield ~ block + N*P*K, npk)

#' # Create an empty list
results <- list()

# Add output to the results list
results <- results %>%
  add_stats(sleep_test) %>%
  add_stats(lm_D9, type = "primary", preregistered = TRUE) %>%
  add_stats(npk_aov, notes = "An ANOVA example")
```

count_data

Count the number of observations

Description

count_data returns the number of observations for categorical variables.

Usage

```
count_data(data, ..., na.rm = FALSE)
```

Arguments

- | | |
|-------|---|
| data | A data frame. |
| ... | One or more unquoted (categorical) column names from the data frame, separated by commas. |
| na.rm | Logical. Should missing values (including NaN) be removed? |

Details

The data frame can be grouped using **dplyr**'s `group_by` so that the number of observations will be calculated for each group level.

Examples

```
# Load dplyr for access to the %>% operator and group_by()
library(dplyr)

# 1 variable
count_data(quote_source, source)

# 2 variables
count_data(quote_source, source, sex)

# Ignore missing values
count_data(quote_source, source, sex, na.rm = TRUE)

# Use group_by() to get percentages within each group
quote_source %>%
  group_by(source) %>%
  count_data(sex)
```

cox

Data of a replication study of C.R. Cox, J. Arndt, T. Pyszczynski, J. Greenberg, A. Abdollahi, S. Solomon (2008, JPSP, 94(4), Exp. 6)

Description

This is the data of a replication study performed by J. Wissink, G. Hoogendoorn, H. Brohmer, M. Verschoor, J. Krijnen, and M. Zeelenberg as part of the Reproducibility Project: Psychology. The target result of this replication was the finding in Experiment 6 of Cox et al. (2008) that participants who scored low on avoidance but high on anxiety demonstrated an increased relative preference for a parent after mortality salience as opposed to dental pain, $b = -32.04$, $SE = 14.47$, $t = -2.22$, $p = .03$.

Usage

cox

Format

A data frame with 200 rows and 67 variables.

ID Participant identifier

sex The participant's sex

age The participant's age

condition The experimental condition: mortality salience or dental pain
avoidance Attachment avoidance score as assessed with the Relationship Scales Questionnaire
anxiety Attachment anxiety score as assessed with the Relationship Scales Questionnaire
affect_positive Sum of positive PANAS items
affect_negative Sum of negative PANAS items
call_parent Minutes allocated (out of 100) to call a parent
call_siblings Minutes allocated (out of 100) to call a sibling
call_partner Minutes allocated (out of 100) to call a romantic partner
call_friend Minutes allocated (out of 100) to call a close friend

Details

For more information on the Reproducibility Project: Psychology, see <https://osf.io/ezcuj/>. The individual scores on the PANAS and RSQ items are not included in this dataset.

Source

<https://osf.io/5tbxf/>

References

Cox, C.R., Arndt, J., Pyszczynski, T., Greenberg, J., Abdollahi, A., & Solomon, S. (2008) Terror management and adults' attachment to their parents: The safe haven remains. *Journal of Personality and Social Psychology*, 94(4), 696-717, <https://dx.doi.org/10.1037/0022-3514.94.4.696>

describe_data	<i>Calculate common descriptive statistics</i>
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Description

describe_data returns a set of common descriptive statistics (e.g., n, mean, sd) for numeric variables.

Usage

```
describe_data(data, column, na.rm = TRUE, short = FALSE)
```

Arguments

data	A data frame.
column	An unquoted (numerical) column name from the data frame.
na.rm	Logical. Should missing values (including NaN) be excluded in calculating the descriptives? The default is TRUE.
short	Logical. Should only a subset of descriptives be reported? If set to TRUE, only the N, M, and SD will be returned. The default is FALSE.

Details

The data can be grouped using `dplyr::group_by` so that descriptives will be calculated for each group level.

When `na.rm` is set to FALSE, a percentage column will be added to the output that contains the percentage of non-missing data.

Skew and kurtosis are based on the `skewness` and `kurtosis` functions of the `moments` package (Komsta & Novomestky, 2015).

Percentages are calculated based on the total of non-missing observations. When `na.rm` is set to FALSE, percentages are based on the total of missing and non-missing observations.

Examples

```
# Load the dplyr package for access to the %>% operator and group_by()
library(dplyr)

# Inspect descriptives of the response column from the 'quote_source' data
# frame included in tidystats
describe_data(quote_source, response)

# Repeat the former, now for each level of the source column
quote_source %>%
  group_by(source) %>%
  describe_data(response)

# Only inspect the total N, mean, and standard deviation
quote_source %>%
  group_by(source) %>%
  describe_data(response, short = TRUE)
```

quote_source

A many labs replication of Lorge & Curtiss (1936)

Description

Data of multiple replication studies of Lorge & Curtiss (1936) from the Many Labs project (Klein et al., 2014).

Usage

`quote_source`

Format

A data frame with 6343 rows and 15 columns:

ID participant number

source attributed source of the quote: Washington or Bin Laden
response evaluation of the quote on a 9-point Likert scale, with 1 indicating disagreement and 9 indicating agreement
age participant's age
sex participant's sex
citizenship participant's citizenship
race participant's race
major participant's major
native_language participant's native language
referrer location of where the study was conducted
compensation how the participant was compensated for their participation
recruitment how the participant was recruited
separated_or_not description of how the study was administered in terms of participant isolation
us_or_international whether the study was conducted in the US or outside of the US (international)
lab_or_online whether the study was conducted in the lab or online

Details

Lorge and Curtiss (1936) examined how a quotation is perceived when it is attributed to a liked or disliked individual. The quotation of interest was, "I hold it that a little rebellion, now and then, is a good thing, and as necessary in the political world as storms are in the physical world." In one condition the quotation was attributed to Thomas Jefferson, a liked individual, and in the other it was attributed to Vladimir Lenin, a disliked individual. More agreement was observed when the quotation was attributed to Jefferson than Lenin. In the replication studies, the quotation was attributed to either George Washington, the liked individual, or Osama Bin Laden, the disliked individual.

References

- Lorge, I., & Curtiss, C. C. (1936). Prestige, suggestion, and attitudes. *The Journal of Social Psychology*, 7, 386-402. <https://doi.org/10.1080/00224545.1936.9919891>
- Klein, R.A. et al. (2014) Investigating Variation in Replicability: A "Many Labs" Replication Project. *Social Psychology*, 45(3), 142-152. <https://dx.doi.org/10.1027/1864-9335/a000178>

read_stats

Read a .json file that was produced with write_stats

Description

read_stats can read in a .json file containing the statistical output that was produced with write_stats. It returns a list containing the results, with the identifier as the name for each list element.

Usage

```
read_stats(file)
```

Arguments

file	Path to the tidy stats data file
------	----------------------------------

Examples

```
results <- read_stats(system.file("results.json", package = "tidystats"))
```

tidy_stats

Tidy the output of a statistics object

Description

`tidy_stats` is used to convert the output of a statistical object to a list of organized statistics. This output can then be added to a list using the `add_stats` function of this package. The `tidy_stats` function is automatically run when `add_stats` is used, so there is generally no need to use this function explicitly. It can be used, however, to quickly peek at how the output of a specific analysis will be organized. Please note that not all statistical tests are supported. See 'Details' below for a list of supported statistical tests.

Usage

```
tidy_stats(x)

## S3 method for class 'htest'
tidy_stats(x)

## S3 method for class 'lm'
tidy_stats(x)

## S3 method for class 'lmerMod'
tidy_stats(x)

## S3 method for class 'lmerModLmerTest'
tidy_stats(x)

## S3 method for class 'aov'
tidy_stats(x)

## S3 method for class 'aoeplist'
tidy_stats(x)

## S3 method for class 'tidystats_descriptives'
```

```
tidy_stats(x)

## S3 method for class 'tidystats_counts'
tidy_stats(x)

## S3 method for class 'anova'
tidy_stats(x)

## S3 method for class 'BFBayesFactor'
tidy_stats(x)

## S3 method for class 'afex_aov'
tidy_stats(x)
```

Arguments

- x The output of a statistical test.

Details

Currently supported functions:

- t.test()
- cor.test()
- chisq.test()
- wilcox.test()
- fisher.test()
- oneway.test()
- aov()
- lm()

Methods (by class)

- htest: tidy_stats method for class 'htest'
- lm: tidy_stats method for class 'lm'
- lmerMod: tidy_stats method for class 'lmerMod'
- lmerModLmerTest: tidy_stats method for class 'lmerModLmerTest'
- aov: tidy_stats method for class 'aov'
- aovlist: tidy_stats method for class 'aovlist'
- tidystats_descriptives: tidy_stats method for class 'tidystats_descriptives'
- tidystats_counts: tidy_stats method for class 'tidystats_counts'
- anova: tidy_stats method for class 'anova'
- BFBayesFactor: tidy_stats method for class 'BayesFactor'
- afex_aov: tidy_stats method for class 'afex_aov'

Examples

```
# Conduct statistical tests
# t-test:
sleep_test <- t.test(extra ~ group, data = sleep, paired = TRUE)

# lm:
ctl <- c(4.17,5.58,5.18,6.11,4.50,4.61,5.17,4.53,5.33,5.14)
trt <- c(4.81,4.17,4.41,3.59,5.87,3.83,6.03,4.89,4.32,4.69)
group <- gl(2, 10, 20, labels = c("Ctl","Trt"))
weight <- c(ctl, trt)
lm_D9 <- lm(weight ~ group)

# ANOVA:
npk_aov <- aov(yield ~ block + N*P*K, npk)

# Tidy the statistics and store each analysis in a separate variable
list_sleep_test <- tidy_stats(sleep_test)
list_lm_D9 <- tidy_stats(lm_D9)
list_npkaov <- tidy_stats(npk_aov)

# Now you can inspect each of these variables, e.g.:
names(list_sleep_test)
str(list_sleep_test)
```

tidy_stats_to_data_frame

Convert a tidystats list to a data frame

Description

`tidy_stats_to_data_frame` converts a tidystats list to a data frame, which can then be used to easily extract specific statistics using standard subsetting functions (e.g., `dplyr::filter`).

Usage

```
tidy_stats_to_data_frame(x)
```

Arguments

x	A tidystats list.
---	-------------------

Examples

```
# Load dplyr for access to the piping operator
library(dplyr)

# Conduct statistical tests
t_test_1 <- t.test(1:10, y = c(7:20))
```

```
t_test_2 <- t.test(1:10, y = c(7:20, 200))
t_test_3 <- t.test(extra ~ group, data = sleep)

#' # Create an empty list
results <- list()

# Add tests to the empty list
results <- results %>%
  add_stats(t_test_1) %>%
  add_stats(t_test_2) %>%
  add_stats(t_test_3)

# Convert the list to a data frame
results_df <- tidy_stats_to_data_frame(results)

# Select all the p-values
results_df %>%
  filter(statistic == "p") %>%
  pull(value)
```

write_stats	<i>Write a tidystats list to a file</i>
-------------	---

Description

write_stats writes a tidystats list to a .json file.

Usage

```
write_stats(x, path, digits = 6)
```

Arguments

x	A tidystats list.
path	Path or connection to write to.
digits	The number of decimal places to use.

Examples

```
# Load dplyr for access to the piping operator
library(dplyr)

# Conduct statistical tests
# t-test:
sleep_test <- t.test(extra ~ group, data = sleep, paired = TRUE)

# lm:
ctl <- c(4.17,5.58,5.18,6.11,4.50,4.61,5.17,4.53,5.33,5.14)
```

```
trt <- c(4.81,4.17,4.41,3.59,5.87,3.83,6.03,4.89,4.32,4.69)
group <- gl(2, 10, 20, labels = c("Ctl","Trt"))
weight <- c(ctl, trt)
lm_D9 <- lm(weight ~ group)

# ANOVA:
npk_aov <- aov(yield ~ block + N*P*K, npk)

#' # Create an empty list
results <- list()

# Add output to the results list
results <- results %>%
  add_stats(sleep_test) %>%
  add_stats(lm_D9, type = "primary", preregistered = TRUE) %>%
  add_stats(npk_aov, notes = "An ANOVA example")

# Save the results
dir <- tempdir()
write_stats(results, file.path(dir, "results.json"))
```

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