

# Package ‘cardx’

March 18, 2024

**Title** Extra Analysis Results Data Utilities

**Version** 0.1.0

**Description** Create extra Analysis Results Data (ARD) summary objects. The package supplements the simple ARD functions from the 'cards' package, exporting functions to put statistical results in the ARD format. These objects are used and re-used to construct summary tables, visualizations, and written reports.

**License** Apache License 2.0

**URL** <https://github.com/insightsengineering/cardx>

**BugReports** <https://github.com/insightsengineering/cardx/issues>

**Depends** R (>= 4.1)

**Imports** cards (>= 0.1.0), cli (>= 3.6.1), dplyr (>= 1.1.2), glue (>= 1.6.2), rlang (>= 1.1.1), tidyr (>= 1.3.0)

**Suggests** broom (>= 1.0.5), broom.helpers (>= 1.13.0), spelling, testthat (>= 3.2.0), withr

**Config/Needs/website** insightsengineering/nesttemplate

**Config/testthat/edition** 3

**Config/testthat/parallel** true

**Encoding** UTF-8

**Language** en-US

**RoxygenNote** 7.3.1

**NeedsCompilation** no

**Author** Daniel Sjoberg [aut, cre],  
F. Hoffmann-La Roche AG [cph, fnd]

**Maintainer** Daniel Sjoberg <daniel.d.sjoberg@gmail.com>

**Repository** CRAN

**Date/Publication** 2024-03-18 14:30:02 UTC

## R topics documented:

ard_chisqtest . . . . .	2
ard_fishtest . . . . .	3
ard_kruskaltest . . . . .	3
ard_mcnemartest . . . . .	4
ard_moodtest . . . . .	5
ard_proportion_ci . . . . .	6
ard_regression . . . . .	7
ard_ttest . . . . .	7
ard_wilcoxtest . . . . .	8
proportion_ci . . . . .	10
<b>Index</b>	<b>13</b>

---

ard_chisqtest	<i>ARD Chi-squared Test</i>
---------------	-----------------------------

---

### Description

Analysis results data for Pearson's Chi-squared Test. Calculated with `chisq.test(x = data[[variable]], y = data[[by]], ...)`

### Usage

```
ard_chisqtest(data, by, variable, ...)
```

### Arguments

data	(data.frame) a data frame.
by, variable	(tidy-select) column names to compare
...	additional arguments passed to <code>fisher.test(...)</code>

### Value

ARD data frame

### Examples

```
cards::ADSL |>
  ard_chisqtest(by = "ARM", variable = "AGEGR1")
```

---

ard_fishertest	<i>ARD Fisher's Exact Test</i>
----------------	--------------------------------

---

**Description**

Analysis results data for Fisher's Exact Test. Calculated with `fisher.test(x = data[[variable]], y = data[[by]], ...)`

**Usage**

```
ard_fishertest(data, by, variable, ...)
```

**Arguments**

<code>data</code>	( <code>data.frame</code> ) a data frame.
<code>by, variable</code>	( <code>tidy-select</code> ) column names to compare
<code>...</code>	additional arguments passed to <code>fisher.test(...)</code>

**Value**

ARD data frame

**Examples**

```
cards::ADSL[1:30, ] |>
  ard_fishertest(by = "ARM", variable = "AGEGR1")
```

---

ard_kruskaltest	<i>ARD Kruskal-Wallis Test</i>
-----------------	--------------------------------

---

**Description**

Analysis results data for Kruskal-Wallis Rank Sum Test.  
Calculated with `kruskal.test(data[[variable]], data[[by]], ...)`

**Usage**

```
ard_kruskaltest(data, by, variable)
```

**Arguments**

data	(data.frame) a data frame.
by	( <a href="#">tidy-select</a> ) column name to compare by
variable	( <a href="#">tidy-select</a> ) column name to be compared

**Value**

ARD data frame

**Examples**

```
cards::ADSL |>
  ard_kruskaltest(by = "ARM", variable = "AGE")
```

---

ard_mcnemartest	<i>ARD McNemar's Test</i>
-----------------	---------------------------

---

**Description**

Analysis results data for McNemar's statistical test.

**Usage**

```
ard_mcnemartest(data, by, variable, ...)
```

**Arguments**

data	(data.frame) a data frame. See below for details.
by	( <a href="#">tidy-select</a> ) column name to compare by.
variable	( <a href="#">tidy-select</a> ) column name to be compared.
...	arguments passed to <code>stats::mcnemar.test(...)</code>

**Details**

For the `ard_mcnemartest()` function, the data is expected to be one row per subject. The data is passed as `stats::mcnemar.test(x = data[[variable]], y = data[[by]], ...)`. Please use `table(x = data[[variable]], y = data[[by]])` to check the contingency table.

**Value**

ARD data frame

**Examples**

```
cards::ADSL |>
  ard_mcnemartest(by = "SEX", variable = "EFFFL")
```

---

ard_moodtest	<i>ARD Mood Test</i>
--------------	----------------------

---

**Description**

Analysis results data for Mood two sample test of scale. Note this not to be confused with the Brown-Mood test of medians.

**Usage**

```
ard_moodtest(data, by, variable, ...)
```

**Arguments**

data	(data.frame) a data frame. See below for details.
by	(tidy-select) column name to compare by.
variable	(tidy-select) column name to be compared.
...	arguments passed to mood.test(...)

**Details**

For the ard\_moodtest() function, the data is expected to be one row per subject. The data is passed as mood.test(data[[variable]] ~ data[[by]], ...).

**Value**

ARD data frame

**Examples**

```
cards::ADSL |>
  ard_moodtest(by = "SEX", variable = "AGE")
```

---

ard\_proportion\_ci      *ARD Proportion Confidence Intervals*

---

## Description

### [Experimental]

Calculate confidence intervals for proportions.

## Usage

```
ard_proportion_ci(
  data,
  variables,
  by = dplyr::group_vars(data),
  conf.level = 0.95,
  strata,
  weights = NULL,
  max.iterations = 10,
  method = c("waldcc", "wald", "clopper-pearson", "wilson", "wilsoncc", "strat_wilson",
            "strat_wilsoncc", "agresti-coull", "jeffreys")
)
```

## Arguments

data	(data.frame) a data frame
variables	(tidy-select) columns to include in summaries. Columns must be class <logical> or <numeric> values coded as c(0, 1).
by	(tidy-select) columns to stratify calculations by
conf.level	(numeric) a scalar in (0, 1) indicating the confidence level. Default is 0.95
strata, weights, max.iterations	arguments passed to proportion_ci_strat_wilson(), when method='strat_wilson'
method	(string) string indicating the type of confidence interval to calculate. Must be one of 'waldcc', 'wald', 'clopper-pearson', 'wilson', 'wilsoncc', 'strat_wilson', 'strat_wilsoncc', 'agresti-coull', 'jeffreys'. See ?proportion_ci for details.

## Value

an ARD data frame

## Examples

```
ard_proportion_ci(mtcars, variables = c(vs, am), method = "wilson")
```

---

ard_regression	<i>Regression ARD</i>
----------------	-----------------------

---

**Description**

Function takes a regression model object and converts it to a ARD structure using the `broom.helpers` package.

**Usage**

```
ard_regression(x, ...)  
  
## Default S3 method:  
ard_regression(x, tidy_fun = broom.helpers::tidy_with_broom_or_parameters, ...)
```

**Arguments**

x	regression model object
...	Arguments passed to <code>broom.helpers::tidy_plus_plus()</code>
tidy_fun	(function) a tidier. Default is <code>broom.helpers::tidy_with_broom_or_parameters</code>

**Value**

data frame

**Examples**

```
lm(AGE ~ ARM, data = cards::ADSL) |>  
  ard_regression(add_estimate_to_reference_rows = TRUE)
```

---

ard_ttest	<i>ARD t-test</i>
-----------	-------------------

---

**Description**

Analysis results data for paired and non-paired t-tests.

**Usage**

```
ard_ttest(data, by, variable, ...)  
  
ard_paired_ttest(data, by, variable, id, ...)
```

**Arguments**

data	(data.frame) a data frame. See below for details.
by	(tidy-select) column name to compare by
variable	(tidy-select) column name to be compared
...	arguments passed to <code>t.test(...)</code>
id	(tidy-select) column name of the subject or participant ID

**Details**

For the `ard_ttest()` function, the data is expected to be one row per subject. The data is passed as `t.test(data[[variable]] ~ data[[by]], paired = FALSE, ...)`.

For the `ard_paired_ttest()` function, the data is expected to be one row per subject per by level. Before the t-test is calculated, the data are reshaped to a wide format to be one row per subject. The data are then passed as `t.test(x = data_wide[[<by level 1>]], y = data_wide[[<by level 2>]], paired = TRUE, ...)`.

**Value**

ARD data frame

**Examples**

```
cards::ADSL |>
  dplyr::filter(ARM %in% c("Placebo", "Xanomeline High Dose")) |>
  ard_ttest(by = ARM, variable = AGE)

# constructing a paired data set,
# where patients receive both treatments
cards::ADSL[c("ARM", "AGE")] |>
  dplyr::filter(ARM %in% c("Placebo", "Xanomeline High Dose")) |>
  dplyr::mutate(.by = ARM, USUBJID = dplyr::row_number()) |>
  dplyr::arrange(USUBJID, ARM) |>
  ard_paired_ttest(by = ARM, variable = AGE, id = USUBJID)
```

---

ard\_wilcoxtest

*ARD Wilcoxon Rank-Sum Test*


---

**Description**

Analysis results data for paired and non-paired Wilcoxon Rank-Sum tests.



**Usage**

```
ard_wilcoxtest(data, by, variable, ...)
```

```
ard_paired_wilcoxtest(data, by, variable, id, ...)
```

**Arguments**

data	(data.frame) a data frame. See below for details.
by	(tidy-select) column name to compare by.
variable	(tidy-select) column name to be compared.
...	arguments passed to wilcox.test(...)
id	(tidy-select) column name of the subject or participant ID.

**Details**

For the `ard_wilcoxtest()` function, the data is expected to be one row per subject. The data is passed as `wilcox.test(data[[variable]] ~ data[[by]], paired = FALSE, ...)`.

For the `ard_paired_wilcoxtest()` function, the data is expected to be one row per subject per level. Before the test is calculated, the data are reshaped to a wide format to be one row per subject.

The data are then passed as `wilcox.test(x = data_wide[[<by level 1>]], y = data_wide[[<by level 2>]], paired = TRUE, ...)`.

**Value**

ARD data frame

**Examples**

```
cards::ADSL |>
  dplyr::filter(ARM %in% c("Placebo", "Xanomeline High Dose")) |>
  ard_wilcoxtest(by = "ARM", variable = "AGE")

# constructing a paired data set,
# where patients receive both treatments
cards::ADSL[c("ARM", "AGE")] |>
  dplyr::filter(ARM %in% c("Placebo", "Xanomeline High Dose")) |>
  dplyr::mutate(.by = ARM, USUBJID = dplyr::row_number()) |>
  dplyr::arrange(USUBJID, ARM) |>
  ard_paired_wilcoxtest(by = ARM, variable = AGE, id = USUBJID)
```

**Description**

Functions to calculate different proportion confidence intervals for use in `ard_proportion()`.

**Usage**

```

proportion_ci_wald(x, conf.level = 0.95, correct = FALSE)

proportion_ci_wilson(x, conf.level = 0.95, correct = FALSE)

proportion_ci_clopper_pearson(x, conf.level = 0.95)

proportion_ci_agresti_coull(x, conf.level = 0.95)

proportion_ci_jeffreys(x, conf.level = 0.95)

proportion_ci_strat_wilson(
  x,
  strata,
  weights = NULL,
  conf.level = 0.95,
  max.iterations = 10L,
  correct = FALSE
)

```

**Arguments**

<code>x</code>	vector of a binary values, i.e. a logical vector, or numeric with values $c(0, 1)$
<code>conf.level</code>	(numeric) a scalar in $(0, 1)$ indicating the confidence level. Default is 0.95
<code>correct</code>	(flag) include the continuity correction. For further information, see for example <a href="#">stats::prop.test()</a> .
<code>strata</code>	(factor) variable with one level per stratum and same length as <code>x</code> .
<code>weights</code>	(numeric or NULL) weights for each level of the strata. If NULL, they are estimated using the iterative algorithm that minimizes the weighted squared length of the confidence interval.
<code>max.iterations</code>	(count) maximum number of iterations for the iterative procedure used to find estimates of optimal weights.

**Value**

Confidence interval of a proportion.

**Functions**

- `proportion_ci_wald()`: Calculates the Wald interval by following the usual textbook definition for a single proportion confidence interval using the normal approximation.
- `proportion_ci_wilson()`: Calculates the Wilson interval by calling `stats::prop.test()`. Also referred to as Wilson score interval.
- `proportion_ci_clopper_pearson()`: Calculates the Clopper-Pearson interval by calling `stats::binom.test()`. Also referred to as the exact method.
- `proportion_ci_agresti_coull()`: Calculates the Agresti-Coull interval (created by Alan Agresti and Brent Coull) by (for 95% CI) adding two successes and two failures to the data and then using the Wald formula to construct a CI.
- `proportion_ci_jeffreys()`: Calculates the Jeffreys interval, an equal-tailed interval based on the non-informative Jeffreys prior for a binomial proportion.
- `proportion_ci_strat_wilson()`: Calculates the stratified Wilson confidence interval for unequal proportions as described in Xin YA, Su XG. Stratified Wilson and Newcombe confidence intervals for multiple binomial proportions. *Statistics in Biopharmaceutical Research*. 2010;2(3).

**Examples**

```
x <- c(
  TRUE, TRUE, TRUE, TRUE, TRUE,
  FALSE, FALSE, FALSE, FALSE, FALSE
)

proportion_ci_wald(x, conf.level = 0.9)
proportion_ci_wilson(x, correct = TRUE)
proportion_ci_clopper_pearson(x)
proportion_ci_agresti_coull(x)
proportion_ci_jeffreys(x)
# Stratified Wilson confidence interval with unequal probabilities

set.seed(1)
rsp <- sample(c(TRUE, FALSE), 100, TRUE)
strata_data <- data.frame(
  "f1" = sample(c("a", "b"), 100, TRUE),
  "f2" = sample(c("x", "y", "z"), 100, TRUE),
  stringsAsFactors = TRUE
)
strata <- interaction(strata_data)
n_strata <- ncol(table(rsp, strata)) # Number of strata

proportion_ci_strat_wilson(
  x = rsp, strata = strata,
  conf.level = 0.90
)
```

```
# Not automatic setting of weights
proportion_ci_strat_wilson(
  x = rsp, strata = strata,
  weights = rep(1 / n_strata, n_strata),
  conf.level = 0.90
)
```

# Index

ard\_chisqtest, [2](#)  
ard\_fishertest, [3](#)  
ard\_kruskaltest, [3](#)  
ard\_mcnemartest, [4](#)  
ard\_moodtest, [5](#)  
ard\_paired\_ttest (ard\_ttest), [7](#)  
ard\_paired\_wilcoxtest (ard\_wilcoxtest),  
[8](#)  
ard\_proportion\_ci, [6](#)  
ard\_regression, [7](#)  
ard\_ttest, [7](#)  
ard\_wilcoxtest, [8](#)

broom.helpers::tidy\_with\_broom\_or\_parameters,  
[7](#)

proportion\_ci, [10](#)  
proportion\_ci\_agresti\_coull  
(proportion\_ci), [10](#)  
proportion\_ci\_clopper\_pearson  
(proportion\_ci), [10](#)  
proportion\_ci\_jeffreys (proportion\_ci),  
[10](#)  
proportion\_ci\_strat\_wilson  
(proportion\_ci), [10](#)  
proportion\_ci\_wald (proportion\_ci), [10](#)  
proportion\_ci\_wilson (proportion\_ci), [10](#)

stats::binom.test(), [11](#)  
stats::prop.test(), [10, 11](#)