

# Package: walrus (via r-universe)

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**Type** Package

**Title** Robust Statistical Methods

**Version** 2.0.0

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**Description** A toolbox of common robust statistical tests, including robust descriptives, robust t-tests, and robust ANOVA. It is also available as a module for 'jamovi' (see <<https://www.jamovi.org>> for more information). Walrus is based on the WRS2 package by Patrick Mair, which is in turn based on the scripts and work of Rand Wilcox. These analyses are described in depth in the book 'Introduction to Robust Estimation & Hypothesis Testing'.

**License** GPL-3

**Encoding** UTF-8

**LazyData** true

**Imports** WRS2, ggplot2, jmvcore (>= 2.2), R6

**Suggests** MASS

**URL** <https://github.com/jamovi/walrus>

**BugReports** <https://github.com/jamovi/walrus/issues>

**RoxygenNote** 6.1.1

**Repository** <https://jamovi.r-universe.dev>

**RemoteUrl** <https://github.com/jamovi/walrus>

**RemoteRef** HEAD

**RemoteSha** ef1a8baea2eb28d7d058ed37689fd25071be7f6a

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walrus-package	<i>Walrus</i>
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## Description

A toolbox of common robust statistical tests, including robust descriptives, robust t-tests, and robust ANOVA. It is also available as a module for 'jamovi' (see [www.jamovi.org](http://www.jamovi.org) for more information). Walrus is based on the WRS2 package by Patrick Mair, which is in turn based on the scripts and work of Rand Wilcox. These analyses are described in depth in the book [Introduction to Robust Estimation & Hypothesis Testing](#).

## Details

Box & Violin Plots	<code>rplots()</code>
Robust Descriptives	<code>rdesc()</code>
Robust Independent Samples T-Test	<code>rttestIS()</code>
Robust Paired Samples T-Test	<code>rttestIS()</code>
Robust ANOVA	<code>ranova()</code>

Ravi:  
"Should we create a logo for walrus?"

Jonathon:  
"Yeah, I guess. Maybe a walrus, or a skewed distribution?  
Bonus points if it somehow contains both."

Ravi gets bonus points



**See Also**

Useful links:

- <https://github.com/jamovi/walrus>
- Report bugs at <https://github.com/jamovi/walrus/issues>

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ranova

*Robust ANOVA*


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**Description**

Robust Analysis of Variance

**Usage**

```
ranova(
  data,
  dep,
  factors = NULL,
  method = "trim",
  ph = FALSE,
  tr = 0.2,
  est = "mom",
  nboot = 599,
  dist = "proj"
)
```

**Arguments**

data	the data as a data frame
dep	a string naming the dependent variable from data; the variable must be numeric
factors	a vector of strings naming the fixed factors from data
method	'median', 'trim' (default) or 'boot'; the method to use, median, trimmed means, or bootstrapped
ph	TRUE or FALSE (default), provide post hoc tests
tr	a number between 0 and 0.5, (default: 0.2), the proportion of measurements to trim from each end, when using the trim and bootstrap methods
est	'onestep', 'mom' (default) or 'median', the M-estimator to use; One-step, Modified one-step or Median respectively
nboot	a number (default: 599) specifying the number of bootstrap samples to use when using the bootstrap method
dist	'maha' or 'proj' (default), whether to use Mahalanobis or Projection distances respectively

**Value**

A results object containing:

results\$main	the table of ANOVA results
results\$phs	the table of posthoc tests

Tables can be converted to data frames with `asDF` or `as.data.frame`. For example:

```
results$main$asDF
as.data.frame(results$main)
```

**Examples**

```
data('goggles', package='WRS2')

ranova(goggles,
       dep = 'attractiveness',
       factors = c('gender', 'alcohol'),
       ph = TRUE)

#
# ROBUST ANOVA
#
# Robust ANOVA
# -----
#               Q           p
# -----
# gender           1.67    0.209
# alcohol          48.28    0.001
# gender:alcohol   26.26    0.001
# -----
# Note. Method of trimmed means,
# trim level 0.2
#
# POST HOC TESTS
#
# Post Hoc Tests - gender
# -----
#               psi-hat    p      Lower    Upper
# -----
# Female   Male    10.0    0.209    -6.00    26.0
# -----
#
# Post Hoc Tests - alcohol
# -----
#               psi-hat    p      Lower    Upper
# -----
# None      2 Pints   -3.33    0.611   -20.5    13.8
```

```

#   None      4 Pints   35.83   < .001   19.3   52.3
#   2 Pints   4 Pints   39.17   < .001   22.5   55.9
# -----
#

```

rdesc

*Robust Descriptives***Description**

Robust Descriptives

**Usage**

```

rdesc(
  data,
  vars,
  splitBy = NULL,
  mean = TRUE,
  trim = TRUE,
  tr = 0.2,
  win = FALSE,
  wl = 0.2,
  mest = FALSE,
  bend = 1.28,
  med = FALSE
)

```

**Arguments**

<code>data</code>	the data as a data frame
<code>vars</code>	a vector of strings naming the variables in data of interest
<code>splitBy</code>	a string naming the variable in data to split the data by
<code>mean</code>	TRUE (default) or FALSE, provide a 'normal' arithmetic mean
<code>trim</code>	TRUE (default) or FALSE, provide a trimmed mean
<code>tr</code>	a number between 0 and 0.5 (default: 0.2); the proportion of measurements to trim from each end when producing trimmed means
<code>win</code>	TRUE or FALSE (default), provide a 'Winsorized' mean
<code>wl</code>	a number between 0 and 0.5 (default: 0.2); the level of 'winsorizing' when producing winsorized means
<code>mest</code>	TRUE or FALSE (default), provide an 'M-estimated' value
<code>bend</code>	a number (default: 1.28), the bending constant to use when using M-estimators
<code>med</code>	TRUE or FALSE (default), provide medians

**Value**

A results object containing:

`results$table`                      the table of descriptives

Tables can be converted to data frames with `asDF` or `as.data.frame`. For example:

```
results$table$asDF
as.data.frame(results$table)
```

**Examples**

```
data('eurosoccer', package='WRS2')

SpainGermany <- subset(eurosoccer, eurosoccer$League == 'Spain' | eurosoccer$League == 'Germany')
SpainGermany <- droplevels(SpainGermany)

walrus::rdesc(
  data = SpainGermany,
  vars = "GoalsGame",
  splitBy = "League",
  med = TRUE)

#
# ROBUST DESCRIPTIVES
#
# Robust Descriptives
# -----
#
#                               SE
# -----
#   GoalsGame   Germany   Mean      1.46   0.105
#               Germany   Trimmed mean 1.45   0.1341
#               Germany   Median      1.43   0.1599
#
#               Spain     Mean      1.45   0.101
#               Spain     Trimmed mean 1.33   0.0601
#               Spain     Median      1.30   0.0766
# -----
#
```

---

rplots

*Box & Violin Plots*

---

**Description**

Box & Violin Plots

**Usage**

```
rplots(  
  data,  
  vars,  
  splitBy = NULL,  
  violin = TRUE,  
  boxplot = FALSE,  
  dot = TRUE,  
  dotType = "stack"  
)
```

**Arguments**

data	the data as a data frame
vars	a vector of strings naming the variables in data of interest
splitBy	a string naming the variable in data to split the data by
violin	TRUE (default) or FALSE, provide violin plots
boxplot	TRUE or FALSE (default), provide box plots
dot	TRUE (default) or FALSE, plot each measurement as a dot
dotType	'jitter' or 'stack' (default); whether data dots are jittered or stacked

**Value**

A results object containing:

results\$plots                      an array of images

**Examples**

```
data('eurosoccer', package='WRS2')  
  
# violin plots  
  
walrus::rplots(  
  data = eurosoccer,  
  vars = "GoalsGame",  
  splitBy = "League")  
  
# box plots  
  
walrus::rplots(  
  data = eurosoccer,  
  vars = "GoalsGame",  
  splitBy = "League",  
  violin = FALSE,  
  boxplot = TRUE,  
  dot = FALSE)
```

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 rttestIS

*Robust Independent Samples T-Test*


---

### Description

Robust Independent Samples T-Test

### Usage

```
rttestIS(
  data,
  deps,
  group,
  yuen = TRUE,
  tr = 0.2,
  mest = FALSE,
  method = "mom",
  yuenbt = FALSE,
  nboot = 599,
  md = FALSE,
  ci = FALSE,
  es = FALSE,
  esci = FALSE
)
```

### Arguments

data	the data as a data frame
deps	a vector of strings naming the dependent variables in data
group	a string naming the grouping variable in data; must have 2 levels
yuen	TRUE (default) or FALSE, use the Yuen's trim method
tr	a number between 0 and 0.5, (default: 0.2), the proportion of measurements to trim from each end, when using the trim and bootstrap methods
mest	TRUE or FALSE (default), use an M-estimator
method	'onestep', 'mom' (default) or 'median', the M-estimator to use; One-step, Modified one-step or Median respectively
yuenbt	TRUE or FALSE (default), use the Yuen's bootstrap method
nboot	a number (default: 599) specifying the number of bootstrap samples to use when using the bootstrap method
md	TRUE or FALSE (default), provide the mean difference
ci	TRUE or FALSE (default), provide a 95% confidence interval on the mean difference
es	TRUE or FALSE (default), provide the effect-size
esci	TRUE or FALSE (default), provide a 95% confidence interval on the effect-size



**Arguments**

<code>data</code>	the data as a data frame
<code>pairs</code>	a list of lists specifying the pairs of measurement in data
<code>tr</code>	a number between 0 and 0.5, (default: 0.2), the proportion of measurements to trim from each end, when using the trim and bootstrap methods
<code>md</code>	TRUE or FALSE (default), provide means and standard errors
<code>es</code>	TRUE or FALSE (default), provide effect sizes
<code>ci</code>	TRUE or FALSE (default), provide confidence intervals

**Value**

A results object containing:

`results$tttest` the table of t-test results

Tables can be converted to data frames with `asDF` or `as.data.frame`. For example:

```
results$tttest$asDF
as.data.frame(results$tttest)
```

**Examples**

```
data(anorexia, package='MASS')
anorexiaFT <- subset(anorexia, subset = Treat == "FT")

rttestPS(anorexiaFT,
  pairs = list(
    list(i1 = 'Prewt', i2 = 'Postwt')))

#
# ROBUST PAIRED SAMPLES T-TEST
#
# Robust Paired Samples T-Test
# -----
#               t      df      p
# -----
# Prewt  Postwt  -3.83   10.0   0.003
# -----
#
#
```

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