

# Package ‘peramo’

May 31, 2023

**Type** Package

**Title** Permutation Tests for Randomization Model

**Version** 0.1.3

**Description** Perform permutation-based hypothesis testing for randomized experiments as suggested in Ludbrook & Dudley (1998) <[doi:10.2307/2685470](https://doi.org/10.2307/2685470)> and Ernst (2004) <[doi:10.1214/088342304000000396](https://doi.org/10.1214/088342304000000396)>, introduced in Pham et al. (2022) <[doi:10.1016/j.chemosphere.2022.136736](https://doi.org/10.1016/j.chemosphere.2022.136736)>.

**Depends** R (>= 4.3.0)

**Imports** magrittr (>= 2.0.3), dplyr (>= 1.1.2), lme4 (>= 1.1-33), parameters (>= 0.21.0), emmeans (>= 1.8.6), stats, utils

**Suggests** multcompView (>= 0.1-9)

**BugReports** <https://github.com/phamdn/peramo/issues>

**License** GPL-3

**Encoding** UTF-8

**RoxygenNote** 7.2.3

**LazyData** true

**NeedsCompilation** no

**Author** Duy Nghia Pham [aut, cre] (<<https://orcid.org/0000-0003-1349-1710>>),  
Inna M. Sokolova [ths] (<<https://orcid.org/0000-0002-2068-4302>>)

**Maintainer** Duy Nghia Pham <[nghiapham@yandex.com](mailto:nghiapham@yandex.com)>

**Repository** CRAN

**Date/Publication** 2023-05-31 16:20:02 UTC

## R topics documented:

peramo-package . . . . .	2
AB . . . . .	3
ctm . . . . .	4
difffcalc . . . . .	4
ensulizole . . . . .	5

nolessor . . . . .	5
owl . . . . .	6
owlStat . . . . .	7
twl . . . . .	8
twlStat . . . . .	9
tw_complex . . . . .	10
XY . . . . .	12
<b>Index</b>	<b>13</b>

---

peramo-package	<i>peramo: Permutation Tests for Randomization Model</i>
----------------	--

---

## Description

Perform permutation-based hypothesis testing for randomized experiments as suggested in Ludbrook & Dudley (1998) [doi:10.2307/2685470](https://doi.org/10.2307/2685470) and Ernst (2004) [doi:10.1214/088342304000000396](https://doi.org/10.1214/088342304000000396), introduced in Pham et al. (2022) [doi:10.1016/j.chemosphere.2022.136736](https://doi.org/10.1016/j.chemosphere.2022.136736).

## Copyright

peramo: Permutation Tests for Randomization Model.  
Copyright (C) 2022-2023 Duy Nghia Pham & Inna M. Sokolova

peramo is free software: you can redistribute it and/or modify it under the terms of the GNU General Public License as published by the Free Software Foundation, either version 3 of the License, or (at your option) any later version.

peramo is distributed in the hope that it will be useful, but WITHOUT ANY WARRANTY; without even the implied warranty of MERCHANTABILITY or FITNESS FOR A PARTICULAR PURPOSE. See the GNU General Public License for more details.

You should have received a copy of the GNU General Public License along with peramo. If not, see <https://www.gnu.org/licenses/>.

## Author(s)

Duy Nghia Pham & Inna M. Sokolova

---

AB *Two-Group Permutation Test*

---

**Description**

AB performs A/B testing for two-group experiments.

**Usage**

```
AB(a, b, rand = 9999, seed = 1)
```

**Arguments**

a	the measurement of responses of the first group.
b	the measurement of responses of the second group.
rand	an integer, the number of randomization samples. The default value is 9999.
seed	an integer, the seed for random number generation. Setting a seed ensures the reproducibility of the result. See <a href="#">set.seed</a> for more details.

**Value**

AB returns a one-row data frame with 6 columns:

nA	the sample size of the first group.
mean.A	the mean responses of the first group.
nB	the sample size of the second group.
mean.B	the mean responses of the second group.
mean.diff	the difference between two mean responses.
pval	the p-value.

**References**

Ernst, M. D. (2004). Permutation Methods: A Basis for Exact Inference. *Statistical Science*, 19(4), 676–685. [doi:10.1214/088342304000000396](https://doi.org/10.1214/088342304000000396).

**Examples**

```
AB(c(19, 22, 25, 26), c(23, 33, 40))
```

---

ctm

*Biomarker Responses of the Ragworms to Copper and Warming*

---

**Description**

Biomarker Responses of the Ragworms to Copper and Warming

**Usage**

ctm\_Cu

ctm\_worm

**Format**

An object of class data.frame with 60 rows and 7 columns.

An object of class data.frame with 210 rows and 26 columns.

---

diffcalc

*Calculate the Differences between Means*

---

**Description**

Calculate the Differences between Means

**Usage**

```
diffcalc(vec, control)
```

**Arguments**

vec            a numeric vector, the mean responses.

control        a logical, whether the control group exists.

---

ensulizole

*Biomarker Responses of the Blue Mussels to Organic UV Filters*

---

**Description**

Biomarker Responses of the Blue Mussels to Organic UV Filters

**Usage**

mussel\_SoS

mussel\_gill

mussel\_digest

**Format**

An object of class `data.frame` with 15 rows and 7 columns.

An object of class `data.frame` with 120 rows and 24 columns.

An object of class `data.frame` with 120 rows and 24 columns.

**Source**

Pham, D. N., Sokolov, E. P., Falfushynska, H., & Sokolova, I. M. (2022). Gone with sunscreens: Responses of blue mussels (*Mytilus edulis*) to a wide concentration range of a UV filter ensulizole. *Chemosphere*, 309, 136736. doi:10.1016/j.chemosphere.2022.136736.

---

nolesser

*Compare the Differences with Critical Values*

---

**Description**

Compare the Differences with Critical Values

**Usage**

```
nolesser(obs, cric)
```

**Arguments**

`obs` a numeric, the observed difference.

`cric` a numeric, the critical values of maximum absolute differences.

owl

*One-Way Layout Permutation Test***Description**

owl performs the global test and multiple comparisons for single factor experiments.

**Usage**

```
owl(df, rand = 9999, alpha.post = 0.05, type.post = "control", seed = 1)
```

**Arguments**

df	a data frame with the name of experimental groups as the first column and the measurement of responses as the remaining columns.
rand	an integer, the number of randomization samples. The default value is 9999.
alpha.post	a numeric, the Type I error rate for multiple comparisons. The default value is 0.05.
type.post	the way of multiple comparisons, "all" for pairwise comparisons or "control" for only comparisons with the control group.
seed	an integer, the seed for random number generation. Setting a seed ensures the reproducibility of the result. See <a href="#">set.seed</a> for more details.

**Details**

The first name appearing in the first column will determine the control group. The other names will be treatment groups.

**Value**

owl returns a list with 9 components:

n.obs	the sample sizes.
avg.obs	the mean responses.
T.obs	the T statistic for global test.
pval	the p-value for global test.
pval.round	the reported form of p-value.
main.test	the strength of evidence against the null hypothesis.
d.multi.obs	the differences in means for multiple comparisons.
mad.cric	the critical value of maximum absolute differences in means.
post.test	TRUE if the differences are significant.

## References

Ernst, M. D. (2004). Permutation Methods: A Basis for Exact Inference. *Statistical Science*, 19(4), 676–685. doi:10.1214/088342304000000396.

Muff, S., Nilsen, E. B., O’Hara, R. B., & Nater, C. R. (2022). Rewriting results sections in the language of evidence. *Trends in Ecology & Evolution*, 37(3), 203–210. doi:10.1016/j.tree.2021.10.009.

## Examples

```
ernst2004 <- data.frame(
  group = factor(rep(c("style1", "style2", "style3"), each = 5 ),
  levels = c("style1", "style2", "style3")),
  speed = c( 135,91,111,87, 122, 175,130,514,283, NA,105,147,159,107,194))
owl(ernst2004, type.post = "all")
```

---

owlStat

*Test Statistics for One-Way Layout Permutation Test*

---

## Description

owlStat computes statistics for owl. This is not meant to be called directly.

## Usage

```
owlStat(lov, env = parent.frame())
```

## Arguments

lov	a list of vectors, responses by experimental groups.
env	an environment, to access outer scope variables.

## Value

owlStat returns a list with 5 components:

n	the sample sizes.
avg	the mean responses.
T	the T statistic for global test.
d.multi	the differences in means for multiple comparisons.
mad	the maximum absolute differences in means.

.

## References

Ernst, M. D. (2004). Permutation Methods: A Basis for Exact Inference. *Statistical Science*, 19(4), 676–685. doi:10.1214/088342304000000396.

---

twl	<i>Two-Way Layout Permutation Test</i>
-----	--

---

### Description

twl performs the global test and multiple comparisons for two-factor experiments.

### Usage

```
twl(  
  df,  
  rand = 4999,  
  seed = 1,  
  mult = FALSE,  
  simple = TRUE,  
  control = TRUE,  
  alpha = 0.05  
)
```

### Arguments

df	a data frame with the first and second columns containing the levels of the two main factors and the third column containing the measurement of responses.
rand	an integer, the number of randomization samples. The default value is 4999.
seed	an integer, the seed for random number generation. Setting a seed ensures the reproducibility of the result. See <a href="#">set.seed</a> for more details.
mult	a logical, whether to perform multiple comparisons.
simple	a logical, whether to perform comparisons for simple effects.
control	a logical, whether to perform only comparisons with the control group.
alpha	a numeric, the Type I error rate for multiple comparisons. The default value is 0.05.

### Details

The first levels appearing in the first and second columns will determine the control groups (if any). The other levels will be treatment groups.

### Value

twl returns a list with possible components:

n, n.main1, and n.main2	the sample sizes.
avg, avg.main1, and avg.main2	the mean responses.



`Fs` the F statistics, p-values, reported form of p-value, and strength of evidence against the null hypotheses.

`d.main1sub` and `d.main2sub` or `d.main1` and `d.main2`  
the differences in means for multiple comparisons.

`mad.main1sub.cric` and `mad.main2sub.cric` or `mad.main1.cric` and `mad.main2.cric`  
the critical value of maximum absolute differences in means.

`mult.test.main1sub` and `mult.test.main2sub` or `mult.test.main1` and `mult.test.main2`  
TRUE if the differences are significant.

## References

Manly, B. F. J. (2007). Randomization, bootstrap, and Monte Carlo methods in biology (3rd ed). Chapman & Hall/ CRC.

Ernst, M. D. (2004). Permutation Methods: A Basis for Exact Inference. *Statistical Science*, 19(4), 676–685. doi:10.1214/088342304000000396.

Muff, S., Nilsen, E. B., O'Hara, R. B., & Nater, C. R. (2022). Rewriting results sections in the language of evidence. *Trends in Ecology & Evolution*, 37(3), 203–210. doi:10.1016/j.tree.2021.10.009.

Motulsky, H. (2020). GraphPad Statistics Guide. GraphPad Software Inc. <https://www.graphpad.com/guides/prism/latest/statistics/index.htm>.

## Examples

```
manly2007 <- data.frame(
  month = factor(rep(c("jun", "jul", "aug", "sep"), each = 6 ),
  levels = c("jun", "jul", "aug", "sep")),
  size = factor(rep(c("small", "large"), each = 3, times = 4),
  levels = c("small", "large")),
  consume = c( 13,242,105,182,21,7,8,59,20,24,312,68,515,488,88,460,1223,990,18,44,21,140,40,27))
twl(manly2007)
twl(manly2007, mult = TRUE, simple = TRUE, control = FALSE)
#might take more than 5s in some machines
```

---

twlStat

*Test Statistics for Two-Way Layout Permutation Test*

---

## Description

twlStat computes statistics for twl. This is not meant to be called directly.

## Usage

```
twlStat(df, env = parent.frame())
```

**Arguments**

df	a data frame with the levels of the two main factors as the first and second columns and the measurement of responses as the third column.
env	an environment, to access outer scope variables.

**Value**

twlStat returns a list with at least 4 components:

Fs	the F statistics for global test.
F.main1 and F.main2	the F statistics for the first main factor and the second main factor.
F.int	the F statistic for the interaction.

In case of multiple comparisons, additional components are:

avg or avg.main1 and avg.main2	the mean responses for multiple comparisons.
d.main1sub and d.main2sub or d.main1 and d.main2	the differences in means.
mad.main1sub and mad.main2sub or mad.main1 and mad.main2	the maximum absolute differences in means.

**References**

Manly, B. F. J. (2007). Randomization, bootstrap, and Monte Carlo methods in biology (3rd ed). Chapman & Hall/ CRC.

---

tw\_complex

*Permutation Test for Two-Way Layout with Extra Factors*


---

**Description**

tw\_complex performs the permutation test for ANOVA of two-factor experiments with complex design.

**Usage**

```
tw_complex(df, res, mains, nested, nuis, seed = 1, rand = 1999, emm = FALSE)
```

## Arguments

df	a data frame with at least three columns.
res	a character string, name of response variable.
mains	two character strings, names of two main factors.
nested	(optional) a character string, name of the nested factor.
nuis	(optional) a character string, name of the nuisance factor.
seed	an integer, the seed for random number generation. Setting a seed ensures the reproducibility of the result. See <a href="#">set.seed</a> for more details.
rand	an integer, the number of randomization samples. The default value is 1999.
emm	a logical, whether to compute estimated marginal means.

## Details

res, mains, nested, and nuis refer to column names in df. While nuis column must be a numeric vector, mains and nested columns must be factors. res can be a numeric or logical vector.

tw\_complex currently support linear models with only mains, generalized linear mixed-effects models with mains and nested, and linear mixed-effects models with mains, nested, and nuis.

## Value

tw\_complex returns a list with 3 main components:

lm, glmer, or lmer	model results.
anova	anova table.
perm	permutation test results with F-statistics, p-values, and strength of evidence.

## References

Manly, B. F. J. (2007). Randomization, bootstrap, and Monte Carlo methods in biology (3rd ed). Chapman & Hall/ CRC.

Ernst, M. D. (2004). Permutation Methods: A Basis for Exact Inference. *Statistical Science*, 19(4), 676–685. doi:10.1214/088342304000000396.

Anderson, M., & Braak, C. T. (2003). Permutation tests for multi-factorial analysis of variance. *Journal of Statistical Computation and Simulation*, 73(2), 85–113. doi:10.1080/00949650215733.

## See Also

[lm](#), [glmer](#), and [lmer](#).

**Examples**

```
tw_complex(df = subset(ctm_Cu, run == "Jan",
  select = c("copper", "temp", "sediment")),
  res = "sediment",
  mains = c("copper", "temp"))
```

```
#might take more than 5s in some machines
```

---

 XY

---

*Permutation Test for Correlation Between Paired Samples*


---

**Description**

XY performs permutation test on correlation coefficients.

**Usage**

```
XY(a, b, rand = 9999, seed = 1, method = c("pearson", "kendall", "spearman"))
```

**Arguments**

a	a numeric vector, the first variable.
b	a numeric vector, the second variable.
rand	an integer, the number of randomization samples. The default value is 9999.
seed	an integer, the seed for random number generation. Setting a seed ensures the reproducibility of the result. See <a href="#">set.seed</a> for more details.
method	correlation coefficient, "pearson", "kendall", or "spearman".

**Value**

XY returns an one-row data frame with 2 columns:

cor	the correlation coefficient.
pval	the p-value.

**See Also**

[cor.test](#)

**Examples**

```
with(subset(ctm_Cu, run == "Jan"), XY(sediment, porewater))
```

# Index

## \* datasets

ctm, 4

ensulizole, 5

AB, 3

cor.test, 12

ctm, 4

ctm\_Cu(ctm), 4

ctm\_worm(ctm), 4

diffcalc, 4

ensulizole, 5

glmer, 11

lm, 11

lmer, 11

mussel\_digest(ensulizole), 5

mussel\_gill(ensulizole), 5

mussel\_SoS(ensulizole), 5

nolesser, 5

owl, 6

owlStat, 7

peramo-package, 2

set.seed, 3, 6, 8, 11, 12

tw\_complex, 10

twl, 8

twlStat, 9

XY, 12