

# Package ‘hce’

July 22, 2025

**Type** Package

**Title** Design and Analysis of Hierarchical Composite Endpoints

**Version** 0.8.0

**Description** Simulate and analyze hierarchical composite endpoints. Includes implementation for the kidney hierarchical composite endpoint as defined in Heerspink HL et al (2023) “Development and validation of a new hierarchical composite end point for clinical trials of kidney disease progression” (Journal of the American Society of Nephrology 34 (2): 2025–2038, <[doi:10.1681/ASN.000000000000243](https://doi.org/10.1681/ASN.000000000000243)>). Win odds, also called Wilcoxon-Mann-Whitney or success odds, is the main analysis method. Other win statistics (win probability, win ratio, net benefit) are also implemented in the univariate case, provided there is no censoring. The win probability analysis is based on the Brunner-Munzel test and uses the DeLong-DeLong-Clarke-Pearson variance estimator, as described by Brunner and Konietzschke (2025) in “An unbiased rank-based estimator of the Mann-Whitney variance including the case of ties” (Statistical Papers 66 (1): 20, <[doi:10.1007/s00362-024-01635-0](https://doi.org/10.1007/s00362-024-01635-0)>). Stratification and covariate adjustment are performed based on the methodology presented by Koch GG et al. in “Issues for covariance analysis of dichotomous and ordered categorical data from randomized clinical trials and non-parametric strategies for addressing them” (Statistics in Medicine 17 (15-16): 1863–92). For a review, see Gasparyan SB et al (2021) “Adjusted win ratio with stratification: Calculation methods and interpretation” (Statistical Methods in Medical Research 30 (2): 580–611, <[doi:10.1177/0962280220942558](https://doi.org/10.1177/0962280220942558)>).

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|      |  |
|------|--|
| ADET | <i>Event-Time dataset for kidney outcomes.</i> |
|------|--|

---

### Description

A dataset with multiple kidney outcomes over time scale outcomes of 1500 patients in the ADSL dataset.

### Usage

ADET

### Format

a data frame with 604 rows (events) and 6 variables:

**ID** patient identifiers, numeric

**AVAL** occurrence time of the event, numeric

**PARAM** name of the event, character

**PARAMCD** coded name of the event, character

**PARAMN** type of the event, outcomes 1-7, where a higher value means a better outcome, numeric

**TRTPN** treatment values, 1 Active or 2 Placebo, numeric

### Source

Heerspink HL et al "Development and validation of a new hierarchical composite endpoint for clinical trials of kidney disease progression." Journal of the American Society of Nephrology (2023): [doi:10.1681/ASN.000000000000243](https://doi.org/10.1681/ASN.000000000000243).

### Examples

```
head(ADET)
# Number of unique patients
length(unique(ADET$ID))
# Number of events per event type
barplot(table(ADET$PARAM))
```

---

|      |  |
|------|--|
| ADLB | <i>Laboratory dataset for Glomerular Filtration Rate (GFR) measurements.</i> |
|------|--|

---

### Description

A dataset of laboratory measurements of kidney function over time for the 1500 patients in the ADSL dataset.

### Usage

ADLB

### Format

a data frame with 13980 rows and 8 variables:

**ID** patient identifiers, numeric

**TRTPN** treatment values, 1 Active or 2 Placebo, numeric

**AVAL** measurement value, numeric

**ADAY** measurement day in the study, numeric

**AVISITN** hospital visit number, numeric

**PARAM** name of the event, GFR measurements, character

**PARAMCD** coded name of the event, GFR, character

**PARAMN** type of the event is set to 7 for all measurements, numeric

### Source

Heerspink HL et al "Development and validation of a new hierarchical composite endpoint for clinical trials of kidney disease progression." Journal of the American Society of Nephrology (2023): [doi:10.1681/ASN.000000000000243](https://doi.org/10.1681/ASN.000000000000243).

### Examples

```
head(ADLB)
```

---

|      |   |
|------|---|
| ADSL | <i>Baseline characteristics dataset of patients with kidney function assessments.</i> |
|------|---|

---

**Description**

A data frame with baseline characteristics for 1500 patients used to derive KHCE dataset.

**Usage**

ADSL

**Format**

a data frame with 1500 rows and 4 variables:

**ID** patient identifiers, numeric

**TRTPN** treatment values, 1 Active or 2 Placebo, numeric

**EGFRBL** Baseline GFR values of patients, numeric

**STRATAN** strata 1-4, higher value means a higher risk for kidney disease progression, numeric

**Source**

Heerspink HL et al "Development and validation of a new hierarchical composite endpoint for clinical trials of kidney disease progression." Journal of the American Society of Nephrology (2023): [doi:10.1681/ASN.000000000000243](https://doi.org/10.1681/ASN.000000000000243).

**Examples**

```
head(ADSL)
```

---

|        |   |
|--------|---|
| as_hce | <i>A generic function for coercing data structures to hce objects</i> |
|--------|---|

---

**Description**

A generic function for coercing data structures to hce objects

**Usage**

```
as_hce(x, ...)
```

**Arguments**

x an object used to select a method.

... additional parameters.

**Value**

an hce object.

**See Also**

[as\\_hce.data.frame\(\)](#).

**Examples**

```
### data frames
data(HCE1)
HCE <- as_hce(HCE1)
calcWINS(HCE)
```

---

|                   |   |
|-------------------|---|
| as_hce.data.frame | <i>Coerce a data frame to an hce object</i> |
|-------------------|---|

---

**Description**

Coerce a data frame to an hce object

**Usage**

```
## S3 method for class 'data.frame'
as_hce(x, ...)
```

**Arguments**

|     |                        |
|-----|------------------------|
| x   | a data frame.          |
| ... | additional parameters. |

**Value**

an hce object.

**See Also**

[as\\_hce\(\)](#), [as\\_hce.default\(\)](#).

**Examples**

```
KHCE <- as_hce(KHCE)
calcWO(KHCE)
```

---

|                |   |
|----------------|---|
| as_hce.default | <i>Coerce a data frame to an hce object</i> |
|----------------|---|

---

**Description**

Coerce a data frame to an hce object

**Usage**

```
## Default S3 method:  
as_hce(x, ...)
```

**Arguments**

|     |                        |
|-----|------------------------|
| x   | an object.             |
| ... | additional parameters. |

**Value**

an hce object.

**See Also**

[as\\_hce\(\)](#), [as\\_hce.data.frame\(\)](#).

**Examples**

```
dat <- KHCE  
class(dat) <- "moo" # non-existent class  
as_hce(dat) # tries to convert to an hce object
```

---

|          |  |
|----------|--|
| calcWINS | <i>A generic function for calculating win statistics</i> |
|----------|--|

---

**Description**

A generic function for calculating win statistics

**Usage**

```
calcWINS(x, ...)
```

**Arguments**

|     |  |
|-----|--|
| x   | an object used to select a method.                 |
| ... | further arguments passed to or from other methods. |

**Value**

a data frame containing calculated values.

**See Also**

[calcWINS.hce\(\)](#), [calcWINS.formula\(\)](#), [calcWINS.data.frame\(\)](#) methods.

---

calcWINS.data.frame     *Win statistics calculation using a data frame*

---

**Description**

Win statistics calculation using a data frame

**Usage**

```
## S3 method for class 'data.frame'
calcWINS(
  x,
  AVAL,
  TRTP,
  ref,
  alpha = 0.05,
  WOnull = 1,
  SE_WP_Type = c("biased", "unbiased"),
  ...
)
```

**Arguments**

|            |   |
|------------|---|
| x          | a data frame containing subject-level data.                                   |
| AVAL       | variable in the data with ordinal analysis values.                            |
| TRTP       | the treatment variable in the data.   |
| ref        | the reference treatment group.  |
| alpha      | 2-sided significance level. The default is 0.05.                              |
| WOnull     | the null hypothesis. The default is 1.  |
| SE_WP_Type | biased or unbiased standard error for win probability. The default is biased. |
| ...        | additional parameters.  |

**Details**

When `SE_WP_Type = "unbiased"`, the calculations for win proportion, net benefit, and win odds utilize the unbiased standard error from Brunner-Konietschke (2025) paper which is a reformulation of the original formula proposed by Bamber (1975).



**Value**

a list containing win statistics and their confidence intervals. It contains the following named data frames:

- summary a data frame containing number of wins, losses, and ties of the active treatment group and the overall number of comparisons.
- WP a data frame containing the win probability and its confidence interval.
- NetBenefit a data frame containing the net benefit and its confidence interval. This is just a  $2x-1$  transformation of WP and its CI.
- WO a data frame containing the win odds and its confidence interval.
- WR1 a data frame containing the win ratio and its confidence interval, using the transformed standard error of the gamma statistic.
- WR2 a data frame containing the win ratio and its confidence interval, using the standard error calculated using  $P_{ties}$ .
- gamma a data frame containing Goodman Kruskal's gamma and its confidence interval.
- SE a data frame containing standard errors used to calculate the Confidence intervals for win statistics.

**References**

The theory of win statistics is covered in the following papers:

- Win proportion and win odds confidence interval calculation:

Bamber D (1975) "The area above the ordinal dominance graph and the area below the receiver operating characteristic graph." *Journal of Mathematical Psychology* 12.4: 387-415. doi:10.1016/0022-2496(75)90001-2.

DeLong ER et al. (1988) "Comparing the Areas Under Two or More Correlated Receiver Operating Characteristic Curves: A Nonparametric Approach." *Biometrics* 44.3: 837-845. doi:10.2307/2531595.

Brunner E et al. (2021) "Win odds: an adaptation of the win ratio to include ties." *Statistics in Medicine* 40.14: 3367-3384. doi:10.1002/sim.8967.

Gasparian SB et al. (2021) "Adjusted win ratio with stratification: calculation methods and interpretation." *Statistical Methods in Medical Research* 30.2: 580-611. doi:10.1177/0962280220942558.

Gasparian SB et al. (2021) "Power and sample size calculation for the win odds test: application to an ordinal endpoint in COVID-19 trials." *Journal of Biopharmaceutical Statistics* 31.6: 765-787. doi:10.1080/10543406.2021.1968893.

Brunner E, Konietschke F. (2025) "An unbiased rank-based estimator of the Mann-Whitney variance including the case of ties." *Statistical Papers* 66.20. doi:10.1007/s00362-024-01635-0.

- Win ratio: the first CI utilizes the standard error derived from the gamma statistic standard error as outlined by:

Gasparyan SB, Kowalewski EK, Buenconsejo J, Koch GG. (2023) "Hierarchical Composite Endpoints in COVID-19: The DARE-19 Trial." In Case Studies in Innovative Clinical Trials, Chapter 7, 95–148. Chapman; Hall/CRC. doi:10.1201/9781003288640-7.

- Win ratio: the second CI utilizes the standard error presented by:

Yu RX, Ganju J. (2022) "Sample size formula for a win ratio endpoint." Statistics in Medicine 41.6: 950-63. doi:10.1002/sim.9297.

- Goodman Kruskal's gamma and CI: matches implementation in DescTools::GoodmanKruskalGamma() and based on:

Agresti A. (2002) Categorical Data Analysis. John Wiley & Sons, pp. 57-59. doi:10.1002/0471249688.

Brown MB, Benedetti JK. (1977) "Sampling Behavior of Tests for Correlation in Two-Way Contingency Tables." Journal of the American Statistical Association 72, 309-315. doi:10.1080/01621459.1977.10480995.

Goodman LA, Kruskal WH. (1954) "Measures of association for cross classifications." Journal of the American Statistical Association 49, 732-764. doi:10.1080/01621459.1954.10501231.

Goodman LA, Kruskal WH. (1963) "Measures of association for cross classifications III: Approximate sampling theory." Journal of the American Statistical Association 58, 310-364. doi:10.1080/01621459.1963.10500850.

## See Also

`calcWINS()`, `calcWINS.hce()`, `calcWINS.formula()`.

## Examples

```
calcWINS(x = COVID19b, AVAL = "GROUP", TRTP = "TRTP", ref = "Placebo")
## Biased vs unbiased
n0 <- 5; n1 <- 7; p0 <- 0.2; p1 <- 0.5; x <- 1:20; delta <- 0.5
WP0 <- (p1 - p0)/2 + 0.5
DAT <- NULL
for(i in x){
  dat <- data.frame(AVAL = c(rbinom(n1, size = 1, p1), rbinom(n0, size = 1, p0)),
    TRTP = c(rep("A", n1), rep("P", n0)))
  CL1 <- calcWINS(x = dat, AVAL = "AVAL", TRTP = "TRTP", ref = "P")$WP
  CL1$Type <- "biased"
  CL2 <- calcWINS(x = dat, AVAL = "AVAL", TRTP = "TRTP",
    ref = "P", SE_WP_Type = "unbiased")$WP
  CL2$Type <- "unbiased"
  DAT <- rbind(DAT, CL1, CL2)
}
WP <- DAT$WP[DAT$Type == "unbiased"]
```

```

plot(x, WP, pch = 19, xlab = "Simulations", ylab = "Win Probability", ylim = c(0., 1.1))
points(x + delta, WP, pch = 19)
arrows(x, DAT$LCL[DAT$Type == "unbiased"],
       x, DAT$UCL[DAT$Type == "unbiased"], angle = 90, code = 3, length = 0.05, "green")
arrows(x + delta, DAT$LCL[DAT$Type == "biased"],
       x + delta, DAT$UCL[DAT$Type == "biased"], angle = 90, code = 3, length = 0.05, col = "red")
abline(h = c(WP0, 1), col = "blue", lty = 3)
legend("bottomleft", legend = c("True WP", "Biased", "Unbiased"),
      col = c(4, 2, 3), lty = c(3, 1, 1), cex = 0.75)

```

---

|                  |  |
|------------------|--|
| calcWINS.formula | <i>Win statistics calculation using formula syntax</i> |
|------------------|--|

---

## Description

Win statistics calculation using formula syntax

## Usage

```

## S3 method for class 'formula'
calcWINS(x, data, ...)

```

## Arguments

|      |                             |
|------|-----------------------------|
| x    | an object of class formula. |
| data | a data frame.               |
| ...  | additional parameters.      |

## Value

a list containing win statistics and their confidence intervals. It contains the following named data frames:

- summary a data frame containing number of wins, losses, and ties of the active treatment group and the overall number of comparisons.
- WP a data frame containing the win probability and its confidence interval.
- NetBenefit a data frame containing the net benefit and its confidence interval. This is just a  $2x-1$  transformation of WP and its CI.
- WO a data frame containing the win odds and its confidence interval.
- WR1 a data frame containing the win ratio and its confidence interval, using the transformed standard error of the gamma statistic.
- WR2 a data frame containing the win ratio and its confidence interval, using the standard error calculated using  $P_{ties}$ .
- gamma a data frame containing Goodman Kruskal's gamma and its confidence interval.
- SE a data frame containing standard errors used to calculate the Confidence intervals for win statistics.

## References

The theory of win statistics is covered in the following papers:

- Win proportion and win odds confidence interval calculation:

Bamber D (1975) "The area above the ordinal dominance graph and the area below the receiver operating characteristic graph." *Journal of Mathematical Psychology* 12.4: 387-415. doi:[10.1016/0022-2496\(75\)90001-2](https://doi.org/10.1016/0022-2496(75)90001-2).

DeLong ER et al. (1988) "Comparing the Areas Under Two or More Correlated Receiver Operating Characteristic Curves: A Nonparametric Approach." *Biometrics* 44.3: 837-845. doi:[10.2307/2531595](https://doi.org/10.2307/2531595).

Brunner E et al. (2021) "Win odds: an adaptation of the win ratio to include ties." *Statistics in Medicine* 40.14: 3367-3384. doi:[10.1002/sim.8967](https://doi.org/10.1002/sim.8967).

Gasparyan SB et al. (2021) "Adjusted win ratio with stratification: calculation methods and interpretation." *Statistical Methods in Medical Research* 30.2: 580-611. doi:[10.1177/0962280220942558](https://doi.org/10.1177/0962280220942558).

Gasparyan SB et al. (2021) "Power and sample size calculation for the win odds test: application to an ordinal endpoint in COVID-19 trials." *Journal of Biopharmaceutical Statistics* 31.6: 765-787. doi:[10.1080/10543406.2021.1968893](https://doi.org/10.1080/10543406.2021.1968893).

Brunner E, Konietschke F. (2025) "An unbiased rank-based estimator of the Mann–Whitney variance including the case of ties." *Statistical Papers* 66.20. doi:[10.1007/s00362-024-01635-0](https://doi.org/10.1007/s00362-024-01635-0).

- Win ratio: the first CI utilizes the standard error derived from the gamma statistic standard error as outlined by:

Gasparyan SB, Kowalewski EK, Buenconsejo J, Koch GG. (2023) "Hierarchical Composite Endpoints in COVID-19: The DARE-19 Trial." In *Case Studies in Innovative Clinical Trials*, Chapter 7, 95–148. Chapman; Hall/CRC. doi:[10.1201/9781003288640-7](https://doi.org/10.1201/9781003288640-7).

- Win ratio: the second CI utilizes the standard error presented by:

Yu RX, Ganju J. (2022) "Sample size formula for a win ratio endpoint." *Statistics in Medicine* 41.6: 950-63. doi:[10.1002/sim.9297](https://doi.org/10.1002/sim.9297).

- Goodman Kruskal's gamma and CI: matches implementation in DescTools::GoodmanKruskalGamma() and based on:

Agresti A. (2002) *Categorical Data Analysis*. John Wiley & Sons, pp. 57-59. doi:[10.1002/0471249688](https://doi.org/10.1002/0471249688).

Brown MB, Benedetti JK. (1977) "Sampling Behavior of Tests for Correlation in Two-Way Contingency Tables." *Journal of the American Statistical Association* 72, 309-315. doi:[10.1080/01621459.1977.10480995](https://doi.org/10.1080/01621459.1977.10480995).

Goodman LA, Kruskal WH. (1954) "Measures of association for cross classifications." *Journal of the American Statistical Association* 49, 732-764. doi:[10.1080/01621459.1954](https://doi.org/10.1080/01621459.1954).

10501231.

Goodman LA, Kruskal WH. (1963) "Measures of association for cross classifications III: Approximate sampling theory." Journal of the American Statistical Association 58, 310-364. doi:10.1080/01621459.1963.10500850.

### See Also

[calcWINS\(\)](#), [calcWINS.hce\(\)](#), [calcWINS.data.frame\(\)](#).

### Examples

```
# Example 1
calcWINS(x = GROUP ~ TRTP, data = COVID19b)
# Example 2
calcWINS(x = GROUP ~ TRTP, data = COVID19, ref = "Placebo", alpha = 0.01, WOnull = 1.2)
#' Example 3
calcWINS(x = GROUP ~ TRTP, data = COVID19)$WP
calcWINS(x = GROUP ~ TRTP, data = COVID19, SE_WP_Type = "unbiased")$WP
```

---

calcWINS.hce

*Win statistics calculation for hce objects*

---

### Description

Win statistics calculation for hce objects

### Usage

```
## S3 method for class 'hce'
calcWINS(x, ...)
```

### Arguments

x                    an hce object.  
...                    additional parameters.

### Value

a list containing win statistics and their confidence intervals. It contains the following named data frames:

- summary a data frame containing number of wins, losses, and ties of the active treatment group and the overall number of comparisons.
- WP a data frame containing the win probability and its confidence interval.
- NetBenefit a data frame containing the net benefit and its confidence interval. This is just a  $2x-1$  transformation of WP and its CI.
- WO a data frame containing the win odds and its confidence interval.

- WR1 a data frame containing the win ratio and its confidence interval, using the transformed standard error of the gamma statistic.
- WR2 a data frame containing the win ratio and its confidence interval, using the standard error calculated using  $P_{ties}$ .
- gamma a data frame containing Goodman Kruskal's gamma and its confidence interval.
- SE a data frame containing standard errors used to calculate the Confidence intervals for win statistics.

## References

The theory of win statistics is covered in the following papers:

- Win proportion and win odds confidence interval calculation:

Bamber D (1975) "The area above the ordinal dominance graph and the area below the receiver operating characteristic graph." *Journal of Mathematical Psychology* 12.4: 387-415. doi:10.1016/0022-2496(75)90001-2.

DeLong ER et al. (1988) "Comparing the Areas Under Two or More Correlated Receiver Operating Characteristic Curves: A Nonparametric Approach." *Biometrics* 44.3: 837-845. doi:10.2307/2531595.

Brunner E et al. (2021) "Win odds: an adaptation of the win ratio to include ties." *Statistics in Medicine* 40.14: 3367-3384. doi:10.1002/sim.8967.

Gasparyan SB et al. (2021) "Adjusted win ratio with stratification: calculation methods and interpretation." *Statistical Methods in Medical Research* 30.2: 580-611. doi:10.1177/0962280220942558.

Gasparyan SB et al. (2021) "Power and sample size calculation for the win odds test: application to an ordinal endpoint in COVID-19 trials." *Journal of Biopharmaceutical Statistics* 31.6: 765-787. doi:10.1080/10543406.2021.1968893.

Brunner E, Konietschke F. (2025) "An unbiased rank-based estimator of the Mann-Whitney variance including the case of ties." *Statistical Papers* 66.20. doi:10.1007/s00362-024-01635-0.

- Win ratio: the first CI utilizes the standard error derived from the gamma statistic standard error as outlined by:

Gasparyan SB, Kowalewski EK, Buenconsejo J, Koch GG. (2023) "Hierarchical Composite Endpoints in COVID-19: The DARE-19 Trial." In *Case Studies in Innovative Clinical Trials*, Chapter 7, 95-148. Chapman; Hall/CRC. doi:10.1201/9781003288640-7.

- Win ratio: the second CI utilizes the standard error presented by:

Yu RX, Ganju J. (2022) "Sample size formula for a win ratio endpoint." *Statistics in Medicine* 41.6: 950-63. doi:10.1002/sim.9297.

- Goodman Kruskal's gamma and CI: matches implementation in DescTools: :GoodmanKruskalGamma() and based on:

Agresti A. (2002) Categorical Data Analysis. John Wiley & Sons, pp. 57-59. doi:10.1002/0471249688.

Brown MB, Benedetti JK. (1977) "Sampling Behavior of Tests for Correlation in Two-Way Contingency Tables." Journal of the American Statistical Association 72, 309-315. doi:10.1080/01621459.1977.10480995.

Goodman LA, Kruskal WH. (1954) "Measures of association for cross classifications." Journal of the American Statistical Association 49, 732-764. doi:10.1080/01621459.1954.10501231.

Goodman LA, Kruskal WH. (1963) "Measures of association for cross classifications III: Approximate sampling theory." Journal of the American Statistical Association 58, 310-364. doi:10.1080/01621459.1963.10500850.

### See Also

`calcWINS()`, `calcWINS.formula()`, `calcWINS.data.frame()`.

### Examples

```
# Example 1
COVID19HCE <- hce(GROUP = COVID19$GROUP, TRTP = COVID19$TRTP)
calcWINS(COVID19HCE)
# Example 2
COVID19bHCE <- hce(GROUP = COVID19b$GROUP, TRTP = COVID19b$TRTP)
calcWINS(COVID19bHCE, ref = "Placebo", WOnull = 1.1, alpha = 0.01)
# Example 3
calcWINS(COVID19HCE, SE_WP_Type = "unbiased")$WP
calcWINS(COVID19HCE, SE_WP_Type = "biased")$WP
```

---

calcWO

*A generic function for calculating win odds*

---

### Description

A generic function for calculating win odds

### Usage

```
calcWO(x, ...)
```

### Arguments

x                    an object used to select a method.  
...                   further arguments passed to or from other methods.

**Value**

a data frame containing calculated values.

**See Also**

`calcWO.hce()`, `calcWO.formula()`, `calcWO.data.frame()` methods.

---

|                   |  |
|-------------------|--|
| calcWO.data.frame | <i>Win odds calculation using a data frame</i> |
|-------------------|--|

---

**Description**

Win odds calculation using a data frame

**Usage**

```
## S3 method for class 'data.frame'
calcWO(x, AVAL, TRTP, ref, alpha = 0.05, WOnull = 1, ...)
```

**Arguments**

|        |  |
|--------|--|
| x      | a data frame containing subject-level data.        |
| AVAL   | variable in the data with ordinal analysis values. |
| TRTP   | the treatment variable in the data.                |
| ref    | the reference treatment group.                     |
| alpha  | significance level. The default is 0.05.           |
| WOnull | the null hypothesis. The default is 1.             |
| ...    | additional parameters.                             |

**Value**

a data frame containing the win odds and its confidence interval. It contains the following columns:

- WO calculated win odds.
- LCL lower confidence limit.
- UCL upper confidence limit.
- SE standard error of the win odds.
- WOnull win odds of the null hypothesis (specified in the WOnull argument).
- alpha two-sided significance level for calculating the confidence interval (specified in the alpha argument).
- Pvalue p-value associated with testing the null hypothesis.
- WP calculated win probability.
- LCL\_WP lower confidence limit for WP.



- UCL\_WP upper confidence limit for WP.
- SE\_WP standard error of the win probability.
- SD\_WP standard deviation of the win probability, calculated as SE\_WP multiplied by sqrt(N).
- N total number of patients in the analysis.

## References

Gasparyan SB et al. "Adjusted win ratio with stratification: calculation methods and interpretation." *Statistical Methods in Medical Research* 30.2 (2021): 580-611. doi:10.1177/0962280220942558

## See Also

[calcWO\(\)](#), [calcWO.hce\(\)](#), [calcWO.formula\(\)](#).

## Examples

```
data(HCE4)
calcWO(x = HCE4, AVAL = "AVAL", TRTP = "TRTP", ref = "P")
```

---

|                |  |
|----------------|--|
| calcWO.formula | <i>Win odds calculation using formula syntax</i> |
|----------------|--|

---

## Description

Win odds calculation using formula syntax

## Usage

```
## S3 method for class 'formula'
calcWO(x, data, ...)
```

## Arguments

|      |                             |
|------|-----------------------------|
| x    | an object of class formula. |
| data | a data frame.               |
| ...  | additional parameters.      |

## Value

a data frame containing the win odds and its confidence interval. It contains the following columns:

- WO calculated win odds.
- LCL lower confidence limit.
- UCL upper confidence limit.
- SE standard error of the win odds.
- WOnull win odds of the null hypothesis (specified in the WOnull argument).

- alpha two-sided significance level for calculating the confidence interval (specified in the alpha argument).
- Pvalue p-value associated with testing the null hypothesis.
- WP calculated win probability.
- LCL\_WP lower confidence limit for WP.
- UCL\_WP upper confidence limit for WP.
- SE\_WP standard error of the win probability.
- SD\_WP standard deviation of the win probability, calculated as SE\_WP multiplied by sqrt(N).
- N total number of patients in the analysis.
- formula returning the specified formula in the x argument.
- ref showing how the reference group was selected. Can be modifying by specifying the ref argument.

## References

Gasparyan SB et al. "Adjusted win ratio with stratification: calculation methods and interpretation." *Statistical Methods in Medical Research* 30.2 (2021): 580-611. doi:[10.1177/0962280220942558](https://doi.org/10.1177/0962280220942558)

## See Also

[calcWO\(\)](#), [calcWO.hce\(\)](#), [calcWO.data.frame\(\)](#).

## Examples

```
#Example 1
data(HCE1)
calcWO(AVAL ~ TRTP, data = HCE1)

#Example 2
calcWO(GROUP ~ TRTP, data = COVID19, ref = "Placebo", alpha = 0.01)
```

---

calcWO.hce

*Win odds calculation for hce objects*

---

## Description

Win odds calculation for hce objects

## Usage

```
## S3 method for class 'hce'
calcWO(x, ...)
```

**Arguments**

x                    an hce object.  
...                    additional parameters.

**Value**

a data frame containing the win odds and its confidence interval. It contains the following columns:

- WO calculated win odds.
- LCL lower confidence limit.
- UCL upper confidence limit.
- SE standard error of the win odds.
- WOnull win odds of the null hypothesis (specified in the WOnull argument).
- alpha two-sided significance level for calculating the confidence interval (specified in the alpha argument).
- Pvalue p-value associated with testing the null hypothesis.
- WP calculated win probability.
- LCL\_WP lower confidence limit for WP.
- UCL\_WP upper confidence limit for WP.
- SE\_WP standard error of the win probability.
- SD\_WP standard deviation of the win probability, calculated as SE\_WP multiplied by sqrt(N).
- N total number of patients in the analysis.

**References**

Gasparyan SB et al. "Adjusted win ratio with stratification: calculation methods and interpretation." *Statistical Methods in Medical Research* 30.2 (2021): 580-611. doi:[10.1177/0962280220942558](https://doi.org/10.1177/0962280220942558)

**See Also**

[calcWO\(\)](#), [calcWO.formula\(\)](#), [calcWO.data.frame\(\)](#).

**Examples**

```
Rates_A <- c(1, 1.5)
Rates_P <- c(2, 2)
dat <- simHCE(n = 500, TTE_A = Rates_A, TTE_P = Rates_P, CM_A = 1.25, CM_P = 1)
calcWO(dat)
calcWO(dat, ref = "A", WOnull = 1, alpha = 0.01)
```

---

 COVID19

*COVID-19 ordinal scale dataset (full report).*


---

**Description**

A dataset with COVID-19 ordinal scale outcomes for 1062 patients.

**Usage**

COVID19

**Format**

a data frame with 1062 rows and 2 variables:

**GROUP** type of the event, ordinal outcomes 1-8, where a higher value means a better outcome

**TRTP** treatment values, A Active or P Placebo, character

**Source**

Beigel JH et al. "Remdesivir for the treatment of Covid-19-final report." New England Journal of Medicine 383.19 (2020): 1813-1836. [doi:10.1056/NEJMoa2007764](https://doi.org/10.1056/NEJMoa2007764).

**Examples**

```
#Frequencies
table(COVID19)
mosaicplot(table(COVID19), col = c(1, 8, 6, 2, 4, 5, 3, 7),
xlab = "Treatment", ylab = "Ordinal Scale", main = "COVID-19 ordinal scale")
# Convert to an hce object
COVID19HCE <- hce(GROUP = COVID19$GROUP, TRTP = COVID19$TRTP)
# Summary wins, losses, and ties with win odds
summaryWO(COVID19HCE, ref = "Placebo")
```

---

 COVID19b

*COVID-19 ordinal scale dataset (preliminary report).*


---

**Description**

A dataset with COVID-19 ordinal scale outcomes for 844 patients.

**Usage**

COVID19b

**Format**

a data frame with 844 rows and 2 variables:

**GROUP** type of the event, ordinal outcomes 1-8, where a higher value means a better outcome

**TRTP** treatment values, Active or Placebo, character

**Source**

Beigel JH et al. "Remdesivir for the treatment of Covid-19-final report." New England Journal of Medicine 383.19 (2020): 1813-1836. doi:[10.1056/NEJMoa2007764](https://doi.org/10.1056/NEJMoa2007764).

**Examples**

```
#Frequencies
table(COVID19b)
mosaicplot(table(COVID19b), col = c(1, 8, 6, 2, 4, 5, 3, 7),
xlab = "Treatment", ylab = "Ordinal Scale", main = "COVID-19 ordinal scale")
# Calculate win statistics
calcWINS(x = COVID19b, AVAL = "GROUP", TRTP = "TRTP", ref = "Placebo")
```

---

COVID19plus

*COVID-19 ordinal scale dataset for a combination therapy.*

---

**Description**

A dataset with COVID-19 ordinal scale outcomes for 1033 patients.

**Usage**

COVID19plus

**Format**

a data frame with 1033 rows and 4 variables:

**ID** patient identifiers, numeric

**TRTP** treatment values, A Active or P Placebo, character

**GROUP** type of the event, ordinal outcomes 1-8, where a higher value means a better outcome

**BASE** baseline ordinal values

**Source**

Kalil AC et al. "Baricitinib plus Remdesivir for Hospitalized Adults with Covid-19." New England Journal of Medicine 384.9 (2021): 795-807. doi:[10.1056/NEJMoa2031994](https://doi.org/10.1056/NEJMoa2031994).

**Examples**

```
COVID19HCE <- hce(GROUP = COVID19plus$GROUP, TRTP = COVID19plus$TRTP)
# Summary wins, losses, and ties with win odds
summaryWO(COVID19HCE, ref = "P")
```

---

hce

*Helper function for hce objects*

---

## Description

Helper function for hce objects

## Usage

```
hce(  
  GROUP = character(),  
  TRTP = character(),  
  AVAL0 = 0,  
  ORD = sort(unique(GROUP))  
)
```

## Arguments

|       |  |
|-------|--|
| GROUP | a character vector of the same length as AVAL containing events.   |
| TRTP  | a character vector of the same length as AVAL containing assigned treatment groups.                                |
| AVAL0 | a numeric vector of analysis values within each category. The default is 0.  |
| ORD   | a character vector containing ordered unique values of the GROUP variable for determining the hierarchy of events. |

## Value

an object of class hce. Its is a subject-level data frame (each row corresponds to one subject), containing the following columns:

- SUBJID subject ID.
- GROUP a character vector specifying the type of the outcome the patient experienced - either a TTE (time-to-event) or C (continuous).
- GROUPLN a numeric vector version of the GROUP column.
- AVAL0 original analysis values - time of the time-to-event outcomes or the continuous outcome.
- AVAL derived analysis value  $AVAL = AVAL0 + GROUPLN$ .
- TRTP assigned treatment groups.

## See Also

[as\\_hce\(\)](#) for coercing to hce objects.

**Examples**

```
# Example 1
set.seed(2022)
d <- hce(GROUP = sample(x = c("A", "B", "C"), size = 10, replace = TRUE),
TRTP = rep(c("Active", "Control"), each = 5),
AVAL0 = c(rnorm(5, mean = 1), rnorm(5)), ORD = c("A", "B", "C"))
calcW0(d, ref = "Control")
```

---

|      |  |
|------|--|
| HCE1 | HCE1, HCE2, HCE3, HCE4 <i>datasets for 1000 patients with different treatment effects.</i> |
|------|--|

---

**Description**

A simulated dataset containing the ordinal values and other attributes for 1000 patients. HCE1

**Usage**

HCE1

**Format**

a data frame with 1000 rows and 6 variables:

**ID** subject ID, numbers from 1 to 1000

**TRTP** treatment values, A Active or P Placebo, character

**GROUP** type of the event, either Time-To-Event (TTE) or Continuous (C), character

**GROUPN** type of the event, for the ordering of outcomes in the GROUP variable, numeric

**AVALT** the timing of the time-to-event outcomes, numeric

**AVAL0** original values for each type of the event, time for TTE outcomes, numeric values for Continuous outcomes, numeric

**AVAL**  $AVAL = AVAL0 + GROUPN$ , ordinal analysis values for the HCE analysis. For the continuous outcome the values of AVAL0 are shifted to start always from 0. Numeric, but caution NOT to apply numeric operations; will give meaningless results

**PADY** primary analysis day, the length of fixed follow-up in days, numeric

---

|      |  |
|------|--|
| HCE2 | HCE1, HCE2, HCE3, HCE4 <i>datasets for 1000 patients with different treatment effects.</i> |
|------|--|

---

### Description

A simulated dataset containing the ordinal values and other attributes for 1000 patients. HCE2

### Usage

HCE2

### Format

a data frame with 1000 rows and 6 variables:

**ID** subject ID, numbers from 1 to 1000

**TRTP** treatment values, A Active or P Placebo, character

**GROUP** type of the event, either Time-To-Event (TTE) or Continuous (C), character

**GROUPN** type of the event, for the ordering of outcomes in the GROUP variable, numeric

**AVALT** the timing of the time-to-event outcomes, numeric

**AVAL0** original values for each type of the event, time for TTE outcomes, numeric values for Continuous outcomes, numeric

**AVAL**  $AVAL = AVAL0 + GROUPN$ , ordinal analysis values for the HCE analysis. For the continuous outcome the values of AVAL0 are shifted to start always from 0. Numeric, but caution NOT to apply numeric operations; will give meaningless results

**PADY** primary analysis day, the length of fixed follow-up in days, numeric

---

|      |  |
|------|--|
| HCE3 | HCE1, HCE2, HCE3, HCE4 <i>datasets for 1000 patients with different treatment effects.</i> |
|------|--|

---

### Description

A simulated dataset containing the ordinal values and other attributes for 1000 patients. HCE3

### Usage

HCE3



**Format**

a data frame with 1000 rows and 6 variables:

**ID** subject ID, numbers from 1 to 1000

**TRTP** treatment values, A Active or P Placebo, character

**GROUP** type of the event, either Time-To-Event (TTE) or Continuous (C), character

**GROUPN** type of the event, for the ordering of outcomes in the GROUP variable, numeric

**AVALT** the timing of the time-to-event outcomes, numeric

**AVAL0** original values for each type of the event, time for TTE outcomes, numeric values for Continuous outcomes, numeric

**AVAL**  $AVAL = AVAL0 + GROUPN$ , ordinal analysis values for the HCE analysis. For the continuous outcome the values of AVAL0 are shifted to start always from 0. Numeric, but caution NOT to apply numeric operations; will give meaningless results

**PADY** primary analysis day, the length of fixed follow-up in days, numeric

---

|      |  |
|------|--|
| HCE4 | HCE1, HCE2, HCE3, HCE4 <i>datasets for 1000 patients with different treatment effects.</i> |
|------|--|

---

**Description**

A simulated dataset containing the ordinal values and other attributes for 1000 patients. HCE4

**Usage**

HCE4

**Format**

a data frame with 1000 rows and 6 variables:

**ID** subject ID, numbers from 1 to 1000

**TRTP** treatment values, A Active or P Placebo, character

**GROUP** type of the event, either Time-To-Event (TTE) or Continuous (C), character

**GROUPN** type of the event, for the ordering of outcomes in the GROUP variable, numeric

**AVALT** the timing of the time-to-event outcomes, numeric

**AVAL0** original values for each type of the event, time for TTE outcomes, numeric values for Continuous outcomes, numeric

**AVAL**  $AVAL = AVAL0 + GROUPN$ , ordinal analysis values for the HCE analysis. For the continuous outcome the values of AVAL0 are shifted to start always from 0. Numeric, but caution NOT to apply numeric operations; will give meaningless results

**PADY** primary analysis day, the length of fixed follow-up in days, numeric

IWP

*Calculates patient-level individual win proportions***Description**

Calculates patient-level individual win proportions

**Usage**

```
IWP(data, AVAL, TRTP, ref)
```

**Arguments**

|      |  |
|------|--|
| data | a data frame containing subject-level data.        |
| AVAL | variable in the data with ordinal analysis values. |
| TRTP | the treatment variable in the data.                |
| ref  | the reference treatment group.                     |

**Value**

the input data frame with a new column of individual win proportions named using the input AVAL value with `_`.

**References**

Gasparyan SB et al. "Adjusted win ratio with stratification: calculation methods and interpretation." *Statistical Methods in Medical Research* 30.2 (2021): 580-611. doi:[10.1177/0962280220942558](https://doi.org/10.1177/0962280220942558)

**See Also**

[calcWO\(\)](#), [calcWO.hce\(\)](#), [calcWO.formula\(\)](#).

**Examples**

```
KHCE1 <- IWP(data = KHCE, AVAL = "EGFRBL", TRTP = "TRTPN", ref = 2)
WP <- tapply(KHCE1$EGFRBL_, KHCE1$TRTPN, mean)
VAR <- tapply(KHCE1$EGFRBL_, KHCE1$TRTPN, function(x) (length(x)-1)*var(x)/length(x))
N <- tapply(KHCE1$EGFRBL_, KHCE1$TRTPN, length)
SE <- sqrt(sum(VAR/N))
c(WP = WP[[1]], SE = SE)
calcWO(EGFRBL ~ TRTP, data = KHCE)[c("WP", "SE_WP")]
```

---

KHCE *Kidney Hierarchical Composite Endpoint dataset.*

---

### Description

A dataset with kidney ordinal scale outcomes of 1500 patients in the ADSL dataset.

### Usage

KHCE

### Format

a data frame with 1500 rows and 11 variables:

**ID** patient identifiers, numeric

**TRTPN** treatment values, 1 Active or 2 Placebo, numeric

**AVAL0** original values for each type of the event, time for TTE outcomes 1-6, numeric values for Continuous outcome 7, numeric

**AVAL**  $AVAL = AVAL0 + GROUPN$ , ordinal analysis values for the HCE analysis, numeric, but caution NOT to apply numeric operations; will give meaningless results

**GROUP** name of the event, character

**GROUPN** ordinal outcomes corresponding to PARAMN values, numeric

**PARAMCD** coded name of the event, character

**PARAMN** severity of the event, outcomes 1-7, where a higher value means a better outcome, character

**STRATAN** strata 1-4, higher value means more severe kidney disease, numeric

**EGFRBL** Baseline GFR values of patients, numeric

**TRTP** treatment values, A Active or P Placebo, character

**PADY** primary analysis day (in years), length of the fixed follow-up, numeric

### Source

Heerspink HL et al "Development and validation of a new hierarchical composite endpoint for clinical trials of kidney disease progression." Journal of the American Society of Nephrology (2023): [doi:10.1681/ASN.000000000000243](https://doi.org/10.1681/ASN.000000000000243).

### Examples

```
# Adjusted win odds
res <- regWO(x = KHCE, AVAL = "AVAL", TRTP = "TRTP", COVAR = "STRATAN", ref = "P")
res
```

---

|       |  |
|-------|--|
| minWO | <i>Minimum detectable or WO for alternative hypothesis for given power (no ties)</i> |
|-------|--|

---

### Description

Minimum detectable or WO for alternative hypothesis for given power (no ties)

### Usage

```
minWO(N, power = 0.5, SD = NULL, k = 0.5, alpha = 0.05, WOnull = 1, digits = 2)
```

### Arguments

|        |   |
|--------|---|
| N      | a numeric vector of sample size values (two arms combined).   |
| power  | the given power. The default is 0.5 corresponding to the minimum detectable win odds. A numeric vector of length 1.           |
| SD     | assumed standard deviation of the win proportion. By default uses the conservative SD. A numeric vector of length 1.          |
| k      | proportion of active group in the overall sample size. Default is 0.5 (balanced randomization). A numeric vector of length 1. |
| alpha  | the significance level for the 2-sided test. Default is 0.05. A numeric vector of length 1.                                   |
| WOnull | the win odds value of the null hypothesis (default is 1). A numeric vector of length 1.                                       |
| digits | precision to use for reporting calculated win odds.   |

### Value

a data frame containing the calculated WO with input values.

### References

Gasparyan SB et al. (2021) "Power and sample size calculation for the win odds test: application to an ordinal endpoint in COVID-19 trials." *Journal of Biopharmaceutical Statistics* 31.6: 765-787. [doi:10.1080/10543406.2021.1968893](https://doi.org/10.1080/10543406.2021.1968893)

### See Also

[powerWO\(\)](#), [sizeWO\(\)](#) for WO power and sample size calculation.

**Examples**

```

minWO(N = 100, digits = 5)
minWO(N = 1200, power = 0.9, k = 0.75)
# Compare the minimum detectable win odds from shifted alternatives to max and ordered alternatives
WO <- minWO(N = 1200, k = 0.5, power = 0.67, digits = 7)$WO
powerWO(N = 1200, WO = WO, k = 0.5, alternative = "shift")
powerWO(N = 1200, WO = WO, k = 0.5, alternative = "ordered")
powerWO(N = 1200, WO = WO, k = 0.5, alternative = "max")

```

plot.hce

*A plot method for hce objects***Description**

Ordinal dominance graph for hce objects

**Usage**

```

## S3 method for class 'hce'
plot(x, fill = FALSE, ...)

```

**Arguments**

|      |  |
|------|--|
| x    | an object of class hce created by <code>as_hce()</code> .                |
| fill | logical; if TRUE fill the area above the graph.                          |
| ...  | additional arguments to be passed to <code>base::plot()</code> function. |

**Value**

no return value, called for plotting.

**References**

Bamber D. "The area above the ordinal dominance graph and the area below the receiver operating characteristic graph." *Journal of Mathematical Psychology* 12.4 (1975): 387-415. doi:[10.1016/0022-2496\(75\)90001-2](https://doi.org/10.1016/0022-2496(75)90001-2)

**Examples**

```

d <- as_hce(KHCE)
d$TRTP <- factor(d$TRTP, levels = c("P", "A"))
res <- calcWO(AVAL ~ TRTP, data = d)
# Ordinal Dominance Graph
plot(d, col = 3, type = 'l')
grid()
# Area above the Ordinal Dominance Graph
plot(d, fill = TRUE, col = "#865A4F", type = 'l',
      lwd = 2, xlab = "Control", ylab = "Active")
legend("bottomright", legend = paste0("WP = ", round(res$WP, 5)))
abline(a = 0, b = 1, lwd = 2, lty = 2, col = "#999999")

```

---

plot.hce\_results      *A print method for hce\_results objects*

---

**Description**

A print method for hce\_results objects

**Usage**

```
## S3 method for class 'hce_results'
plot(x, ...)
```

**Arguments**

x                    an object of class hce\_results.  
 ...                  additional arguments to be passed to `base::plot()` function.

**Value**

no return value, called for plotting.

**Examples**

```
WO <- minWO(N = 100:1000)
plot(WO)
POW <- powerWO(N = 100:1000, WO = 1.2)
plot(POW, ylim = c(0, 1))
```

---

powerWO                    *Power calculation for the win odds test (no ties)*

---

**Description**

Power calculation for the win odds test (no ties)

**Usage**

```
powerWO(
  N,
  WO,
  SD = NULL,
  k = 0.5,
  alpha = 0.05,
  WOnull = 1,
  alternative = c("shift", "max", "ordered")
)
```

**Arguments**

|             |   |
|-------------|---|
| N           | a numeric vector of sample size values.   |
| WO          | the given win odds for the alternative hypothesis. A numeric vector of length 1.  |
| SD          | assumed standard deviation of the win proportion. By default uses the conservative SD. A numeric vector of length 1.  |
| k           | proportion of active group in the overall sample size. Default is 0.5 (balanced randomization). A numeric vector of length 1.   |
| alpha       | the significance level for the 2-sided test. Default is 0.05. A numeric vector of length 1.   |
| WOnull      | the win odds value of the null hypothesis (default is 1). A numeric vector of length 1.   |
| alternative | a character string specifying the class of the alternative hypothesis, must be one of "shift" (default), "max" or "ordered". You can specify just the initial letter. |

**Details**

alternative = "max" refers to the maximum variance of the win proportion across all possible alternatives. The maximum variance equals  $WP*(1 - WP)/k$  where the win probability is calculated as  $WP = WO/(WO + 1)$ . alternative = "shift" specifies the variance across alternatives from a shifted family of distributions (Wilcoxon test). The variance formula, as suggested by Noether, is calculated based on the null hypothesis as follows  $1/(12*k*(1 - k))$ . alternative = "ordered" specifies the variance across alternatives from stochastically ordered distributions which include shifted distributions.

**Value**

a data frame containing the calculated power with input values.

**References**

- All formulas were presented in

Bamber D (1975) "The area above the ordinal dominance graph and the area below the receiver operating characteristic graph." *Journal of Mathematical Psychology* 12.4: 387-415. doi:10.1016/0022-2496(75)90001-2.

- Noether's formula for shifted alternatives

Noether GE (1987) "Sample size determination for some common nonparametric tests." *Journal of the American Statistical Association* 82.398: 645-7. doi:10.1080/01621459.1987.10478478.

- For shift alternatives see also

Gasparyan SB et al. (2021) "Power and sample size calculation for the win odds test: application to an ordinal endpoint in COVID-19 trials." *Journal of Biopharmaceutical Statistics* 31.6: 765-787. doi:10.1080/10543406.2021.1968893.

**See Also**

`sizeWO()`, `minWO()` for WO sample size or minimum detectable WO calculation.

**Examples**

```
# Example 1- Use the default standard deviation
powerWO(N = 1000, WO = 1.2)
powerWO(N = seq(500, 1500, 100), WO = 1.2)
# Example 2 - Use data-driven win odds and standard deviation from the COVID19 dataset
res <- calcWO(x = COVID19, AVAL = "GROUP", TRTP = "TRTP", ref = "Placebo")
print(res)
powerWO(N = 500, WO = res$WO, SD = res$SD_WP)
powerWO(N = 500, WO = res$WO) # power with the default standard deviation for the win proportion.
# Example 3 - Non-balanced 3:1 randomization
powerWO(N = 1000, WO = 1.2, k = 0.75)
# Example 4 - Comparison of different alternatives
powerWO(N = 1000, WO = 1.2, alternative = "m")
powerWO(N = 1000, WO = 1.2, alternative = "s")
powerWO(N = 1000, WO = 1.2, alternative = "o")
```

---

print.hce\_results      *A print method for hce\_results objects*

---

**Description**

A print method for hce\_results objects

**Usage**

```
## S3 method for class 'hce_results'
print(x, ...)
```

**Arguments**

`x`                    an object of class hce\_results.  
`...`                  additional arguments to be passed to `base::print()` function.

**Value**

no return value, called for printing.

**Examples**

```
print(powerWO(N = 1000, WO = 1.2))
```



---

|          |  |
|----------|--|
| propWINS | <i>Proportion of wins/losses/ties given the win odds and the win ratio</i> |
|----------|--|

---

### Description

Proportion of wins/losses/ties given the win odds and the win ratio

### Usage

```
propWINS(WO, WR, Overall = 1, alpha = NULL, N = NULL)
```

### Arguments

|         |   |
|---------|---|
| WO      | win odds.   |
| WR      | win ratio.  |
| Overall | number of comparisons, the sample size of the active treatment multiplied by the sample size of the placebo. The default is 1, hence gives the proportion.  |
| alpha   | significance level for the win ratio confidence interval. The default is NULL hence the confidence interval is not produced.  |
| N       | the combined sample size of two treatment groups. The default is NULL. If alpha is specified then either N should be specified or Overall > 1. For given Overall, the pooled sample size is calculated as $N = 2 * \text{sqrt}(\text{Overall})$ . |

### Details

- **Win ratio** defined as  $WR = \frac{W}{L}$ .
- **Win odds** defined as  $WO = \frac{W+0.5T}{L+0.5T} = \frac{WP}{1-WP}$ .
- **Net Benefit** defined as  $NB = \frac{W-L}{O}$ .

Given the overall number of comparisons  $O$ , the win proportion  $WP$  and the win ratio  $WR$ , it is possible to find the total number of wins and losses. Indeed, first the win odds can be found  $WO = \frac{WP}{1-WP}$  and

$$L = O * \frac{2WP - 1}{WR - 1},$$

$$W = WR * O * \frac{2WP - 1}{WR - 1},$$

$$T = O - W - L.$$

### Value

a data frame with a number (or proportion if Overall = 1) of wins/losses/ties. If alpha is specified returns also WR confidence interval.

## References

- For the relationship between win odds and win ratio see  
 Gasparian SB et al. "Hierarchical Composite Endpoints in COVID-19: The DARE-19 Trial". Case Studies in Innovative Clinical Trials, Chapter 7 (2023): 95-148. Chapman and Hall/CRC. doi:10.1201/9781003288640-7.
- The win ratio CI uses the standard error presented in  
 Yu RX, Ganju J. (2022) "Sample size formula for a win ratio endpoint." Statistics in Medicine 41.6: 950-63. doi:10.1002/sim.9297.

## Examples

```
# Example 1
propWINS(WR = 2, WO = 1.5)
# Example 2 - Back-calculation
COVID19HCE <- hce(GROUP = COVID19$GROUP, TRTP = COVID19$TRTP)
res <- calcWINS(COVID19HCE)
WR <- res$WR1$WR
WO <- res$WO$WO
Overall <- res$summary$TOTAL
propWINS(WR = WR, WO = WO, Overall = Overall)
## Verify
res$summary
# Example 3 - Confidence interval
propWINS(WR = 1.4, WO = 1.3, alpha = 0.05, Overall = 2500)
propWINS(WR = 2, WO = 1.5, alpha = 0.01, N = 500)
```

---

regWO

*A generic function for win odds regression*

---

## Description

A generic function for win odds regression

## Usage

```
regWO(x, ...)
```

## Arguments

x                    an object used to select a method.  
 ...                  further arguments passed to or from other methods.

## Value

a data frame containing calculated values.

**See Also**

[regWO.data.frame\(\)](#), [regWO.formula\(\)](#) methods.

---

|                  |   |
|------------------|---|
| regWO.data.frame | <i>Win Odds Regression Using a Data Frame</i> |
|------------------|---|

---

**Description**

This function performs regression analysis for the win odds using a single numeric covariate.

**Usage**

```
## S3 method for class 'data.frame'
regWO(x, AVAL, TRTP, COVAR, ref, alpha = 0.05, WOnull = 1, ...)
```

**Arguments**

|        |   |
|--------|---|
| x      | a data frame containing subject-level data.               |
| AVAL   | a variable in the data with ordinal analysis values.      |
| TRTP   | the treatment variable in the data.                       |
| COVAR  | a numeric covariate.                                      |
| ref    | the reference treatment group.                            |
| alpha  | the significance level, with a default value of 0.05.     |
| WOnull | the null hypothesis value for win odds. The default is 1. |
| ...    | additional parameters.                                    |

**Value**

a data frame containing the calculated win odds and its confidence interval, including:

- WO\_beta adjusted win odds.
- LCL lower confidence limit for adjusted WO.
- UCL upper confidence limit for adjusted WO.
- SE standard error of the adjusted win odds.
- WOnull win odds of the null hypothesis (specified in the WOnull argument).
- alpha two-sided significance level for calculating the confidence interval (specified in the alpha argument).
- Pvalue p-value associated with testing the null hypothesis.
- N total number of patients in the analysis.
- beta adjusted win probability.
- LCL\_beta lower confidence limit for adjusted win probability.
- UCL\_beta upper confidence limit for adjusted win probability.

- SE\_beta standard error for the adjusted win probability.
- SD\_beta standard deviation for the adjusted win probability.
- WP (non-adjusted) win probability.
- SE\_WP standard error of the non-adjusted win probability.
- SD\_WP standard deviation of the non-adjusted win probability.
- WO non-adjusted win odds.
- COVAR\_MEAN\_DIFF mean difference between two treatment groups of the numeric covariate.
- COVAR\_VAR sum of variances of two treatment groups of the numeric covariate.
- COVAR\_COV covariance between the response and the numeric covariate.

## References

Gasparyan SB et al. (2021) "Adjusted win ratio with stratification: calculation methods and interpretation." *Statistical Methods in Medical Research* 30.2: 580-611. doi:[10.1177/0962280220942558](https://doi.org/10.1177/0962280220942558).

## See Also

[regWO\(\)](#), [regWO.formula\(\)](#).

## Examples

```
# A baseline covariate that is highly correlated with the outcome
set.seed(2023)
dat <- COVID19
n <- nrow(dat)
dat$Severity <- ifelse(dat$GROUP > 4, rnorm(n, 0), rnorm(n, 100))
tapply(dat$Severity, dat$TRTP, mean)
regWO(x = dat, AVAL = "GROUP", TRTP = "TRTP", COVAR = "Severity", ref = "Placebo")
# Without adjustment
calcWO(x = dat, AVAL = "GROUP", TRTP = "TRTP", ref = "Placebo")
```

---

regWO.formula

*Win Odds Regression Using a Formula Syntax*

---

## Description

This function performs regression analysis for the win odds using a single numeric covariate.

## Usage

```
## S3 method for class 'formula'
regWO(x, data, ...)
```

**Arguments**

|      |                             |
|------|-----------------------------|
| x    | an object of class formula. |
| data | a data frame.               |
| ...  | additional parameters.      |

**Value**

a data frame containing the calculated win odds and its confidence interval, including:

- WO\_beta adjusted win odds.
- LCL lower confidence limit for adjusted WO.
- UCL upper confidence limit for adjusted WO.
- SE standard error of the adjusted win odds.
- WOnull win odds of the null hypothesis (specified in the WOnull argument).
- alpha two-sided significance level for calculating the confidence interval (specified in the alpha argument).
- Pvalue p-value associated with testing the null hypothesis.
- N total number of patients in the analysis.
- beta adjusted win probability.
- LCL\_beta lower confidence limit for adjusted win probability.
- UCL\_beta upper confidence limit for adjusted win probability.
- SE\_beta standard error for the adjusted win probability.
- SD\_beta standard deviation for the adjusted win probability.
- WP (non-adjusted) win probability.
- SE\_WP standard error of the non-adjusted win probability.
- SD\_WP standard deviation of the non-adjusted win probability.
- WO non-adjusted win odds.
- COVAR\_MEAN\_DIFF mean difference between two treatment groups of the numeric covariate.
- COVAR\_VAR sum of variances of two treatment groups of the numeric covariate.
- COVAR\_COV covariance between the response and the numeric covariate.
- formula returning the specified formula in the x argument.
- ref showing how the reference group was selected. Can be modifying by specifying the ref argument.

**References**

Gasparian SB et al. (2021) "Adjusted win ratio with stratification: calculation methods and interpretation." *Statistical Methods in Medical Research* 30.2: 580-611. doi:[10.1177/0962280220942558](https://doi.org/10.1177/0962280220942558).

**See Also**

[regWO\(\)](#), [regWO.data.frame\(\)](#).

**Examples**

```
regWO(AVAL ~ TRTP, data = KHCE)
regWO(AVAL ~ TRTP + EGFRBL, data = KHCE)
```

---

|        |  |
|--------|--|
| simHCE | <i>Simulate hce object with given event rates of time-to-event outcomes (Weibull), mean and SD of the continuous outcome (normal or log-normal) by treatment group</i> |
|--------|--|

---

**Description**

Simulate hce object with given event rates of time-to-event outcomes (Weibull), mean and SD of the continuous outcome (normal or log-normal) by treatment group

**Usage**

```
simHCE(
  n,
  n0 = n,
  TTE_A,
  TTE_P,
  CM_A,
  CM_P,
  CSD_A = 1,
  CSD_P = CSD_A,
  fixedfy = 1,
  yeardays = 360,
  pat = 100,
  shape = 1,
  theta = 1,
  logC = FALSE,
  seed = NULL,
  dec = 2,
  all_data = FALSE
)
```

**Arguments**

|       |   |
|-------|---|
| n     | sample size in the active treatment group.  |
| n0    | sample size in the placebo group.   |
| TTE_A | event rates per year in the active group for the time-to-event outcomes.  |
| TTE_P | event rates per year in the placebo group for the time-to-event outcomes. Should have the same length as TTE_A. |
| CM_A  | mean value for the continuous outcome of the active group.  |
| CM_P  | mean value for the continuous outcome of the placebo group.   |

|          |  |
|----------|--|
| CSD_A    | standard deviation for the continuous outcome of the active group.   |
| CSD_P    | standard deviation for the continuous outcome of the placebo group.  |
| fixedfy  | length of follow-up in years.  |
| yeardays | number of days in a year.  |
| pat      | scale of provided event rates (per pat-years).   |
| shape    | shape of the Weibull distribution for time-to-event outcomes. Default is exponential distribution with shape = 1.  |
| theta    | Gumbel dependence coefficient of the Weibull distributions for time-to-event outcomes. Default is theta = 1 which assumes independence of time-to-event outcomes. Must be above or equal to 1.   |
| logC     | logical, whether to use log-normal distribution for the continuous outcome.  |
| seed     | for generating random numbers.   |
| dec      | decimal places for the continuous outcome used for rounding. The default is dec = 2.   |
| all_data | logical, whether to return source datasets ADET (an event-time dataset for all time-to-event outcomes per patient) and BDS (a basic data structure for the continuous outcome for all patients). |

### Value

an object of class `hce` containing the following columns:

- ID subject identifier.
- TRTP planned treatment group - "A" for active, "P" for Placebo.
- GROUP type of the outcome, either "TTE" for time-to-event outcomes or "C" for continuous. Only one continuous outcome is possible, but no restriction on the number of "TTE" outcomes.
- GROUPN order of outcomes in GROUP, with a higher value signifying a better outcome.
- AVALT the timing of the time-to-event outcomes.
- AVAL0 numeric values of the continuous outcome and the timing of "TTE" outcomes.
- AVAL analysis values derived as  $AVAL0 + GROUPN$ . For the continuous outcome the values of AVAL0 are shifted to start always from 0.
- seed the seed of the random sample. If not specified in seed argument will be selected based on system time.
- PADY primary analysis day, the length of fixed follow-up in days calculated as `yeardays` multiplied by `fixedfy`.

If `all_data = TRUE`, the function returns a list containing the `hce` dataset, along with its source datasets: ADET (an event-time dataset for all time-to-event outcomes per patient) and BDS (a basic data structure for the continuous outcome for all patients).

### See Also

[hce\(\)](#), [as\\_hce\(\)](#) for the helper a coerce function to `hce` objects.

## Examples

```
# Example 1
Rates_A <- c(1.72, 1.74, 0.58, 1.5, 1)
Rates_P <- c(2.47, 2.24, 2.9, 4, 6)
dat <- simHCE(n = 2500, TTE_A = Rates_A, TTE_P = Rates_P,
             CM_A = -3, CM_P = -6, CSD_A = 16, CSD_P = 15, fixedfy = 3)
head(dat)

# Example 2
Rates_A <- 10
Rates_P <- 15
dat <- simHCE(n = 1000, n0 = 500, TTE_A = Rates_A, TTE_P = Rates_P,
             CM_A = 0.1, CM_P = 0, seed = 5, shape = 0.2, logC = TRUE, dec = 0)
summaryWO(dat)

# Example 3: Comparison of dependent and independent outcomes
Rates_A <- c(10, 20)
Rates_P <- c(20, 20)
dat1 <- simHCE(n = 2500, TTE_A = Rates_A, TTE_P = Rates_P,
             CM_A = -3, CM_P = -6, CSD_A = 15, fixedfy = 3, theta = 1, seed = 1)
dat2 <- simHCE(n = 2500, TTE_A = Rates_A, TTE_P = Rates_P,
             CM_A = -3, CM_P = -6, CSD_A = 15, fixedfy = 3, theta = 1.0001, seed = 1)
calcWO(dat1)
calcWO(dat2)
```

---

 simKHCE

*Simulate a kidney disease hce dataset*


---

## Description

Simulate a kidney disease hce dataset, capturing eGFR (Estimated Glomerular Filtration Rate) progression over time, along with a competing and dependent terminal event: KFRT (Kidney Failure Replacement Therapy)

## Usage

```
simKHCE(
  n,
  CM_A,
  CM_P = -4,
  n0 = n,
  TTE_A = 10,
  TTE_P = TTE_A,
  fixedfy = 2,
  Emin = 20,
  Emax = 100,
  sigma = 8,
  Sigma = 3,
```



```

    m = 10,
    theta = -0.23,
    phi = 0
  )

```

### Arguments

|         |  |
|---------|--|
| n       | sample size in the active treatment group.   |
| CM_A    | annualized eGFR slope in the active group.   |
| CM_P    | annualized eGFR slope in the control group.  |
| n0      | sample size in the control treatment group.  |
| TTE_A   | event rate per year in the active group for KFRT.  |
| TTE_P   | event rate per year in the placebo group for KFRT.   |
| fixedfy | length of follow-up in years.  |
| Emin    | lower limit of eGFR at baseline.   |
| Emax    | upper limit of eGFR at baseline.   |
| sigma   | within-patient standard deviation.   |
| Sigma   | between-patient standard deviation.  |
| m       | number of equidistant visits.  |
| theta   | coefficient of dependence of eGFR values and the risk of KFRT.   |
| phi     | coefficient of proportionality (between 0 and 1) of the treatment effect. The case of 0 corresponds to the uniform treatment effect. |

### Value

a list containing the dataset GFR for longitudinal measurements of eGFR and the competing KFRT events, the dataset ADET for the time-to-event kidney outcomes (sustained declines or sustained low levels of eGFR), and the combined HCE dataset for the kidney hierarchical composite endpoint.

### See Also

[simHCE\(\)](#) for a general function of simulating hce datasets.

### Examples

```

# Example 1
set.seed(2022)
L <- simKHCE(n = 1000, CM_A = -3.25)
dat <- L$HCE
calcWO(dat)

```

---

|        |   |
|--------|---|
| simORD | <i>Simulate ordinal variables for two treatment groups using categorization of beta distributions</i> |
|--------|---|

---

### Description

Simulate ordinal variables for two treatment groups using categorization of beta distributions

### Usage

```
simORD(n, n0 = n, M, alpha1 = 8, beta1 = 7, alpha0 = 5, beta0 = 5)
```

### Arguments

|        |  |
|--------|--|
| n      | sample size in the active treatment group.                       |
| n0     | sample size in the placebo group.                                |
| M      | number of ordinal values to be simulated.                        |
| alpha1 | shape1 parameter for the beta distribution in the active group.  |
| beta1  | shape2 parameter for the beta distribution in the active group.  |
| alpha0 | shape1 parameter for the beta distribution in the placebo group. |
| beta0  | shape2 parameter for the beta distribution in the placebo group. |

### Value

a data frame containing the following columns:

- ID subject identifier.
- TRTP planned treatment group - "A" for active, "P" for Placebo.
- GROUPN ordinal values. The number of unique values is specified by the variable M0.
- tau the theoretical win odds.
- theta the theoretical win probability.

### See Also

[simHCE\(\)](#) for simulating hce objects.

### Examples

```
# Example 1
set.seed(2024)
alpha1 <- 8
beta1 <- 8
alpha0 <- 4
beta0 <- 5
d <- simORD(n = 1500, n0 = 1500, M = 5, alpha1 = alpha1, beta1 = beta1,
```

```

alpha0 = alpha0, beta0 = beta0)
x <- seq(0, 1, 0.01)
plot(x, dbeta(x, shape1 = alpha1, shape2 = beta1),
type = "l", ylab = "Density of beta distribution", col = 2)
lines(x, dbeta(x, shape1 = alpha0, shape2 = beta0), col = 3, lty = 2)
legend("topleft", lty = c(1, 2), col = c(2, 3), legend = c("Control", "Active"))
D <- hce(GROUP = d$GROUPN, TRTP = d$TRTP)
table(D$TRTP, D$GROUPN)
calcWO(D)
# Example 2
set.seed(2024)
d <- simORD(n = 100, n0 = 50, M = 2)
d_hce <- hce(GROUP = d$GROUPN, TRTP = d$TRTP)
calcWO(d_hce)
### compare with the theoretical values of the continuous distributions
c(tau = unique(d$tau), theta = unique(d$theta))
# Example 2 - Convergence of the win odds to its theoretical value
set.seed(2024)
N <- NULL
size <- c(seq(10, 500, 1))
for(i in size){
  d <- simORD(n = i, M = 2)
  d_hce <- hce(GROUP = d$GROUPN, TRTP = d$TRTP)
  TAU <- calcWO(d_hce)
  D <- data.frame(WO = TAU$WO, n = i, tau = unique(d$tau))
  N <- rbind(N, D)
}
plot(N$n, N$WO, log = "y", ylim = c(0.5, 2), ylab = "Win Odds", xlab = "Sample size", type = "l")
lines(N$n, N$tau, col = "darkgreen", lty = 2, lwd = 2)
abline(h = 1, lty = 4, col = "red")
legend("bottomright", legend = c("Theoretical Win Odds", "Null", "Win Odds Estimate"),
lty = c(4, 2, 1), col = c("darkgreen", "red", "black"))
title("Convergence of the win odds to its theoretical value")

```

---

sizeWO

*Sample size calculation for the win odds test (no ties)*


---

## Description

Sample size calculation for the win odds test (no ties)

## Usage

```

sizeWO(
  WO,
  power,
  SD = NULL,
  k = 0.5,
  alpha = 0.05,
  WOnull = 1,

```

```

    alternative = c("shift", "max", "ordered")
  )

```

### Arguments

|             |   |
|-------------|---|
| WO          | a numeric vector of win odds values.  |
| power       | the given power. A numeric vector of length 1.  |
| SD          | assumed standard deviation of the win proportion. By default uses the conservative SD. A numeric vector of length 1.  |
| k           | proportion of active group in the overall sample size. Default is 0.5 (balanced randomization). A numeric vector of length 1.   |
| alpha       | the significance level for the 2-sided test. Default is 0.05. A numeric vector of length 1.   |
| WOnull      | the win odds value of the null hypothesis (default is 1). A numeric vector of length 1.   |
| alternative | a character string specifying the class of the alternative hypothesis, must be one of "shift" (default), "max" or "ordered". You can specify just the initial letter. |

### Details

alternative = "max" refers to the maximum variance of the win proportion across all possible alternatives. The maximum variance equals  $WP*(1 - WP)/k$  where the win probability is calculated as  $WP = WO/(WO + 1)$ . alternative = "shift" specifies the variance across alternatives from a shifted family of distributions (Wilcoxon test). The variance formula, as suggested by Noether, is calculated based on the null hypothesis as follows  $1/(12*k*(1 - k))$ . alternative = "ordered" specifies the variance across alternatives from stochastically ordered distributions which include shifted distributions.

### Value

a data frame containing the sample size with input values.

### References

- All formulas were presented in
  - Bamber D (1975) "The area above the ordinal dominance graph and the area below the receiver operating characteristic graph." *Journal of Mathematical Psychology* 12.4: 387-415. doi:10.1016/0022-2496(75)90001-2.
- Noether's formula for shifted alternatives
  - Noether GE (1987) "Sample size determination for some common nonparametric tests." *Journal of the American Statistical Association* 82.398: 645-7. doi:10.1080/01621459.1987.10478478.
- For shift alternatives see also

Gasparyan SB et al. (2021) "Power and sample size calculation for the win odds test: application to an ordinal endpoint in COVID-19 trials." *Journal of Biopharmaceutical Statistics* 31.6: 765-787. doi:10.1080/10543406.2021.1968893.

### See Also

`powerWO()`, `minWO()` for WO power or minimum detectable WO calculation.

### Examples

```
sizeWO(WO = 1.25, power = 0.9)
sizeWO(WO = 1.25, power = 0.9, k = 0.75)
sizeWO(WO = seq(1.05, 1.5, 0.05), power = 0.9)
# Comparison of different alternatives
x <- seq(1.05, 5, 0.05)
N1 <- sizeWO(WO = x, power = 0.9, alternative = "m")$SampleSize
N2 <- sizeWO(WO = x, power = 0.9, alternative = "o")$SampleSize
N3 <- sizeWO(WO = x, power = 0.9, alternative = "s")$SampleSize
d <- data.frame(WO = x, N_m = N1, N_o = N2, N_s = N3)
## Check the power for the ordered alternative
check <- c()
for(i in seq_along(x)){
  check[i] <- powerWO(N = d[i, "N_o"], WO = d[i, "WO"], alternative = "o")$power
}
d$power_check_o <- check
print(d)
```

---

|        |   |
|--------|---|
| sizeWR | <i>Sample size calculation for the win ratio test (with WR = 1 null hypothesis)</i> |
|--------|---|

---

### Description

Sample size calculation for the win ratio test (with WR = 1 null hypothesis)

### Usage

```
sizeWR(WR, power, WO = NULL, Pties = NULL, k = 0.5, alpha = 0.05)
```

### Arguments

|       |   |
|-------|---|
| WR    | a numeric vector of win odds values.  |
| power | the given power. A numeric vector of length 1.  |
| WO    | win odds. Should be specified only if Pties is not specified. A numeric vector of length 1.                                   |
| Pties | probability of ties. A numeric vector of length 1.  |
| k     | proportion of active group in the overall sample size. Default is 0.5 (balanced randomization). A numeric vector of length 1. |
| alpha | the significance level for the 2-sided test. Default is 0.05. A numeric vector of length 1.                                   |

**Value**

a data frame containing the sample size with input values.

**References**

Yu RX, Ganju J. (2022) "Sample size formula for a win ratio endpoint." *Statistics in Medicine*, 41.6: 950-63. doi:10.1002/sim.9297.

**See Also**

[sizeWO\(\)](#) for WO sample size calculation.

**Examples**

```
sizeWR(WR = 1.35, Pties = 0.125, power = 0.8)
sizeWR(WR = 1.35, WO = 1.3, power = seq(0.5, 0.9, 0.05))
```

---

stratWO

*A generic function for stratified win odds with adjustment*

---

**Description**

A generic function for stratified win odds with adjustment

**Usage**

```
stratWO(x, ...)
```

**Arguments**

x                    an object used to select a method.  
...                   further arguments passed to or from other methods.

**Value**

a list containing the stratified results and results by strata.

**See Also**

[stratWO.data.frame\(\)](#) methods.

---

stratWO.data.frame      *Stratified win odds with adjustment*

---

## Description

Stratified win odds with adjustment

## Usage

```
## S3 method for class 'data.frame'
stratWO(
  x,
  AVAL,
  TRTP,
  STRATA,
  ref,
  COVAR = NULL,
  alpha = 0.05,
  WOnull = 1,
  ...
)
```

## Arguments

|        |  |
|--------|--|
| x      | a data frame containing subject-level data.        |
| AVAL   | variable in the data with ordinal analysis values. |
| TRTP   | the treatment variable in the data.                |
| STRATA | a character variable for stratification.           |
| ref    | the reference treatment group.                     |
| COVAR  | a numeric covariate.                               |
| alpha  | the reference treatment group.                     |
| WOnull | the null hypothesis. The default is 1.             |
| ...    | additional parameters.                             |

## Value

a data frame containing the following columns:

- WO stratified (or adjusted/stratified) win odds.
- LCL lower confidence limit for adjusted (or adjusted/stratified) WO.
- UCL upper confidence limit for adjusted (or adjusted/stratified) WO.
- SE standard error of the adjusted (or adjusted/stratified) win odds.
- WOnull win odds of the null hypothesis (specified in the WOnull argument).

- alpha two-sided significance level for calculating the confidence interval (specified in the alpha argument).
- Pvalue p-value associated with testing the null hypothesis.
- WP adjusted (or adjusted/stratified) win probability.
- LCL\_WP lower confidence limit for adjusted (or adjusted/stratified) WP.
- UCL\_WP upper confidence limit for adjusted (or adjusted/stratified) WP.
- SE\_WP standard error for the adjusted (or adjusted/stratified) win probability.
- SD\_WP standard deviation of the adjusted (or adjusted/stratified) win probability.
- N total number of patients in the analysis.
- Type "STRATIFIED" or "STRATIFIED/ADJUSTED" depending on whether COVAR is specified.

## References

Gasparyan SB et al. (2021) "Adjusted win ratio with stratification: calculation methods and interpretation." *Statistical Methods in Medical Research* 30.2: 580-611. doi:10.1177/0962280220942558.

## See Also

[stratWO\(\)](#).

## Examples

```
# Stratified win odds
res <- stratWO(x = KHCE, AVAL = "AVAL", TRTP = "TRTP",
              STRATA = "STRATAN", ref = "P")

res
## Compare with the non-stratified win odds
res0 <- calcWO(AVAL ~ TRTP, data = KHCE, ref = "P")
res0
## Compare with the win odds in each stratum separately
l <- lapply(split(KHCE, KHCE$STRATAN), calcWO, AVAL = "AVAL", TRTP = "TRTP", ref = "P")
l <- do.call(rbind, l)
l <- l[, c("WO", "LCL", "UCL", "N")]
l$STRATA <- as.numeric(row.names(l))
plot(y = l$WO, x = l$STRATA, ylim = c(0.5, 2.5), log = "y", xlim = c(0, 6),
     ylab = "Win Odds", xlab = "", xaxt = "n")
axis(1, at = 1:6, labels = c(paste0("STR = ", l$STRATA), "Stratified", "Non-stratified"))
arrows(l$STRATA, l$LCL, l$STRATA,
       l$UCL, angle = 90, code = 3, length = 0.05, col = "darkgreen")
points(5, res$WO)
arrows(5, res$LCL, 5, res$UCL, angle = 90, code = 3,
      length = 0.05, col = "darkblue")
abline(h = c(1, res$WO), col = "red", lty = 4)
points(6, res0$WO)
arrows(6, res0$LCL, 6, res0$UCL, angle = 90, code = 3,
      length = 0.05, col = "darkred")
# Stratified and adjusted win odds
res <- stratWO(x = KHCE, AVAL = "AVAL", COVAR = "EGFRBL",
```



```

      TRTP = "TRTP", STRATA = "STRATAN", ref = "P")
res

```

---

summaryWO *A generic function for summarizing win odds*

---

### Description

A generic function for summarizing win odds

### Usage

```
summaryWO(x, ...)
```

### Arguments

x                    an object used to select a method.  
 ...                  further arguments passed to or from other methods.

### Value

a data frame containing calculated values.

### See Also

[summaryWO.hce\(\)](#), [summaryWO.formula\(\)](#), [summaryWO.data.frame\(\)](#) methods.

---

summaryWO.data.frame *Win odds summary for a data frame*

---

### Description

Win odds summary for a data frame

### Usage

```
## S3 method for class 'data.frame'
summaryWO(x, AVAL, TRTP, ref, GROUP = NULL, ...)
```

### Arguments

x                    a data frame containing subject-level data.  
 AVAL                variable in the data with ordinal analysis values.  
 TRTP                the treatment variable in the data.  
 ref                 the reference treatment group.  
 GROUP              an optional variable for grouping.  
 ...                 additional parameters.

**Value**

a list containing the summary of wins, losses, and ties. It contains the following named objects:

- summary a data frame containing number of wins, losses, and ties by treatment group and the overall number of comparisons.
- summary\_by\_GROUP (if GROUP variable is specified) a summary data frame by GROUP.
- WO calculated WO (win odds) and WP (win probability) and their standard errors.

**See Also**

[calcWO\(\)](#), [summaryWO\(\)](#), [summaryWO.formula\(\)](#), [summaryWO.hce\(\)](#) methods.

**Examples**

```
summaryWO(x = HCE3, AVAL = "AVAL", TRTP = "TRTP", ref = "P", GROUP = "GROUP")
```

---

|                   |  |
|-------------------|--|
| summaryWO.formula | <i>Win odds summary using formula syntax</i> |
|-------------------|--|

---

**Description**

Win odds summary using formula syntax

**Usage**

```
## S3 method for class 'formula'
summaryWO(x, data, ...)
```

**Arguments**

|      |                             |
|------|-----------------------------|
| x    | an object of class formula. |
| data | a data frame.               |
| ...  | additional parameters.      |

**Value**

a list containing the summary of wins, losses, and ties. It contains the following named objects:

- summary a data frame containing number of wins, losses, and ties by treatment group and the overall number of comparisons.
- WO calculated WO (win odds) and WP (win probability) and their standard errors.
- formula returning the specified formula in the x argument.
- ref showing how the reference group was selected. Can be modifying by specifying the ref argument.

**See Also**

[calcWO\(\)](#), [summaryWO\(\)](#), [summaryWO.data.frame\(\)](#), [summaryWO.hce\(\)](#) methods.

**Examples**

```
summaryWO(data = COVID19, GROUP ~ TRTP)
summaryWO(data = COVID19, GROUP ~ TRTP, GROUP = "GROUP", ref = "Placebo")
```

---

|               |   |
|---------------|---|
| summaryWO.hce | <i>Win odds summary for hce objects</i> |
|---------------|---|

---

**Description**

Win odds summary for hce objects

**Usage**

```
## S3 method for class 'hce'
summaryWO(x, ...)
```

**Arguments**

```
x          an hce object.
...        additional parameters.
```

**Value**

a list containing the summary of wins, losses, and ties. It contains the following named objects:

- summary a data frame containing number of wins, losses, and ties by treatment group and the overall number of comparisons.
- summary\_by\_GROUP (if GROUP variable is specified) a summary data frame by GROUP.
- WO calculated WO (win odds) and WP (win probability) and their standard errors.

**See Also**

[calcWO\(\)](#), [summaryWO\(\)](#), [summaryWO.data.frame\(\)](#), [summaryWO.formula\(\)](#) methods.

**Examples**

```
dat <- as_hce(HCE4)
summaryWO(dat, ref = "P")
```

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