

# Package ‘MannWhitneyCopula’

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**Type** Package

**Title** Computing Mann-Whitney Effect Based on Copulas

**Version** 0.1.1

**Description** Computing the Mann-Whitney effect based on copula models.

Estimation of the association parameter in survival copula models.

A description of the underlying methods is described in Nakazono et al. (2024) <[doi:10.3390/math12101453](https://doi.org/10.3390/math12101453)> and Nakazono et al. (accepted for publication in Statistical Papers).

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**Author** Kosuke Nakazono [aut, cre]

**Maintainer** Kosuke Nakazono <nakazono@ism.ac.jp>

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**MW.comp***Parametric calculation for the Mann-Whitney effect under survival copula models*

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## Description

`MW.comp` provides a parametric estimator for the Mann-Whitney effect under the parametric survival functions and copulas. See Nakazono, et al.(2024) for details.

## Usage

```
MW.comp(
  copula = c("clayton", "gumbel", "frank", "fgm", "gb"),
  copula.param = 1,
  s1 = c("exponential", "weibull", "gamma", "log-normal", "burr3"),
  S1.param,
  s2 = c("exponential", "weibull", "gamma", "log-normal", "burr3"),
  S2.param
)
```

## Arguments

<code>copula</code>	copula family. Available options include; "clayton", "gumbel", "frank", "fgm", "gb".
<code>copula.param</code>	the copula parameter.
<code>s1</code>	a parametric survival function for S1. Available options include; "exponential", "weibull", "gamma", "log-normal", "burr3".
<code>S1.param</code>	the distribution parameter for S1.
<code>s2</code>	a parametric survival function for S2. Available options include; "exponential", "weibull", "gamma", "log-normal", "burr3".
<code>S2.param</code>	the distribution parameter for S2.

## Details

`copula.param` is restricted as below:

- "clayton";  $\text{copula.param} \geq 0$
- "gumbel";  $\text{copula.param} \geq 0$
- "frank";  $-\infty < \text{copula.param} < \infty$
- "fgm";  $-1 \leq \text{copula.param} \leq 1$
- "gb";  $-1 \leq \text{copula.param} \leq 1$

## Value

estimate of the Mann-Whitney effect.

## References

Nakazono, K., Lin, Y. C., Liao, G. Y., Uozumi, R., & Emura, T. (2024). Computation of the Mann–Whitney effect under parametric survival copula models. *Mathematics*, 12(10), 1453.

## Examples

```
# Under the exponential survival functions and Clayton copula
MW.comp(
  copula = "clayton",
  copula.param = 1,
  s1 = "exponential",
  S1.param = 1,
  s2 = "exponential",
  S2.param = 2
)
```

MW.Copula

*Parametric calculation for the Mann-Whitney effect under the parametric copulas*

## Description

MW.Copula provides a parametric estimator and confidence interval for the Mann-Whitney effect under the parametric survival functions and copulas. The result of this function includes results for logit-transformed estimators. See Nakazono, et al. (2025) for details.

## Usage

```
MW.Copula(
  t.event, event, group,
  copula = c("clayton", "gumbel", "frank", "fgm", "gb"),
  copula.param = 1,
  s1 = c("exponential", "weibull", "gamma", "log-normal", "burr3"),
  s2 = c("exponential", "weibull", "gamma", "log-normal", "burr3"),
  par1 = c(0, 0),
  par2 = c(0, 0),
  alpha = 0.05,
  logit = FALSE
)
```

## Arguments

t.event	a vector of time-to-event data.
event	a vector for event indicator.
group	a vector for group indicator.

<code>copula</code>	copula family.
<code>copula.param</code>	the copula parameter.
<code>s1</code>	a parametric survival function for S1. Available options include; "exponential", "weibull", "gamma", "log-normal", "burr3".
<code>s2</code>	a parametric survival function for S2. Available options include; "exponential", "weibull", "gamma", "log-normal", "burr3".
<code>par1</code>	initial value(s) for the parameters of S1.
<code>par2</code>	initial value(s) for the parameters of S2.
<code>alpha</code>	significance level.
<code>logit</code>	logical; if TRUE, the estimator and CI is logit-transformed.

### Value

parameter: a vector of MLE.  
 estimate: the Mann-Whitney effect estimator  
 SE: the standard error  
 CI: the 1-alpha% confidence interval for Mann-Whitney effect  
 P.value: the P-value for testing the null hypothesis H0: p=1/2.  
 kendall: Kendall's tau  
 logit: the estimator and CI are logit-transformed or not.

### References

Nakazono, K., Uozumi, R., & Emura, T. (2025). Parametric inference for the Mann-Whitney effect under survival copula models, Statistical Papers, in press.

### Examples

```

##Mann-Whitney effect under exponential distributions
#set distribution parameter
lambda1 = 1
lambda2 = 2

#generate time to event
u = runif(100)
t.event1 = -log(u) / lambda1
t.event1 = sort(t.event1)

v = runif(100)
t.event2 = -log(v) / lambda2
t.event2 = sort(t.event2)

#censoring indicator
t1c = runif(100, 0, 1.5)
t.event1 = (t1c >= t.event1) * t.event1 + (t1c < t.event1) * t1c
event1 = 1 * (t1c > t.event1)

```

```
t2c = runif(100, 0, 0.8)
t.event2 = (t2c >= t.event2) * t.event2 + (t2c < t.event2) * t2c
event2 = 1 * (t2c > t.event2)

t.event = c(t.event1, t.event2)
event = c(event1, event2)

#group indicator
group = rep(c(1, 0), each = 100)

MW.Copula(t.event, event, group,
           copula = "clayton", copula.param = 1,
           s1 = "exponential", s2 = "exponential", par1 = c(0, 0), par2 = c(0, 0), logit = FALSE)
```

**MW.plot***Plot method for Mann-Whitney effect under parametric copula models***Description**

`MW.plot` plots the results, the parametric estimator and their confidence intervals, for the Mann-Whitney effect under parametric survival functions and copulas.

**Usage**

```
MW.plot(t.event, event, group,
        copula = c("clayton", "gumbel", "frank", "fgm", "gb"),
        lower = 0, upper = 1,
        s1 = c("exponential", "weibull", "gamma", "log-normal", "burr3"),
        s2 = c("exponential", "weibull", "gamma", "log-normal", "burr3"),
        par1 = c(0, 0),
        par2 = c(0, 0),
        alpha = 0.05,
        logit = FALSE,
        xaxis = 2
      )
```

**Arguments**

- |                      |   |
|----------------------|---|
| <code>t.event</code> | a vector for time-to-event.                                   |
| <code>event</code>   | a vector for event indicator.                                 |
| <code>group</code>   | a vector for group indicator.                                 |
| <code>copula</code>  | copula family.  |
| <code>lower</code>   | the lower end points of the interval of the copula parameter. |
| <code>upper</code>   | the lower end points of the interval of the copula parameter. |

s1	a parametric survival function for S1. Available options include; "exponential", "weibull", "gamma", "log-normal", "burr3".
s2	a parametric survival function for S2. Available options include; "exponential", "weibull", "gamma", "log-normal", "burr3".
par1	initial value(s) for the parameters of S1.
par2	initial value(s) for the parameters of S2.
alpha	significance level.
logit	logical; if TRUE, the estimator and CI is logit-transformed.
xaxis	a indicator specifying whether the xaxis is the copula parameter (xaxis = 1) or Kendall's tau (xaxis = 2).

**Value**

No return value, called for side effects (generates a plot).

**Examples**

```
##Exponential distributions
#set distribution parameter
lambda1 = 1
lambda2 = 2

#generate time to event
u = runif(100)
t.event1 = -log(u) / lambda1
t.event1 = sort(t.event1)

v = runif(100)
t.event2 = -log(v) / lambda2
t.event2 = sort(t.event2)

#censoring indicator
t1c = runif(100, 0, 1.5)
t.event1 = (t1c >= t.event1) * t.event1 + (t1c < t.event1) * t1c
event1 = 1 * (t1c > t.event1)

t2c = runif(100, 0, 0.8)
t.event2 = (t2c >= t.event2) * t.event2 + (t2c < t.event2) * t2c
event2 = 1 * (t2c > t.event2)

t.event = c(t.event1, t.event2)
event = c(event1, event2)

#group indicator
group = rep(c(1, 0), each = 100)

MW.plot(t.event, event, group,
        copula = "clayton",
        lower = 0.2, upper = 0.8,
        s1 = "exponential", s2 = "exponential",
```

---

```
par1 = c(0, 0), par2 = c(0, 0), alpha = 0.05, logit = FALSE, xaxis = 2)
```

---

**survival.mle***Maximum likelihood estimation under parametric survival function***Description**

`survival.mle` provides the maximum likelihood estimator and their variance-covariance matrix under parametric survival functions. This function also provides AIC and Kolmogorov-Smirnov distance to evaluate the model fitting.

**Usage**

```
survival.mle(
  t.event,
  event,
  distribution = c("exponential", "weibull", "gamma", "log-normal", "burr3"),
  par = c(0,0))
```

**Arguments**

<code>t.event</code>	a vector for time-to-event.
<code>event</code>	a vector for event indicator.
<code>distribution</code>	a parametric distribution for survival function. Available options include; "exponential", "weibull", "gamma", "log-normal", "burr3".
<code>par</code>	initial value(s) for the distribution parameters.

**Value**

`estimate`: a vector of MLE.  
`var`: variance-covariance matrix of MLE.  
`AIC`: Akaike information criteria.  
`KS`: Kolmogorov-Smirnov distance between MLE and KM estimator.

**Examples**

```
#MLE under exponential distribution
#set distribution parameter
lambda = 1

#generate time to event
u = runif(100)
t.event = -log(u)/lambda
t.event = sort(t.event)
```

```
#censoring indicator  
tc = runif(100, 0, 1.5)  
t.event = (tc >= t.event) * t.event + (tc < t.event) * tc  
event = 1 * (tc > t.event)  
  
survival.mle(t.event, event, distribution = "exponential")
```

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