

Package ‘rankICC’

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Type Package

Title Rank Intraclass Correlation for Clustered Data

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Description Estimates the rank intraclass correlation coefficient (ICC) for clustered continuous and ordinal data. See Tu et al. (2023) <[DOI:10.1002/sim.9864](https://doi.org/10.1002/sim.9864)> for details.

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Collate 'rankICC.R' 'rankICC3levels.R'

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NeedsCompilation no

Author Shengxin Tu [aut, cre],
Chun Li [aut],
Donglin Zeng [aut],
Bryan Shepherd [aut]

Maintainer Shengxin Tu <shengxin.tu@vanderbilt.edu>

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rankICC

*Rank ICC with two hierarchies***Description**

rankICC computes the rank intraclass correlation coefficient (ICC) of a two-level hierarchical distribution. It can be used with any orderable variable, including continuous and discrete variables. Different weighting methods are provided, including methods assigning equal weights to observations or to clusters.

Usage

```
rankICC(
  x,
  cluster,
  weights = c("obs", "clusters", "ess", "combination"),
  conf.int = 0.95,
  fisher = FALSE,
  na.rm = FALSE,
  ...
)
```

Arguments

<code>x</code>	a numeric or factor vector.
<code>cluster</code>	a vector of cluster index corresponding to <code>x</code> .
<code>weights</code>	a character string indicating which weighting method is used. Or an optional vector of user-defined weights to be used. Should be one of the strings "obs", "clusters", "ess", "combination", or a numeric vector. Default is "obs". See Details.
<code>conf.int</code>	numeric specifying confidence interval level.
<code>fisher</code>	logical, indicating whether to apply Fisher transformation to compute confidence intervals.
<code>na.rm</code>	logical. Should missing values be removed?
<code>...</code>	additional arguments to be passed to the iteration function if <code>weights</code> is "ess" or "combination". Specifying the tolerance via "tol" and the maximum iteration times via "maxIter".

Details

"obs" assigns equal weights to observations; $p_{ij} = 1/N$, where N is the total number of observations. "clusters" assigns equal weights to clusters; $p_{ij} = 1/(nk_i)$, where n is the total number of clusters and k_i is the cluster size. "ess" and "combination" implement iterations until convergence; $p_{ij}(\gamma_I) = 1/(1+k_i\gamma_I)/\{\sum_{j=1}^n k_j/(1+k_j\gamma_I)\}$ for "ess", $p_{ij}(\gamma_I) = (1-\gamma_I)/N + \gamma_I/(nk_i)$ for "combination".

Value

a vector with following components.

rankICC	the rank ICC.
SE	the standard error.
Lower, Upper	the lower and upper bound of the confidence interval.

References

Tu, S., Li, C., Zeng, D., and Shepherd, B. E. (2023). Rank intraclass correlation for clustered data. *Statistics in Medicine* 42, 4333-4348.

Examples

```
k <- 50; m <- 5
sigma.u <- 1; sigma.e <- 2
u <- rnorm(k, 5, sigma.u)
x1 <- matrix(NA, k, m)
for (i in 1:k){
  x1[i,] <- u[i] + rnorm(5, 0, sigma.e)
}
x <- as.vector(t(x1))
cluster <- rep(1:k, each=5)
rankICC(x, cluster, weights = "clusters")
rankICC(x, cluster, weights = "ess", tol = 1e-4, maxIter = 10)
```

rankICC3levels	<i>Rank ICC with three hierarchies</i>
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Description

rankICC3levels computes the rank intraclass correlation coefficient (ICC) with three hierarchies. Starting from the innermost level, the three levels are named level 1, level 2, and level 3. The rank ICC at level 2 evaluates the rank correlation between a random pair from the same level-2 unit. The rank ICC at level 3 evaluates the rank correlation between a random pair from the same level-3 unit but different level-2 units.

Usage

```
rankICC3levels(
  x,
  level2,
  level3,
  weights = c("level1", "level2", "level3"),
  conf.int = 0.95,
```

```

    fisher = FALSE,
    na.rm = FALSE
  )

```

Arguments

<code>x</code>	a numeric or factor vector.
<code>level2</code>	a vector indicating level-2 membership.
<code>level3</code>	a vector indicating level-3 membership.
<code>weights</code>	a character string indicating which weighting method is used. Or an optional vector of user-defined weights to be used. Should be one of the strings "level1", "level2", "level3", or a numeric vector. Default is "level1". See Details.
<code>conf.int</code>	numeric specifying confidence interval level.
<code>fisher</code>	logical, indicating whether to apply Fisher transformation to compute confidence intervals.
<code>na.rm</code>	logical. Should missing values be removed?

Details

"level1" assigns equal weights to level-1 units; $p_{ijk} = 1/(\sum_{i=1}^n \sum_{j=1}^{n_i} m_{ij})$, where n is the total number of level-3 units, n_i is the number of level-2 units in the i th level-3 unit, and m_{ij} is the number of level-1 units in the j th level-2 unit and the i th level-3 unit. "level2" assigns equal weights to level-2 units; $p_{ijk} = 1/(m_{ij} \sum_{i=1}^n n_i)$. "level3" assigns equal weights to level-3 units; $p_{ijk} = 1/(n n_i m_{ij})$.

Value

a matrix with two rows. The first row is for rank ICC at level 2 and the second row is for rank ICC at level 3. Each row has the following components.

rankICC	the rank ICC.
SE	the standard error.
Lower, Upper	the lower and upper bound of the confidence interval.

References

Tu, S., Li, C., Zeng, D., and Shepherd, B. E. (2023). Rank intraclass correlation for clustered data. *Statistics in Medicine* 42, 4333-4348.

Examples

```

k <- 50; m <- 5
sigma.u <- 1; sigma.e <- 2
u <- rnorm(k, 5, sigma.u)
x1 <- matrix(NA, k, m)

```

```
for (i in 1:k){  
  x1[i,] <- u[i] + rnorm(5, 0, sigma.e)  
}  
x <- as.vector(t(x1))  
level2 <- rep(1:k, each=5)  
level3 <- round(level2 / 10)  
rankICC3levels(x, level2, level3, weights = "level2")
```

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