

Package: ESTER (via r-universe)

September 11, 2024

Title Efficient Sequential Testing with Evidence Ratios

Version 0.2.1

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Description An implementation of sequential testing that uses evidence ratios computed from the weights of a set of models. These weights correspond either to Akaike weights computed from the Akaike Information Criterion (AIC) or the Bayesian Information Criterion (BIC) and following Burnham & Anderson (2004, <[doi:10.1177/0049124104268644](https://doi.org/10.1177/0049124104268644)>) recommendations, or to pseudo-BMA weights computed from the WAIC or the LOOIC of models fitted with 'brms' and following Yao et al. (2017, <[arXiv:1704.02030v3](https://arxiv.org/abs/1704.02030v3)>).

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LazyData yes

RoxygenNote 6.0.1

Depends R (>= 3.3.0)

Imports brms, lme4, dplyr, magrittr, tidyr, ggplot2, rlang

URL <https://github.com/lNALBORczyk/ESTER>

BugReports <https://github.com/lNALBORczyk/ESTER/issues>

Suggests knitr, rmarkdown, testthat

VignetteBuilder knitr

Repository <https://lNALBORczyk.r-universe.dev>

RemoteUrl <https://github.com/lNALBORczyk/ester>

RemoteRef HEAD

RemoteSha eee73e59b3e62caa936d64d563b6fa9d69e593b7

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<i>aic</i>	<i>Computes the Akaike Information Criterion</i>
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Description

Computes the Akaike Information Criterion (AIC) of a model, or the second-order bias correction for small samples (AICc), as suggested by Burnham & Anderson (2002, 2004).

Usage

```
aic(mod, correction = TRUE)
```

Arguments

<code>mod</code>	A fitted model of class <code>lm</code> or <code>merMod</code> .
<code>correction</code>	Should we apply the second-order correction (default to TRUE) ?

Author(s)

Ladislav Nalborczyk <<ladislav.nalborczyk@gmail.com>>

References

Burnham, K. P., \& Anderson, D. R. (2002). Model Selection and Multimodel Inference: A Practical Information-Theoretical Approach. 2d ed. New York: Springer-Verlag.

Burnham, K. P., \& Anderson, D. R. (2004). Multimodel inference: Understanding AIC and BIC in model selection. *Sociological Methods and Research*, 33(2), 261-304.

See Also

[bic](#), [ictab](#)

Examples

```
data(mtcars)
mod1 <- lm(mpg ~ cyl, mtcars)
aic(mod1)
```

bic	<i>Computes the Bayesian Information Criterion</i>
-----	--

Description

Computes the Bayesian Information Criterion of a model (Schwarz, 1978).

Usage

```
bic(mod)
```

Arguments

mod A fitted model of class `lm` or `merMod`.

Author(s)

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References

Schwarz, G. (1978). Estimating the dimension of a model. *Annals of Statistics*, 6, 461-464.

See Also

[aic](#), [ictab](#)

Examples

```
data(mtcars)
mod1 <- lm(mpg ~ cyl, mtcars)
bic(mod1)
```

ESTER	<i>Efficient Sequential Testing with Evidence Ratios</i>
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Description

The **ESTER** package implements sequential testing based on evidence ratios computed from the Akaike weights of a set of models. These weights are being computed using either the Akaike Information Criterion (AIC) or the Bayesian Information Criterion (BIC).

Details

See `vignette("ESTER")` for a general introduction and overview.

Author(s)

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See Also

[ictab](#), [seqtest](#)

ictab	<i>Computes Akaike weights or pseudo-BMA weights of a set of models</i>
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Description

Returns a table with weights of a set of models, based on various information criteria. Currently, `ictab` supports the computation of Akaike weights from the `aic` or the `bic` computed on `lm` or `merMod` models, as well as the computation of pseudo-BMA weights, computed from the WAIC or LOOIC of `brmsfit` models.

Usage

```
ictab(mods, ic, ...)
```

Arguments

<code>mods</code>	Should be a named list of models, of class <code>lm</code> , <code>merMod</code> or <code>brmsfit</code> .
<code>ic</code>	Indicates which information criterion to use. Current supported information criteria include <code>aic</code> and <code>bic</code> for <code>lm</code> and <code>merMod</code> models, as well as WAIC and LOO for <code>brmsfit</code> models.
<code>...</code>	Additional parameters to be passed to the <code>ic</code> function.

Value

An object of class `data.frame`, which contains the value of the information criterion (either AIC, BIC, WAIC or LOOIC), the number of parameters (`k` for AIC and BIC or `p` for WAIC or LOOIC), the `delta_IC` (for AIC and BIC) or the `elpd` for models compared with WAIC or LOOIC, and the weight of each model (Akaike weights for AIC or BIC and pseudo-BMA weights for WAIC or LOOIC).

Author(s)

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References

- Burnham, K. P., \& Anderson, D. R. (2002). Model Selection and Multimodel Inference: A Practical Information-Theoretical Approach. 2d ed. New York: Springer-Verlag.
- Burnham, K. P., \& Anderson, D. R. (2004). Multimodel inference: Understanding AIC and BIC in model selection. *Sociological Methods and Research*, 33(2), 261-304.
- Yao, Y. P., Vehtari, A., Simpson, D., \& Gelman, A. (2017). Using stacking to average Bayesian predictive distributions.

See Also

[aic](#), [bic](#)

Examples

```
data(mtcars)
mod1 <- lm(mpg ~ cyl, mtcars)
mod2 <- lm(mpg ~ cyl + vs, mtcars)
mod3 <- lm(mpg ~ cyl * vs, mtcars)
mods <- list(mod1 = mod1, mod2 = mod2, mod3 = mod3)
ictab(mods, aic)

## Not run:
library(brms)
mod1 <- brm(mpg ~ cyl, mtcars)
mod2 <- brm(mpg ~ cyl + vs, mtcars)
mods <- list(m1 = mod1, m2 = mod2)
ictab(mods, LOO, reloo = TRUE, k_threshold = 0.6, cores = 2)

## End(Not run)
```

seqtest

Sequential testing with evidence ratios

Description

Computes sequential evidence ratios, either based on the AIC, BIC, WAIC, or LOOIC. Supported models currently include `lm`, `merMod`, or `brmsfit` models. When data involve repeated measures (and so multiple lines per subject), a column indicating the subject "id" should be provided to the `id` argument. If nothing is passed to the `id` argument, `seqtest` will suppose that there is only one observation (i.e., one line) per subject.

Usage

```
seqtest(ic = aic, mod1, mod2, nmin = 10, id = NULL, boundary = Inf,
        blind = FALSE, nsims = NULL)
```

Arguments

ic	Indicates whether to use the aic or the bic.
mod1	A model of class lm or lmerMod.
mod2	A model of class lm or lmerMod (of the same class of mod1).
nmin	Minimum sample size from which start to compute sequential evidence ratios.
id	If applicable (i.e., repeated measures), name of the "id" column of your dataframe, in character string.
boundary	The Evidence Ratio (or its reciprocal) at which the run is stopped as well
blind	If true, the function only returns a "continue or stop" message
nsims	Number of permutation samples to evaluate (is ignored if blind = TRUE)

Author(s)

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See Also

[ictab](#)

Examples

```
## Not run:
# A first simple example
data(mtcars)
mod1 <- lm(mpg ~ cyl, mtcars)
mod2 <- lm(mpg ~ cyl + disp, mtcars)
seqtest(ic = aic, mod1, mod2, nmin = 10)

# Plotting the results
seqtest(ic = aic, mod1, mod2, nmin = 10) %>% plot

# Example with 10 permutation samples
seqtest(ic = aic, mod1, mod2, nmin = 10, nsims = 10)

# Example with blinding
seqtest(ic = aic, mod1, mod2, nmin = 10, boundary = 10, blind = TRUE)

# Example with repeated measures
library(lme4)
data(sleepstudy)
mod1 <- lmer(Reaction ~ Days + (1|Subject), sleepstudy)
mod2 <- lmer(Reaction ~ Days + I(Days^2) + (1|Subject), sleepstudy)
seqtest(ic = aic, mod1, mod2, nmin = 10, id = "Subject", nsims = 10)

# Example with brmsfit models
library(brms)
mod1 <- brm(Reaction ~ Days + (1|Subject), sleepstudy)
mod2 <- brm(Reaction ~ Days + I(Days^2) + (1|Subject), sleepstudy)
seqtest(ic = WAIC, mod1, mod2, nmin = 10, id = "Subject")
```

seqtest

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End(Not run)

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