Package 'MuMIn'

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Description Model selection and model averaging based on information criteria (AICc and alikes).

License GPL

Suggests lme4

LazyLoad yes

R topics documented:

MuMIn-package *Multi-model inference*

Description

The package MuMIn contains functions for (automated) model selection and model averaging based on information criteria (AIC alike).

Details

User level functions include:

model.avg - does model averaging.

get.models - evaluates models from the table returned by dredge.

dredge - runs models with combinations of terms of the supplied 'global' model.

AICc - calculates second-order Akaike information criterion for one or several fitted model objects.

Author(s)

Kamil Bartoń <kamil.barton@go2.pl>

References

Burnham, K. P. and Anderson, D. R (2002) *Model selection and multimodel inference: a practical information-theoretic approach*. 2nd ed.

See Also

AIC, step

Examples

```
fm1 <- lm(Fertility ~ . , data = swiss)

dd <- dredge(fm1)
dd
#list of models with delta.aicc < 4
top.models.1 <- get.models(dd, subset = delta < 4)
model.avg(top.models.1) # get averaged coefficients

#or as a 95% confidence set:
top.models.2 <- get.models(dd, cumsum(weight) <= .95)

model.avg(top.models.2) # get averaged coefficients

# Mixed models:

data(Orthodont, package="nlme")

require(nlme)
fm2 <- lme(distance ~ age + Sex, data = Orthodont, random = ~ 1 | Subject,
method="ML")
```

```
dredge(fm2)


require(lme4)
fm3 <- lmer(distance ~ age + Sex + (1 | Subject), data = Orthodont, REML=FALSE)
dd3 <- dredge(fm3)

# Get top-most models, but fitted by REML:
(top.models.3 <- get.models(dd3, subset = delta < 4, REML=TRUE))
# use: method = "REML" for older versions of lme4
```

`AICc` *Second-order Akaike Information Criterion*

Description

Calculates second-order Akaike information criterion for one or several fitted model objects (AIC for small samples).

Usage

```
AICc(object, ..., k = 2)
```

Arguments

`object`	a fitted model object
`...`	optionally more fitted model objects
`k`	the "penalty" per parameter to be used; the default `k` = 2 is the classical `AIC`

Value

If just one object is provided, returns a numeric value with the corresponding AICc; if more than one object are provided, returns a `data.frame` with rows corresponding to the objects and columns representing the number of parameters in the model (df), AICc and the `AIC`.

Author(s)

Kamil Bartoń

References

Burnham, K. P. and Anderson, D. R (2002) *Model selection and multimodel inference: a practical information-theoretic approach*. 2nd ed.

See Also

Akaike's An Information Criterion: `AIC`

Cement *Cement hardening data*

Description

Cement hardening data from Woods et al (1939).

Usage

```
data(Cement)
```

Format

`Cement` is a data frame with 5 variables. x1-x4 are four predictor variables expressed as a percentage of weight.

X1 calcium aluminate

X2 tricalcium silicate

X3 tetracalcium alumino ferrite

X4 dicalcium silicate

y calories of heat evolved per gram of cement after 180 days of hardening

Author(s)

Kamil Bartoń

Source

Woods H., Steinour H.H., Starke H.R. (1932) Effect of composition of Portland cement on heat evolved during hardening. *Industrial & Engineering Chemistry* 24, 1207-1214

References

Burnham, K. P. and Anderson, D. R (2002) *Model selection and multimodel inference: a practical information-theoretic approach*. 2nd ed.

dredge *Evaluate "all possible" models*

Description

runs models with all possible combinations of the explanatory variables in the supplied model.

Usage

```
 dredge(global.model, beta = FALSE, eval = TRUE, rank = "AICc",
fixed = NULL, m.max = NA, ...)
```

Arguments

global.model	a fitted 'global' model object. Currently, it can be a lm, glm, gam, lme, lmer, sarlm or spautolm, but also other types are likely to work (untested).
beta	logical should standardized coefficients be returned rather than normal ones?
eval	whether to evaluate and rank the models. If FALSE, a list of all possible model formulas is returned
rank	optional custom rank function (information criterion) to be used instead AICc, e.g. QAIC or BIC, See 'Details'
fixed	optional, either a single sided formula or a character vector giving names of terms to be included in all models
m.max	optional, maximum number of terms to be included in single model, defaults to the number of terms in global.model
...	optional arguments for the rank function

Details

Models are run one by one by calling update with modified formula argument. This method, while robust in that it can be applied to a variety of different models, is not very efficient, so may be time (and memory) consuming.

Handling interactions, dredge respects marginality constraints, so "all possible combinations" do not include models containing interactions without their respective main effects.

rank is found by a call to match.fun and typically is specified as a function or a symbol (e.g. a backquoted name) or a character string specifying a function to be searched for from the environment of the call to lapply.

Function rank must be able to accept model as a first argument and must always return a scalar.

Value

dredge returns a data.frame of class model.selection with models' coefficients, k, deviance/RSS, r-squared, AIC, AICc, etc. This depends on a type of model. Models are ordered according to AICc (lowest on top), or by rank function if specified. It has also a formulas attribute - a list containing all the model formulas.

Note

Use the lmer from **lme4** library rather than the old lmer from **Matrix** package. Make sure there is no a na.action set to na.omit in global.model. This can result with models fitted to different data sets, if there are NA's present. dredge cannot handle nested model designs (formulas such as y ~ a/b) properly.

Author(s)

Kamil Bartoń

References

Burnham, K. P. and Anderson, D. R (2002) *Model selection and multimodel inference: a practical information-theoretic approach*. 2nd ed.

See Also

get.models, model.avg. QAIC has examples of using custom rank function.

Examples

```
# Example from Burnham and Anderson (2002), page 100:
data(Cement)
lm1 <- lm(y ~ ., data = Cement)
dd <- dredge(lm1)
dd

#models with delta.aicc < 4
model.avg(get.models(dd, subset = delta < 4)) # get averaged coefficients

#or as a 95% confidence set:
top.models <- get.models(dd, cumsum(weight) <= .95)

model.avg(top.models) # get averaged coefficients

#topmost model:
top.models[[1]]
```

---

get.models *Get models*

Description

Gets list of models from a model.selection object

Usage

```
get.models(dd, subset = delta <= 4, ...)
```

Arguments

dd	object returned by dredge
subset	subset of models
...	additional parameters passed to update, for example, in lme/lmer one may want to use method = "REML" while using "ML" for model selection

Value

list of models.

Author(s)

Kamil Bartoń

See Also

dredge, model.avg

Examples

```
# Mixed models:

require(nlme)
fm2 <- lme(distance ~ age + Sex, data = Orthodont,
random = ~ 1 | Subject, method="ML")
dd2 <- dredge(fm2)

# Get top-most models, but fitted by REML:
(top.models.2 <- get.models(dd2, subset = delta < 4, method = "REML"))
```

---

miscellaneous *Helper functions*

Description

`beta.weights` - computes standardized coefficients (beta weights) for a model;

`coeffs` - extracts model coefficients;

`getAllTerms` - extracts independent variable names from a model object;

`tTable` - extracts a table of coefficients, standard errors, and p-values from a model object;

`Weights` - calculates Akaike weights (normalized relative likelihoods)

Usage

```
beta.weights(model)
coeffs(model)
getAllTerms(x, ...)
tTable(model, ...)
Weights(aic, ...)

cbindDataFrameList(x)
rbindDataFrameList(x)
```

Arguments

`model`	a fitted model object
`x`	a fitted model object or a `formula`. for `*bindDataFrameList`, a list of `data.frames`
`...`	other arguments, not used
`aic`	a vector of AIC (or other information criterion) values

Note

`coeffs`'s value is in most cases identical to that returned by `coef`, the only difference is that it returns fixed effects' coefficients for mixed models.

Functions `*bindDataFrameList` are not exported from the name space, use e.g. `MuMIn:::cbindDataFrameLi` to access them.

Author(s)

Kamil Bartoń

See Also

dredge

`model.avg` *Model averaging*

Description

Model averaging based on an information criterion.

Usage

```
model.avg(m1, ..., beta = FALSE, method = c("0", "NA"), rank = NULL,
rank.args = NULL, alpha = 0.05)

## S3 method for class 'averaging':
coef(object, ...)

## S3 method for class 'averaging':
predict(object, newdata, se.fit = NULL, interval = NULL,
type=NULL, ...)
```

Arguments

`m1`	A fitted model object or a list of such objects.
`beta`	Logical, should standardized coefficients be returned rather than normal ones?
`method`	If set to "0" (default), terms missing in one model are assumed to be 0's, otherwise they are omitted from the weighted average.
`rank`	Custom rank function (information criterion) to use instead of `AICc`, e.g. `QAIC` or `BIC`, may be omitted if `m1` is a list returned by `dredge`. See 'Details'.
`rank.args`	Optional `list` of arguments for the `rank` function.
`alpha`	Significance level for calculatinq confidence intervals.
`object`	An object returned by `model.avg`.
`newdata`	An optional data frame in which to look for variables with which to predict. If omitted, the fitted values are used.
`se.fit,interval`	Currently not used.
`type`	Ignored. Only predictions on the link scale are allowed. Warning is given if user tries something else here.
`...`	for `model.avg` - more fitted model objects, for `predict` - arguments to be passed to respective `predict` method

Details

rank is found by a call to match.fun and typically is specified as a function or a symbol (e.g. a backquoted name) or a character string specifying a function to be searched for from the environment of the call to lapply.
Function rank must be able to accept model as a first argument and must always return a scalar. \
Apart from predict and coef, other default methods, such as formula and residuals may be used.

Value

model.avg returns a list with elements:

summary	Model table with deviance, AICc, Delta and weight.
coefficients	the model coefficients
variance	variance of coefficients
avg.model	averaged model summary (data.frame with columns: coef - averaged coefficients, var - unconditional variance estimator, ase - adjusted standard error estimator, lci, uci - unconditional confidence intervals)
relative.importance	relative variable importances
variable.codes	Variable names with numerical codes used in the summary
relative.importance	Relative importance of variables
weights	
beta	(logical) were standardized coefficients used?
model	the model matrix, analogical to one that would be used in a single model.
residuals	the residuals (response minus fitted values).

Author(s)

Kamil Bartoń

References

Burnham, K. P. and Anderson, D. R (2002) *Model selection and multimodel inference: a practical information-theoretic approach*. 2nd ed.

See Also

dredge, get.models. QAIC has examples of using custom rank function.

Examples

```
# Example from Burnham and Anderson (2002), page 100:
data(Cement)
lm1 <- lm(y ~ ., data = Cement)
dd <- dredge(lm1)
dd

#models with delta.aicc < 4
```

```
model.avg(get.models(dd, subset = delta < 4)) # get averaged coefficients

#or as a 95% confidence set:
top.models <- get.models(dd, cumsum(weight) <= .95)

model.avg(top.models) # get averaged coefficients

#topmost model:
top.models[[1]]

## Not run:
# using BIC (Schwarz's Bayesian criterion) to rank the models
BIC <- function(x) AIC(x, k = log(length(residuals(x))))
mav <- model.avg(top.models, rank=BIC)

## End(Not run)


# Predicted values
nseq <- function(x, len=length(x)) seq(min(x, na.rm=TRUE), max(x, na.rm=TRUE),
length=len)

# New predictors: X1 along the range of original data, other variables held
# constant at their means
newdata <- as.data.frame(lapply(lapply(Cement[1:5], mean), rep, 25))
newdata$X1 <- nseq(Cement$X1, 25)

# Predictions from each of the models in a set:
pred <- sapply(top.models, predict, newdata=newdata)
# Add predictions from the averaged model:
pred <- cbind(pred, averaged=predict(model.avg(top.models), newdata))

matplot(x = newdata$X1, y = pred, type="l", lwd=c(rep(1,ncol(pred)-1), 2))

legend("topleft", legend=c(lapply(top.models, formula), "Averaged model"),
   col=seq(ncol(pred)), lty=1:5, lwd=c(rep(1,ncol(pred)-1), 2))
```

`par.avg` *Parameter averaging*

Description

Averages single parameter based on provided weights

Usage

```
par.avg(x, se, npar, weight, alpha = 0.05)
```

Arguments

`x`	vector of parameters
`se`	vector of standard errors
`npar`	vector giving numbers of estimated parameters

`weight`	vector of weights
`alpha`	significance level for calculatinq confidence intervals

Value

par.avg returns a vector with named elements:

`Coefficient`	model coefficients
`Variance`	unconditional variance of coefficients
`Unconditional SE`	
`Lower CI, Upper CI`	relative variable importances

Author(s)

Kamil Bartoń

References

Burnham, K. P. and Anderson, D. R (2002) *Model selection and multimodel inference: a practical information-theoretic approach*. 2nd ed.

See Also

`model.avg` for averaging models.

`QAIC` *Quasi AIC*

Description

Calculates "quasi AIC" for one or several fitted model objects. This function is provided just as an example of custom rank function for use with `model.avg` and `dredge`

Usage

```
QAIC(object, ..., chat)
```

Arguments

`object`	a fitted model object.
`...`	optionally more fitted model objects.
`chat`	c - hat

Details

`rank` is specified as a function or a symbol (e.g. a backquoted name) or a character string specifying a function.

Function `rank` must be able to accept model as a first argument and must always return a scalar.

Value

If just one object is provided, returns a numeric value with the corresponding QAIC; if more than one object are provided, returns a data.frame with rows corresponding to the objects.

Author(s)

Kamil Bartoń

Examples

```
budworm <- data.frame(ldose = rep(0:5, 2), numdead = c(1, 4, 9, 13, 18, 20, 0,
2, 6, 10, 12, 16), sex = factor(rep(c("M", "F"), c(6, 6))))

budworm$SF <- cbind(budworm$numdead, 20 - budworm$numdead)

budworm.qlg <- glm(SF ~ sex*ldose, family = quasibinomial, data = budworm)

dd1 <- dredge(budworm.qlg, rank = "QAIC",
chat = summary(budworm.qlg)$dispersion)
gm1 <- get.models(dd1, 1:4)

model.avg(gm1)

model.avg(gm1[[1]], gm1[[2]], rank = "QAIC", rank.args = list(chat = 1))
```

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