Package 'actuaRE'

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```
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Description Using this package, you can fit a random effects model using either the hierarchical credi-
     bility model, a combination of the hierarchical credibility model with a generalized lin-
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Description

Using this package, you can fit a random effects model using either the hierarchical credibility model, a combination of the hierarchical credibility model with a generalized linear model or a Tweedie generalized linear mixed model. See Campo, B.D.C. and Antonio, K. (2023) < doi:10.1080/03461238.2022.2161413

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References

Campo, B.D.C. and Antonio, Katrien (2023). Insurance pricing with hierarchically structured data an illustration with a workers' compensation insurance portfolio. *Scandinavian Actuarial Journal*, doi: 10.1080/03461238.2022.2161413

Dannenburg, D. R., Kaas, R. and Goovaerts, M. J. (1996). *Practical actuarial credibility models*. Amsterdam: IAE (Institute of Actuarial Science and Econometrics of the University of Amsterdam).

Jewell, W. S. (1975). The use of collateral data in credibility theory: a hierarchical model. Laxenburg: IIASA.

Ohlsson, E. (2005). Simplified estimation of structure parameters in hierarchical credibility. *Presented at the Zurich ASTIN Colloquium*.

Ohlsson, E. (2008). Combining generalized linear models and credibility models in practice. *Scandinavian Actuarial Journal* **2008**(4), 301–314.

See Also

hierCredibility hierCredGLM hierCredTweedie tweedieGLMM BalanceProperty

Examples

```
library(actuaRE)
# Vignette of the package
vignette(package = "actuaRE")

# Load data
data(hachemeisterLong)
data(dataCar)

# Hierarchical credibility model of Jewell
fit = hierCredibility(ratio, weight, cohort, state, hachemeisterLong)

# Combination of the hierarchical credibility model with a GLM (Ohlsson, 2008)
fit = hierCredGLM(Y ~ area + (1 | VehicleType / VehicleBody), dataCar, weights = w,
p = 1.7)
```

.addREs

Add random effects to the data frame

Description

Internal function

Usage

```
.addREs(obj, newdata)
```

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Arguments

obj	object with model fit

newdata an object coercible to data. table.

adjustIntercept Adjust the intercept to regain the balance property

Description

This function updates the intercept term of the model fit such that the balance property is satisfied.

Usage

```
adjustIntercept(obj, data)
```

Arguments

```
obj an object of type glm, cpglm or cpglmm containing the model fit.

data a data. frame or data. table object that was used to fit the model.
```

Value

The object with the adjusted (fixed effects) coefficients.

References

Campo, B.D.C. and Antonio, Katrien (2023). Insurance pricing with hierarchically structured data an illustration with a workers' compensation insurance portfolio. *Scandinavian Actuarial Journal*, doi: 10.1080/03461238.2022.2161413

Wüthrich, M. V. (2020). Bias regularization in neural network models for general insurance pricing. *European actuarial journal* **10**(1), 179–202.

Examples

```
library(statmod)
datas = dataCar[1:1e3, ]
Fit = glm(Y ~ area + gender, data = datas, weights = datas$w, family = tweedie(1.75, 0),
model = TRUE, control = glm.control(epsilon = 1e-4, maxit = 5e2))
w = weights(Fit, "prior")
y = Fit$y
sum(w * y) == sum(w * fitted(Fit))
adjFit = adjustIntercept(Fit, datas)
coef(adjFit)
sum(w * y) == sum(w * fitted(adjFit))
```

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BalanceProperty

Balance property

Description

Function to assess whether the balance property holds

Usage

```
BalanceProperty(obj)
```

Arguments

obj

an object containing the model fit

Value

a list with the slots call (the original call), BalanceProperty (logical indicating whether the balance property is satisfied) and Alpha (Ratio total observed damage to total predicted damage).

References

Campo, B.D.C. and Antonio, Katrien (2023). Insurance pricing with hierarchically structured data an illustration with a workers' compensation insurance portfolio. *Scandinavian Actuarial Journal*, doi: 10.1080/03461238.2022.2161413

Wüthrich, M. V. (2020). Bias regularization in neural network models for general insurance pricing. *European actuarial journal* **10**(1), 179–202.

Examples

```
fit = hierCredGLM(Y \sim area + (1 | VehicleType / VehicleBody), dataCar, weights = w, p = 1.75, epsilon = 1e-6) BalanceProperty(fit)
```

dataCar

data Car

Description

This data set is taken from the dataCar data set of the insuranceData package and slightly adjusted (see the code in examples for reproducing this data set). The original data set is based on one-year vehicle insurance policies taken out in 2004 or 2005. There are 67566 policies, of which 4589 (6.8%) had at least one claim.

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Usage

```
data(dataCar)
```

Format

```
A data frame with 67566 observations on the following 15 variables.
```

```
veh_value vehicle value, in $10,000s
exposure 0-1
clm occurrence of claim (0 = no, 1 = yes)
numclaims number of claims
claimcst0 claim amount (0 if no claim)
veh_body vehicle body, coded as BUS CONVT COUPE HBACK HDTOP MCARA MIBUS PANVN RDSTR SEDAN
     STNWG TRUCK UTE
veh_age 1 (youngest), 2, 3, 4
gender a factor with levels F M
area a factor with levels ABCDEF
agecat 1 (youngest), 2, 3, 4, 5, 6
X_OBSTAT_ a factor with levels 01101 0 0 0
Y the loss ratio, defined as the number of claims divided by the exposure
w the exposure, identical to exposure
VehicleType type of vehicle, common vehicle or uncommon vehicle
VehicleBody vehicle body, identical to veh_body
```

Details

Adjusted data set dataCar, where we removed claims with a loss ratio larger than 1 000 000. In addition, we summed the exposure per vehicle body and removed those where the summed exposure was less than 100. Hereby, we ensure that there is sufficient exposure for each vehicle body category.

Source

http://www.acst.mq.edu.au/GLMsforInsuranceData

References

De Jong P., Heller G.Z. (2008), Generalized linear models for insurance data, Cambridge University Press

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Examples

```
# How to construct the data set using the original dataCar data set from the insuranceData package
library(plyr)
library(magrittr)
data("dataCar", package = "insuranceData")
dataCar$Y = with(dataCar, claimcst0 / exposure)
dataCar$w = dataCar$exposure
         = dataCar[which(dataCar$Y < 1e6), ]</pre>
Yw = ddply(dataCar, .(veh_body), function(x) c(crossprod(x$Y, x$w) / sum(x$w), sum(x$w)))
dataCar = dataCar[!dataCar$veh_body %in% Yw[Yw$V2 < 1e2, "veh_body"], ]</pre>
dataCar$veh_body %<>% droplevels()
dataCar$VehicleType = sapply(tolower(dataCar$veh_body), function(x) {
  if(x %in% c("sedan", "ute", "hback"))
    "Common vehicle"
  else
    "Uncommon vehicle"
})
dataCar$VehicleBody = dataCar$veh_body
```

findbars

Determine random-effects expressions from a formula

Description

From the right hand side of a formula for a mixed-effects model, determine the pairs of expressions that are separated by the vertical bar operator. Also expand the slash operator in grouping factor expressions and expand terms with the double vertical bar operator into separate, independent random effect terms.

Usage

```
findbars(term)
```

Arguments

term

a mixed-model formula

Value

pairs of expressions that were separated by vertical bars

Note

This function is called recursively on individual terms in the model, which is why the argument is called term and not a name like form, indicating a formula.

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See Also

```
formula, model.frame, model.matrix.

Other utilities: mkRespMod, mkReTrms, nlformula, nobars, subbars
```

Examples

```
findbars(f1 <- Reaction ~ Days + (Days | Subject))
## => list( Days | Subject )
## These two are equivalent:% tests in ../inst/tests/test-doubleVertNotation.R
findbars(y ~ Days + (1 | Subject) + (0 + Days | Subject))
findbars(y ~ Days + (Days || Subject))
## => list of length 2: list (1 | Subject , 0 + Days | Subject)
findbars(~ 1 + (1 | batch / cask))
## => list of length 2: list (1 | cask:batch , 1 | batch)
```

fixef

Extract fixed-effects estimates

Description

Extract the fixed-effects estimates

Arguments

object

any fitted model object from which fixed effects estimates can be extracted.

Details

Extract the estimates of the fixed-effects parameters from a fitted model.

Value

a named, numeric vector of fixed-effects estimates.

Examples

fixef-actuaRE 9

fixef-actuaRE

Extract the fixed-effects estimates from a fitted random effects model

Description

A generic function to extract the fixed effects (i.e. the company-specific effects) estimates from a fitted random effects model.

Usage

```
## S3 method for class 'hierCredGLM'
fixef(object, ...)
## S3 method for class 'hierCredTweedie'
fixef(object, ...)
```

Arguments

```
object an object of type hierCredGLM or hierCredTweedie ... ignored.
```

Value

a named, numeric vector of fixed-effects estimates.

Examples

```
fit = hierCredGLM(Y \sim area + (1 | VehicleType / VehicleBody), dataCar, weights = w, p = 1.75, epsilon = 1e-6) fixef(fit)
```

hachemeisterLong

Hachemeister Data Set

Description

Long format of the Hachemeister (1975) data set giving average claim amounts in private passenger bodily injury insurance. We have data of five U.S. states over 12 quarters between July 1970 and June 1973 and we have the corresponding number of claims. To obtain a hierarchical structure, we created an artificial variable cohort. With this, we created a hierarchical multi-level factor, with cohort as the first hierarchical level and state as the second hierarchical level, nested within cohort.

Usage

hachemeisterLong

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Format

```
A data.frame with 60 rows and the following 5 columns: cohort artificially created variable; state the state number; time time variable (quarter of the observation); ratio the average claim amount; weight the corresponding number of claims.
```

Source

Hachemeister, C. A. (1975), *Credibility for regression models with application to trend*, Proceedings of the Berkeley Actuarial Research Conference on Credibility, Academic Press.

hierCredGLM

Combining the hierarchical credibility model with a GLM (Ohlsson, 2008)

Description

Fit a random effects model using Ohlsson's methodology. In this function you explicitly specify the power parameter p. See hierCredTweedie when you also want to estimate the p.

Usage

```
hierCredGLM(
formula,
data,
weights,
p = 1.5,
link.power = 0,
muHatGLM = TRUE,
epsilon = 1e-04,
maxiter = 500,
maxiterGLM = 500,
verbose = FALSE,
returnData = TRUE,
balanceProperty = TRUE,
y = TRUE,
...
)
```

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Arguments

formula object of type formula that specifies which model should be fitted. Syntax is the

same as for lmer and glmer. For example, Yijkt ~ x1 + x2 + (1 | Industry /

Branch).

data an object that is coercible by as.data.table, containing the variables in the

model.

weights variable name of the exposure weight.

p the value for the power parameter of the Tweedie distribution, which is passed

to tweedie. Default is 1.5.

link.power index of power link function, which is passed to tweedie. link.power = 0

produces a log-link. Defaults to the canonical link, which is 1 - p.

muHatGLM indicates which estimate has to be used in the algorithm for the intercept term.

Default is TRUE, which used the intercept as estimated by the GLM. If FALSE,

the estimate of the hierarchical credibility model is used.

epsilon positive convergence tolerance ϵ ; the iterations converge when $7 ||\theta[k] - \theta[k-1]||$

1|||²[[2]]/|| $\theta[k-1]$ ||²[[2]] $< \epsilon$. Here, $\theta[k]$ is the parameter vector at the k^{th}

iteration.

maxiter maximum number of iterations.

maxiterGLM maximum number of iterations when fitting the GLM part. Passed to glm.

verbose logical indicating if output should be produced during the algorithm.

returnData logical indicating if input data has to be returned.

balanceProperty

logical indicating if the balance property should be satisfied.

y logical indicating whether the response vector should be returned as a compo-

nent of the returned value.

... arguments passed to glm

Value

An object of type hierCredGLM with the following slots:

call the matched call

HierarchicalResults

results of the hierarchical credibility model.

fitGLM the results from fitting the GLM part.

iter total number of iterations.

Converged logical indicating whether the algorithm converged.

LevelsCov object that summarizes the unique levels of each of the contract-specific covari-

ates.

fitted.values the fitted mean values, resulting from the model fit.

prior.weights the weights (exposure) initially supplied.

y if requested, the response vector. Default is TRUE.

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References

Campo, B.D.C. and Antonio, Katrien (2023). Insurance pricing with hierarchically structured data an illustration with a workers' compensation insurance portfolio. *Scandinavian Actuarial Journal*, doi: 10.1080/03461238.2022.2161413

Ohlsson, E. (2008). Combining generalized linear models and credibility models in practice. *Scandinavian Actuarial Journal* **2008**(4), 301–314.

See Also

hierCredGLM-class, fitted.hierCredGLM, predict.hierCredGLM, ranef-actuaRE, weights-actuaRE, hierCredibility, hierCredTweedie, plotRE, adjustIntercept, BalanceProperty

Examples

```
data("dataCar")
fit = hierCredGLM(Y ~ area + (1 | VehicleType / VehicleBody), dataCar, weights = w,
p = 1.7)
fit
summary(fit)
ranef(fit)
fixef(fit)
```

hierCredGLM-class

Class "hierCredGLM" of fitted random effects models estimated with Ohlsson's GLMC algorithm

Description

Class "hierCredGLM" of fitted random effects models estimated with Ohlsson's GLMC algorithm

Usage

```
## S3 method for class 'hierCredGLM'
print(x, ...)
## S3 method for class 'hierCredGLM'
summary(object, ...)
## S3 method for class 'hierCredGLM'
fitted(object, ...)
```

Arguments

```
x an object of class hierCredGLM
... currently ignored.
object an object of class hierCredGLM
```

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Value

The function hierCredGLM returns an object of class hierCredGLM, which has the following slots:

call the matched call

HierarchicalResults

results of the hierarchical credibility model.

fitGLM the results from fitting the GLM part.

iter total number of iterations.

Converged logical indicating whether the algorithm converged.

LevelsCov object that summarizes the unique levels of each of the contract-specific covari-

ates.

fitted.values the fitted mean values, resulting from the model fit.

prior.weights the weights (exposure) initially supplied.

y if requested, the response vector. Default is TRUE.

S3 methods

print: Prints the call, the estimated variance parameters, the unique number of categories of the hierarchical MLF and the output of the GLM part. The ... argument is currently ignored. Returns an invisible copy of the original object.

summary: In addition to the output of the print.hierCredGLM function, the summary function also prints the random effect estimates and a summary of the GLM (see summary.glm). Returns an invisible copy of the original object.

fitted: Returns the fitted values.

See Also

hierCredGLM

hierCredibility Hierarchical credibility model of Jewell

Description

Fit a random effects model, without contract-specific risk factors, using the hierarchical credibility model of Jewell.

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Usage

```
hierCredibility(
   Yijkt,
   wijkt,
   sector,
   group,
   data,
   muHat = NULL,
   type = c("additive", "multiplicative"),
   returnData = FALSE
)
```

Arguments

Yijkt variable name of the response variable (the loss cost within actuarial applica-

tions).

wijkt variable name of the exposure weight.

sector variable name of the first hierarchical level.

group variable name of the second hierarchical level that is nested within the first hier-

archical level.

data an object that is coercible by as.data.table, containing the variables in the

model.

muHat estimate for the intercept term. Default is NULL and in this case, the estimator as

given in Ohlsson (2005) is used.

type specifies whether the additive (Dannenburg, 1996) or multiplicative (Ohlsson,

2005) formulation of the hierarchical credibility model is used. Default is addi-

tive.

returnData Logical, indicates whether the data object has to be returned. Default is FALSE.

Value

An object of type hierCredibility with the following slots:

call the matched call

type Whether additive or multiplicative hierarchical credibility model is used.

Variances The estimated variance components. s2 is the estimated variance of the indi-

vidual contracts, tausq the estimate of Var(V[j]) and nusq is the estimate of

Var(V[jk]).

Means The estimated averages at the portfolio level (intercept term μ), at the first hi-

erarchical level $(bar(Y)[\%.\%j\%.\%\%.\%]^z)$ and at the second hierarchical level

(bar(Y)[%.%jk%.%]).

Weights The weights at the first hierarchical level z[j%.%] and at the second hierarchical

level w[%.%jk%.%].

Credibility The credibility weights at the first hierarchical level q[j%.%] and at the second

hierarchical level z[jk].

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Premiums	The overall expectation $widehat(\mu)$, sector expectation $widehat(V)[j]$ and group expectation $widehat(V)[jk]$.
Relativity	The estimated random effects $widehat(U)[j]$ and $widehat(U)[jk]$ of the sector and group, respectively.
RawResults	Objects of type data.table with all intermediate results.
fitted.values	the fitted mean values, resulting from the model fit.

References

Campo, B.D.C. and Antonio, Katrien (2023). Insurance pricing with hierarchically structured data an illustration with a workers' compensation insurance portfolio. *Scandinavian Actuarial Journal*, doi: 10.1080/03461238.2022.2161413

Dannenburg, D. R., Kaas, R. and Goovaerts, M. J. (1996). *Practical actuarial credibility models*. Amsterdam: IAE (Institute of Actuarial Science and Econometrics of the University of Amsterdam).

Jewell, W. S. (1975). The use of collateral data in credibility theory: a hierarchical model. Laxenburg: IIASA.

Ohlsson, E. (2005). Simplified estimation of structure parameters in hierarchical credibility. *Presented at the Zurich ASTIN Colloquium*.

See Also

hierCredibility-class, fitted.hierCredibility, predict.hierCredibility, ranef-actuaRE, weights-actuaRE, hierCredTweedie, hierCredGLM, cpglm, plotRE

Examples

```
library(actuar)
library(actuaRE)
data("hachemeister", package = "actuar")
Df = as.data.frame(hachemeister)
X = as.data.frame(cbind(cohort = c(1, 2, 1, 2, 2), hachemeister))
Df = reshape(X, idvar = "state", varying = list(paste0("ratio.", 1:12),
    paste0("weight.", 1:12)), direction = "long")
fitActuar = cm(~ cohort + cohort:state, data = X, ratios = ratio.1:ratio.12,
    weights = weight.1:weight.12, method = "Ohlsson")
fitActuaRE = hierCredibility(ratio.1, weight.1, cohort, state, Df)
summary(fitActuar)
summary(fitActuaRE)
```

hierCredibility-class Class "hierCredibility" of fitted hierarchical credibility models

Description

Class "hierCredibility" of fitted hierarchical credibility models

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Usage

```
## S3 method for class 'hierCredibility'
print(x, ...)
## S3 method for class 'hierCredibility'
summary(object, ...)
## S3 method for class 'hierCredibility'
fitted(object, ...)
```

Arguments

x an object of class hierCredibility

... currently ignored.

object an object of class hierCredibility

Value

The function hierCredibility returns an object of class hierCredibility, which has the following slots:

call the matched call

type Whether additive or multiplicative hierarchical credibility model is used.

Variances The estimated variance components. s2 is the estimated variance of the indi-

vidual contracts, tausq the estimate of Var(V[j]) and nusq is the estimate of

Var(V[jk]).

Means The estimated averages at the portfolio level (intercept term μ), at the first hi-

erarchical level $(bar(Y)[\%.\%j\%.\%\%.\%]^z)$ and at the second hierarchical level

(bar(Y)[%.%jk%.%]).

Weights The weights at the first hierarchical level z[j%.%] and at the second hierarchical

level w[%.%jk%.%].

Credibility The credibility weights at the first hierarchical level q[j%.%] and at the second

hierarchical level z[jk].

Premiums The overall expectation $widehat(\mu)$, sector expectation widehat(V)[j] and group

expectation widehat(V)[jk].

Relativity The estimated random effects widehat(U)[j] and widehat(U)[jk] of the sector

and group, respectively.

RawResults Objects of type data. table with all intermediate results.

fitted.values the fitted mean values, resulting from the model fit.

S3 methods

print: Prints the call, the estimated variance parameters and the unique number of categories of the hierarchical MLF. The . . . argument is currently ignored. Returns an invisible copy of the original object.

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summary: In addition to the output of the print.hierCredibility function, the summary function prints the random effect estimates as well. Returns an invisible copy of the original object.

fitted: Returns the fitted values.

See Also

hierCredibility

hierCredTweedie Combining the hierarchical credibility model with a GLM (Ohlsson, 2008)

Description

Fit a random effects model using Ohlsson's methodology. In this function you estimate the power parameter p. See hierCredGLM when you want fix p.

Usage

```
hierCredTweedie(
  formula,
  data,
  weights,
  muHatGLM = TRUE,
  epsilon = 1e-04,
  maxiter = 500,
  verbose = FALSE,
  returnData = TRUE,
  cpglmControl = list(bound.p = c(1.01, 1.99)),
  balanceProperty = TRUE,
  optimizer = "bobyqa",
  y = TRUE,
  ...
)
```

Arguments

formula	object of type formula that specifies which model should be fitted. Syntax is the same as for lmer and glmer. For example, $Yijkt \sim x1 + x2 + (1 \mid Industry / Branch)$.
data	an object that is coercible by as.data.table, containing the variables in the model.
weights	variable name of the exposure weight.
muHatGLM	indicates which estimate has to be used in the algorithm for the intercept term. Default is TRUE, which used the intercept as estimated by the GLM. If FALSE, the estimate of the hierarchical credibility model is used.

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epsilon positive convergence tolerance ϵ ; the iterations converge when $||\theta[k] - \theta[k -$

1|| $|^2$ [[2]]/|| $\theta[k-1]$ || $|^2$ [[2]] $< \epsilon$. Here, $\theta[k]$ is the parameter vector at the k^{th}

iteration.

maxiter maximum number of iterations.

verbose logical indicating if output should be produced during the algorithm.

returnData logical indicating if input data has to be returned.

cpglmControl a list of parameters to control the fitting process in the GLM part. By default,

cpglmControl = list(bound.p = c(1.01, 1.99)) which restricts the range of the power parameter p to [1.01, 1.99] in the fitting process. This list is passed to

cpglm.

balanceProperty

logical indicating if the balance property should be satisfied.

optimizer a character string that determines which optimization routine is to be used in es-

timating the index and the dispersion parameters. Possible choices are "nlminb" (the default, see nlminb), "bobyqa" (bobyqa) and "L-BFGS-B" (optim).

y logical indicating whether the response vector should be returned as a compo-

nent of the returned value.

... arguments passed to cpglm.

Details

When estimating the GLM part, this function uses the cpglm function from the cplm package.

Value

An object of type hierCredTweedie with the following slots:

call the matched call

HierarchicalResults

results of the hierarchical credibility model.

fitGLM the results from fitting the GLM part.

iter total number of iterations.

Converged logical indicating whether the algorithm converged.

LevelsCov object that summarizes the unique levels of each of the contract-specific covari-

ates.

fitted.values the fitted mean values, resulting from the model fit.

prior.weights the weights (exposure) initially supplied.

y if requested, the response vector. Default is TRUE.

References

Ohlsson, E. (2008). Combining generalized linear models and credibility models in practice. *Scandinavian Actuarial Journal* **2008**(4), 301–314.

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See Also

hierCredTweedie-class, fitted.hierCredTweedie, predict.hierCredTweedie, ranef-actuaRE, weights-actuaRE, hierCredibility, hierCredGLM, cpglm, plotRE, adjustIntercept, BalanceProperty @references Campo, B.D.C. and Antonio, Katrien (2023). Insurance pricing with hierarchically structured data an illustration with a workers' compensation insurance portfolio. *Scandinavian Actuarial Journal*, doi: 10.1080/03461238.2022.2161413

Examples

```
data("dataCar")
fit = hierCredTweedie(Y ~ area + (1 | VehicleType / VehicleBody), dataCar,
weights = w, epsilon = 1e-6)
fit
summary(fit)
ranef(fit)
fixef(fit)
```

hierCredTweedie-class Class "hierCredTweedie" of fitted random effects models estimated with Ohlsson's GLMC algorithm

Description

Class "hierCredTweedie" of fitted random effects models estimated with Ohlsson's GLMC algorithm

Usage

```
## S3 method for class 'hierCredTweedie'
print(x, ...)
## S3 method for class 'hierCredTweedie'
summary(object, ...)
## S3 method for class 'hierCredTweedie'
fitted(object, ...)
```

Arguments

```
x an object of class hierCredTweedie... currently ignored.object an object of class hierCredTweedie
```

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Value

The function hierCredGLM returns an object of class hierCredGLM, which has the following slots:

call the matched call

HierarchicalResults

results of the hierarchical credibility model.

fitGLM the results from fitting the GLM part.

iter total number of iterations.

Converged logical indicating whether the algorithm converged.

LevelsCov object that summarizes the unique levels of each of the contract-specific covari-

ates.

fitted.values the fitted mean values, resulting from the model fit.

prior.weights the weights (exposure) initially supplied.

y if requested, the response vector. Default is TRUE.

S3 methods

print: Prints the call, the estimated variance parameters, the unique number of categories of the hierarchical MLF and the output of the GLM part. The ... argument is currently ignored. Returns an invisible copy of the original object.

summary: In addition to the output of the print.hierCredTweedie function, the summary function also prints the random effect estimates and a summary of the GLM (see summary.glm). Returns an invisible copy of the original object.

fitted: Returns the fitted values.

See Also

hierCredTweedie

is.formula Formula

Description

Checks if the object is of the type formula

Usage

is.formula(x)

Arguments

x the object.

isNested 21

Value

logical indicating if the object is a formula or not.

Examples

```
f = formula(y ~ x)
is.formula(f)
```

isNested

Is f1 nested within f2?

Description

Does every level of f1 occur in conjunction with exactly one level of f2? The function is based on converting a triplet sparse matrix to a compressed column-oriented form in which the nesting can be quickly evaluated.

Usage

```
isNested(f1, f2)
```

Arguments

```
f1 factor 1
f2 factor 2
```

Value

TRUE if factor 1 is nested within factor 2

Examples

```
library(lme4)
with(Pastes, isNested(cask, batch)) ## => FALSE
with(Pastes, isNested(sample, batch)) ## => TRUE
```

22 modular

modular

Modular Functions for Mixed Model Fits

Description

Modular functions for mixed model fits

Usage

Arguments

subset

formula a two-sided linear formula object describing both the fixed-effects and random-

effects parts of the model, with the response on the left of a \sim operator and the terms, separated by + operators, on the right. Random-effects terms are distinguished by vertical bars ("|") separating expressions for design matrices

from grouping factors.

data an optional data frame containing the variables named in formula. By default

the variables are taken from the environment from which 1mer is called. While data is optional, the package authors *strongly* recommend its use, especially when later applying methods such as update and drop1 to the fitted model (*such methods are not guaranteed to work properly if* data *is omitted*). If data is omitted, variables will be taken from the environment of formula (if specified

as a formula) or from the parent frame (if specified as a character vector).

an optional expression indicating the subset of the rows of data that should be used in the fit. This can be a logical vector, or a numeric vector indicating which observation numbers are to be included, or a character vector of the row names

to be included. All observations are included by default.

weights an optional vector of 'prior weights' to be used in the fitting process. Should be

NULL or a numeric vector.

na.action a function that indicates what should happen when the data contain NAs. The de-

fault action (na.omit, inherited from the 'factory fresh' value of getOption("na.action"))

strips any observations with any missing values in any variables.

offset this can be used to specify an *a priori* known component to be included in the

linear predictor during fitting. This should be NULL or a numeric vector of length equal to the number of cases. One or more offset terms can be included in the formula instead or as well, and if more than one is specified their sum is used.

See model.offset.

contrasts an optional list. See the contrasts.arg of model.matrix.default.

control a list giving

for [g]lFormula: all options for running the model, see lmerControl;

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for mkLmerDevfun, mkGlmerDevfun: options for the inner optimization step; for optimizeLmer and optimizeGlmer: control parameters for nonlinear optimizer (typically inherited from the ... argument to lmerControl). start starting values (see lmer; for glFormula, should be just a numeric vector of fixed-effect coefficients) family a GLM family; see glm and family. optional starting values on the scale of the conditional mean; see glm for details. mustart etastart optional starting values on the scale of the unbounded predictor; see glm for details. other potential arguments; for optimizeLmer and optimizeGlmer, these are passed to internal function optwrap, which has relevant parameters calc. derivs and use.last.params (see lmerControl).

Details

These functions make up the internal components of an [gn]lmer fit.

- [g]lFormula takes the arguments that would normally be passed to [g]lmer, checking for errors and processing the formula and data input to create a list of objects required to fit a mixed model.
- mk(G1|L)merDevfun takes the output of the previous step (minus the formula component) and creates a deviance function
- optimize(G1|L)mer takes a deviance function and optimizes over theta (or over theta and beta, if stage is set to 2 for optimizeGlmer
- updateGlmerDevfun takes the first stage of a GLMM optimization (with nAGQ=0, optimizing over theta only) and produces a second-stage deviance function
- mkMerMod takes the *environment* of a deviance function, the results of an optimization, a list of random-effect terms, a model frame, and a model all and produces a [g]lmerMod object.

Value

1Formula and glFormula return a list containing components:

fr model frame

X fixed-effect design matrix

reTrms list containing information on random effects structure: result of mkReTrms

mkLmerDevfun and mkGlmerDevfun return a function to calculate deviance (or restricted deviance) as a function of the theta (random-effect) parameters. updateGlmerDevfun returns a function to calculate the deviance as a function of a concatenation of theta and beta (fixed-effect) parameters. These deviance functions have an environment containing objects required for their evaluation. CAUTION: The environment of functions returned by mk(Gl|L)merDevfun contains reference class objects (see ReferenceClasses, merPredD-class, lmResp-class), which behave in ways that may surprise many users. For example, if the output of mk(Gl|L)merDevfun is naively copied, then modifications to the original will also appear in the copy (and vice versa). To avoid this behavior one must make a deep copy (see ReferenceClasses for details).

optimizeLmer and optimizeGlmer return the results of an optimization.

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Examples

```
library(lme4)
### Fitting a linear mixed model in 4 modularized steps
## 1. Parse the data and formula:
lmod <- lFormula(Reaction ~ Days + (Days|Subject), sleepstudy)</pre>
names(lmod)
## 2. Create the deviance function to be optimized:
(devfun <- do.call(mkLmerDevfun, lmod))</pre>
ls(environment(devfun)) # the environment of 'devfun' contains objects
# required for its evaluation
## 3. Optimize the deviance function:
opt <- optimizeLmer(devfun)</pre>
opt[1:3]
## 4. Package up the results:
mkMerMod(environment(devfun), opt, lmod$reTrms, fr = lmod$fr)
### Same model in one line
lmer(Reaction ~ Days + (Days|Subject), sleepstudy)
### Fitting a generalized linear mixed model in six modularized steps
## 1. Parse the data and formula:
glmod <- glFormula(cbind(incidence, size - incidence) ~ period + (1 | herd),</pre>
                   data = cbpp, family = binomial)
#.... see what've got :
str(glmod, max=1, give.attr=FALSE)
## 2. Create the deviance function for optimizing over theta:
(devfun <- do.call(mkGlmerDevfun, glmod))</pre>
ls(environment(devfun)) # the environment of devfun contains lots of info
## 3. Optimize over theta using a rough approximation (i.e. nAGQ = 0):
(opt <- optimizeGlmer(devfun))</pre>
## 4. Update the deviance function for optimizing over theta and beta:
(devfun <- updateGlmerDevfun(devfun, glmod$reTrms))</pre>
## 5. Optimize over theta and beta:
opt <- optimizeGlmer(devfun, stage=2)</pre>
str(opt, max=1) # seeing what we'got
## 6. Package up the results:
(fMod <- mkMerMod(environment(devfun), opt, glmod$reTrms, fr = glmod$fr))</pre>
### Same model in one line
fM <- glmer(cbind(incidence, size - incidence) ~ period + (1 | herd),</pre>
            data = cbpp, family = binomial)
all.equal(fMod, fM, check.attributes=FALSE, tolerance = 1e-12)
# ---- even tolerance = 0 may work
```

NrUnique 25

Description

Remove the random-effects terms from a mixed-effects formula, thereby producing the fixed-effects formula.

Usage

```
nobars(term)
```

Arguments

term

the right-hand side of a mixed-model formula

Value

the fixed-effects part of the formula

Note

This function is called recursively on individual terms in the model, which is why the argument is called term and not a name like form, indicating a formula.

See Also

```
formula, model.frame, model.matrix.
Other utilities: findbars, mkRespMod, mkReTrms, nlformula, subbars
```

Examples

```
nobars(Reaction ~ Days + (Days|Subject)) ## => Reaction ~ Days
```

NrUnique

Number of unique elements in a vector

Description

Number of unique elements in a vector

Usage

```
NrUnique(x, na.rm = TRUE)
```

Arguments

x object of type vector.

na.rm logical indicating if missing values have to be removed. Default to TRUE.

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Value

vector with the number of unique elements

Examples

```
set.seed(1)
x = sample(letters, 50, TRUE)
NrUnique(x)
```

plotRE

Visualizing the random effect estimates using ggplot2

Description

Using this function, you can create plots of the random effect estimates from fitted random effects models. To make the plots, we rely on the ggplot2 package.

Usage

```
plotRE(
  obj,
  levelRE = c("all", "first", "second"),
  colour = "black",
  plot = TRUE
)
```

Arguments

obj an object of type hierCredibility, hierCredGLM or hierCredTweedie

levelRE indicates which hierarchical level has to be used. "all" plots both levels in

the hierarchy, "first" the first level in the hierarchy and "second" the second

level.

colour colour for geom_point

plot logical indicating if the ggplot objects have to be plotted.

Value

a list with ggplot objects.

Examples

```
fitHGLM <- hierCredGLM(Y \sim area + gender + (1 \mid VehicleType / VehicleBody), dataCar, weights = w) \\plotRE(fitHGLM)
```

predict.hierCredGLM 27

Description

Obtain predictions based on the model fit with hierCredGLM

Usage

```
## S3 method for class 'hierCredGLM'
predict(object, newdata = NULL, ...)
```

Arguments

object a model object for which prediction is desired.

newdata optionally, a data frame in which to look for variables with which to predict. If

omitted, the fitted values are used.

... arguments passed to glm

Details

The random effects are taken into account by specifying these as an offset in the predict.glm function.

Value

If newdata is omitted the predictions are based on the data used for the fit.

See Also

hierCredGLM

```
predict.hierCredibility
```

Model predictions

Description

Obtain predictions based on a model fit with hierCredibility

Usage

```
## S3 method for class 'hierCredibility'
predict(object, newdata = NULL, ...)
```

Arguments

object a model object for which prediction is desired.

newdata optionally, a data frame in which to look for variables with which to predict. If

omitted, the fitted values are used.

... ignored.

Value

If newdata is omitted the predictions are based on the data used for the fit.

See Also

```
hierCredibility
```

```
predict.hierCredTweedie
```

Model predictions

Description

Obtain predictions based on the model fit with hierCredTweedie

Usage

```
## S3 method for class 'hierCredTweedie'
predict(object, newdata = NULL, ...)
```

Arguments

object a model object for which prediction is desired.

newdata optionally, a data frame in which to look for variables with which to predict. If

omitted, the fitted values are used.

... arguments passed to cpglm

Details

The random effects are taken into account by specifying these as an offset in the predict.cpglm function.

Value

If newdata is omitted the predictions are based on the data used for the fit.

See Also

hierCredTweedie

print.BalanceProperty 29

print.BalanceProperty Print method for an object of class BalanceProperty

Description

Print method for an object of class BalanceProperty

Usage

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

Arguments

x an object of type BalanceProperty

... Currently ignored.

Value

Prints the call and whether the balance property is satisfied or not. Returns an invisible copy of the original object.

See Also

BalanceProperty

ranef

Extract the modes of the random effects

Description

A generic function to extract the conditional modes of the random effects from a fitted model object. For linear mixed models the conditional modes of the random effects are also the conditional means.

Arguments

object

an object of a class of fitted models with random effects.

Details

If grouping factor i has k levels and j random effects per level the ith component of the list returned by ranef is a data frame with k rows and j columns.

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Value

• From ranef: An object composed of a list of data frames, one for each grouping factor for the random effects. The number of rows in the data frame is the number of levels of the grouping factor. The number of columns is the dimension of the random effect associated with each level of the factor.

Examples

```
library(lattice) ## for dotplot, qqmath
library(lme4)
fm1 <- lmer(Reaction ~ Days + (Days|Subject), sleepstudy)
fm2 <- lmer(Reaction ~ Days + (1|Subject) + (0+Days|Subject), sleepstudy)
fm3 <- lmer(diameter ~ (1|plate) + (1|sample), Penicillin)
ranef(fm1)</pre>
```

ranef-actuaRE

Extract the random effect estimates from a fitted random effects model

Description

A generic function to extract the estimates/predictions of the random effects from a fitted random effects model.

Usage

```
## S3 method for class 'hierCredibility'
ranef(object, ...)
## S3 method for class 'hierCredGLM'
ranef(object, ...)
## S3 method for class 'hierCredTweedie'
ranef(object, ...)
```

Arguments

```
object an object of type hierCredibility, hierCredGLM or hierCredTweedie
... Currently ignored.
```

Value

A list of data frames, one for each grouping factor for the random effects. The number of rows in the data frame is the number of levels of the grouping factor. The first (two) columns correspond(s) to the grouping factor. The last column corresponds to the estimated random effect.

simulatedclustereddata 31

simulatedclustereddata

Simulated data sets to illustrate the package functionality

Description

Both the tweedietraindata and tweedietestdata dataframe are synthetically generated data sets to illustrate the functionality of the package. The tweedietraindata has 250 000 observations and the tweedietestdata has 250 000 observations. The same settings were used to generate both data sets.

Usage

```
data(tweedietraindata)
data(tweedietestdata)
```

Format

```
y the tweedie distributed outcome variable cluster the cluster subcluster the subcluster nested within cluster x1 covariate 1 x2 covariate 2 x3 covariate 3 x4 covariate 4 x5 covariate 5
```

Details

See the examples for how the data sets were generated.

Examples

```
# The data sets were generated as follows
lapply(c("magrittr", "dplyr", "data.table", "tweedie"), library, character.only = TRUE)
set.seed(1)

# Simulate training data
set.seed(1)
nClusters = 5
nSubclusters = 5
p = 5
Uj = scale(rnorm(nClusters))
Ujk = do.call("c", lapply(seq_len(nClusters), function(x) scale(rnorm(nSubclusters))))
nPop = 1e6
nSample = 50
```

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```
nTest
             = 1e3
             = replicate(p, rnorm(nPop))
Beta
             = rnorm(p)
             = sample(seq_len(nClusters), nPop, TRUE)
cluster
subcluster
            = NULL
uniqueCl
             = sort(unique(cluster))
for(cl in uniqueCl)
  subcluster[cluster == cl] <- sample(</pre>
    1 - seq_len(nSubclusters) + which(cl == uniqueCl) * nSubclusters,
    sum(cluster == cl),
    TRUE)
table(subcluster, cluster)
          = X %*% Beta + Uj[match(cluster, seq_len(nClusters))] +
            Ujk[match(subcluster, seq_len(nClusters * nSubclusters))]
          = rtweedie(nPop, mu = exp(as.vector(eta)), phi = 1, power = 1.5)
У
wt
          = runif(nPop)
          = data.frame(y, X, wt, cluster, subcluster)
colnames(Dt) %<>% tolower
tweedietraindata = Dt %>%
  group_by(subcluster) %>%
  sample_n(size = nSample) %>%
  as.data.table
tweedietestdata = Dt %>%
  group_by(subcluster) %>%
  sample_n(size = nSample) %>%
  as.data.table
```

tweedieGLMM

Fitting a Tweedie GLMM, using the initial estimates of hier-CredTweedie

Description

This function first estimates the random effects model using Ohlsson's GLMC algorithm (Ohlsson, 2008) and then uses these estimates as initial estimates when fitting a Tweedie GLMM.

Usage

```
tweedieGLMM(
  formula,
  data,
  weights,
  muHatGLM = FALSE,
  epsilon = 1e-04,
  maxiter = 500,
  verbose = FALSE,
  balanceProperty = TRUE
)
```

tweedieGLMM 33

Arguments

formula	object of type formula that specifies which model should be fitted. Syntax is the same as for lmer and glmer. For example, $Yijkt \sim x1 + x2 + (1 \mid Industry / Branch)$.
data	an object that is coercible by as.data.table, containing the variables in the model.
weights	variable name of the exposure weight.
muHatGLM	indicates which estimate has to be used in the algorithm for the intercept term. Default is TRUE, which used the intercept as estimated by the GLM. If FALSE, the estimate of the hierarchical credibility model is used.
epsilon	positive convergence tolerance ϵ ; the iterations converge when $ \theta[k] - \theta[k-1] ^2[[2]]/ \theta[k-1] ^2[[2]] < \epsilon$. Here, $\theta[k]$ is the parameter vector at the k^{th} iteration.
maxiter	maximum number of iterations.
verbose	logical indicating if output should be produced during the algorithm.
balancePropert	y
	logical indicating if the balance property should be satisfied.

Value

an object of class cpglmm, containing the model fit.

References

Campo, B.D.C. and Antonio, Katrien (2023). Insurance pricing with hierarchically structured data an illustration with a workers' compensation insurance portfolio. *Scandinavian Actuarial Journal*, doi: 10.1080/03461238.2022.2161413

See Also

cpglmm and hierCredTweedie

Examples

```
data("dataCar")
fitTweedieGLMM = tweedieGLMM(Y ~ area + gender + (1 | VehicleType / VehicleBody), dataCar,
   weights = w, verbose = TRUE, epsilon = 1e-4)
```

34 weights-actuaRE

weights-actuaRE

Extract the model weights

Description

weights is a generic function which extracts fitting weights from objects returned by modeling functions. Methods can make use of napredict methods to compensate for the omission of missing values. The default methods does so.

Usage

```
## S3 method for class 'cpglm'
weights(object, type = c("prior", "working"), ...)
## S3 method for class 'hierCredGLM'
weights(object, type = c("prior", "working"), ...)
## S3 method for class 'hierCredTweedie'
weights(object, type = c("prior", "working"), ...)
```

Arguments

object an object for which the extraction of model weights is meaningful. Can be either cpglm, glm, hierCredibility, hierCredGLM or hierCredTweedie type indicates if prior or working weights need to be extracted.
... ignored

Value

Weights extracted from the object object: the default method looks for component "weights" and if not NULL calls napredict on it.

See Also

weights, cpglm, glm, hierCredibility, hierCredGLM or hierCredTweedie

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