

Package ‘galamm’

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Title Generalized Additive Latent and Mixed Models

Version 0.2.0

Description Estimates generalized additive latent and mixed models using maximum marginal likelihood, as defined in Sorensen et al. (2023) [doi:10.1007/s11336-023-09910-z](https://doi.org/10.1007/s11336-023-09910-z), which is an extension of Rabe-Hesketh and Skrondal (2004)'s unifying framework for multilevel latent variable modeling [doi:10.1007/BF02295939](https://doi.org/10.1007/BF02295939). Efficient computation is done using sparse matrix methods, Laplace approximation, and automatic differentiation. The framework includes generalized multilevel models with heteroscedastic residuals, mixed response types, factor loadings, smoothing splines, crossed random effects, and combinations thereof. Syntax for model formulation is close to 'lme4' (Bates et al. (2015) [doi:10.18637/jss.v067.i01](https://doi.org/10.18637/jss.v067.i01)) and 'PLmixed' (Rockwood and Jeon (2019) [doi:10.1080/00273171.2018.1516541](https://doi.org/10.1080/00273171.2018.1516541)).

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URL <https://github.com/LCBC-UiO/galamm>,
<https://lcbc-uo.github.io/galamm/>

BugReports <https://github.com/LCBC-UiO/galamm/issues>

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R topics documented:

anova.galamm	3
coef.galamm	4
cognition	5
confint.galamm	6
deviance.galamm	7
diet	8
epilep	9
extract_optim_parameters.galamm	10
factor_loadings.galamm	11
family.galamm	12
fitted.galamm	13
fixef	14
formula.galamm	15
galamm	16
galamm_control	22
hsced	24
latent_covariates	24
latent_covariates_long	25
lifespan	26
logLik.galamm	27
mresp	28
mresp_hsced	28
nobs.galamm	29
plot.galamm	30
plot_smooth.galamm	31
predict.galamm	32
print.summary.galamm	33
print.VarCorr.galamm	34
ranef.galamm	36
residuals.galamm	37

<code>anova.galammm</code>	3
<code>s</code>	38
<code>sigma.galammm</code>	39
<code>summary.galammm</code>	40
<code>t2</code>	41
<code>VarCorr</code>	43
<code>vcov.galammm</code>	44
Index	46

<code>anova.galammm</code>	<i>Compare likelihoods of galamm objects</i>
----------------------------	--

Description

Anova function for comparing different GALAMMs fitted on the same data.

Usage

```
## S3 method for class 'galamm'
anova(object, ...)
```

Arguments

<code>object</code>	An object of class <code>galamm</code> returned from <code>galamm</code> .
<code>...</code>	Other fitted models of class <code>galamm</code> . Currently, if no models are provided in this argument, no table will be returned.

Value

A table with model comparison metric.

Author(s)

Some of the source code for this function is adapted from `lme4::anova.merMod`, with authors Douglas M. Bates, Martin Maechler, Ben Bolker, and Steve Walker.

References

Bates DM, Mächler M, Bolker B, Walker S (2015). “Fitting Linear Mixed-Effects Models Using Lme4.” *Journal of Statistical Software*, **67**(1), 1–48. ISSN 1548-7660, doi:[10.18637/jss.v067.i01](https://doi.org/10.18637/jss.v067.i01).

See Also

`summary.galammm()` for the summary method and `anova()` for the generic function.
 Other summary functions: `plot.galammm()`, `plot_smooth.galammm()`, `print.summary.galammm()`, `summary.galammm()`

Examples

```

# Poisson GLMM
count_mod <- galamm(
  formula = y ~ lbas * treat + lage + v4 + (1 | subj),
  data = epilep, family = poisson
)

# Model without interaction
count_mod0 <- galamm(
  formula = y ~ lbas + treat + lage + v4 + (1 | subj),
  data = epilep, family = poisson
)

# Model comparison
anova(count_mod, count_mod0)

```

coef.galammm	<i>Extract galamm coefficients</i>
--------------	------------------------------------

Description

Currently, this function only returns the fixed effects.

Usage

```

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

```

Arguments

`object` An object of class `galamm`, from [galamm](#).
`...` Optional arguments passed on to other methods. Currently not used.

Value

A matrix with the requested coefficients.

See Also

[fixef.galammm\(\)](#) for fixed effects, [ranef.galammm\(\)](#) for random effects, and [coef\(\)](#) for the generic function.

Other details of model fit: [VarCorr\(\)](#), [confint.galammm\(\)](#), [deviance.galammm\(\)](#), [factor_loadings.galammm\(\)](#), [family.galammm\(\)](#), [fitted.galammm\(\)](#), [fixef\(\)](#), [formula.galammm\(\)](#), [llikAIC\(\)](#), [logLik.galammm\(\)](#), [nobs.galammm\(\)](#), [predict.galammm\(\)](#), [print.VarCorr.galammm\(\)](#), [ranef.galammm\(\)](#), [residuals.galammm\(\)](#), [sigma.galammm\(\)](#), [vcov.galammm\(\)](#)

Examples

```
# Poisson GLMM
count_mod <- galamm(
  formula = y ~ lbas * treat + lage + v4 + (1 | subj),
  data = epilep, family = poisson
)

# Extract coefficients
coef(count_mod)
```

cognition

Simulated Data with Measurements of Cognitive Abilities

Description

Simulated dataset mimicking the measurement of abilities in three cognitive domains. The latent traits (cognitive ability in a given domain) are based on the functions in `mgcv::gamSim` (Wood 2017), and depend on the explanatory variable `x`.

Usage

```
cognition
```

Format

`cognition` **A data frame with 14400 rows and 7 columns::**

id Subject ID.

domain Factor variable denoting the cognitive domain.

x Explanatory variable.

timepoint Factor variable denoting the timepoint.

item Factor variable denoting the item within the tests of each cognitive domain.

trials Number of trials, if applicable.

y Response variable. For domain 1 a real number, for domain 2 a binomially distributed variable based on a single trial, for domain 3 a real number.

References

Wood SN (2017). *Generalized Additive Models: An Introduction with R*, 2 edition. Chapman and Hall/CRC.

See Also

Other datasets: [diet](#), [epilep](#), [hsced](#), [latent_covariates](#), [latent_covariates_long](#), [lifespan](#), [mresp](#), [mresp_hsced](#)

confint.galammm *Confidence intervals for model parameters*

Description

Confidence intervals for model parameters

Usage

```
## S3 method for class 'galamm'
confint(object, parm, level = 0.95, method = "Wald", ...)
```

Arguments

object	An object of class <code>galamm</code> returned from <code>galamm</code> .
parm	Parameters for which to compute intervals. Use "theta" to get all variance parameters, "beta" to get all fixed regression coefficients, "lambda" to get all factor loadings, and "weights" to get all weights. The parameter can also be given as a numeric vector with indices specifying the parameters. When given as characters, the arguments are case sensitive.
level	Decimal number specifying the confidence level. Defaults to 0.95.
method	Character of length one specifying the type of confidence interval. Currently only "Wald" is available. The argument is case sensitive.
...	Other arguments passed on to other methods. Currently not used.

Value

A matrix with the requested confidence intervals.

See Also

`fixef.galammm()` for fixed effects, `coef.galammm()` for coefficients more generally, and `vcov.galammm()` for the variance-covariance matrix. `confint()` is the generic function.

Other details of model fit: `VarCorr()`, `coef.galammm()`, `deviance.galammm()`, `factor_loadings.galammm()`, `family.galammm()`, `fitted.galammm()`, `fixef()`, `formula.galammm()`, `llikAIC()`, `logLik.galammm()`, `nobs.galammm()`, `predict.galammm()`, `print.VarCorr.galammm()`, `ranef.galammm()`, `residuals.galammm()`, `sigma.galammm()`, `vcov.galammm()`

Examples

```
# Poisson GLMM
count_mod <- galamm(
  formula = y ~ lbas * treat + lage + v4 + (1 | subj),
  data = epilep, family = poisson
)

confint(count_mod, parm = "beta", level = .99)
```

deviance.galammm	<i>Extract deviance of galamm object</i>
------------------	--

Description

Extract deviance of galamm object

Usage

```
## S3 method for class 'galamm'  
deviance(object, ...)
```

Arguments

object	Object of class galamm, returned from galamm .
...	Other arguments passed on to other methods. Currently not used.

Value

A numeric value giving the deviance of the model fit.

See Also

[logLik.galammm\(\)](#) for a function returning the log likelihood and [deviance\(\)](#) for the generic function.

Other details of model fit: [VarCorr\(\)](#), [coef.galammm\(\)](#), [confint.galammm\(\)](#), [factor_loadings.galammm\(\)](#), [family.galammm\(\)](#), [fitted.galammm\(\)](#), [fixef\(\)](#), [formula.galammm\(\)](#), [llikAIC\(\)](#), [logLik.galammm\(\)](#), [nobs.galammm\(\)](#), [predict.galammm\(\)](#), [print.VarCorr.galammm\(\)](#), [ranef.galammm\(\)](#), [residuals.galammm\(\)](#), [sigma.galammm\(\)](#), [vcov.galammm\(\)](#)

Examples

```
# Linear mixed model with heteroscedastic residuals  
mod <- galamm(  
  formula = y ~ x + (1 | id),  
  weights = ~ (1 | item),  
  data = hsced  
)  
  
# Extract deviance  
deviance(mod)
```

diet

Diet Data

Description

Longitudinal epilepsy data from Morris et al. (1977). This documentation is based on Chapter 14.2 of Skrondal and Rabe-Hesketh (2004), where the dataset is used. See also Rabe-Hesketh et al. (2003).

Usage

diet

Format

diet **A data frame with 236 rows and 7 columns::**

id Subject ID.

age Age (standardized).

bus Dummy variable indicating whether the subject is a bus driver or banking staff.

item Integer indicating whether the outcome is fiber intake at time 1 (item = 1), fiber intake at time 2 (item = 2), or coronary heart disease (item = 3).

y Outcome.

chd Dummy variable indicating whether y is an indicator for coronary heart disease, coded as 0/1.

fiber Dummy variable indicating whether y is a fiber measurement at either timepoint 1 or 2.

fiber2 Dummy variable indicating whether y is a fiber measurement at timepoint 2.

Source

<http://www.gllamm.org/books/readme.html#14.2>

References

Morris JN, Marr JW, Clayton DG (1977). “Diet and Heart: A Postscript.” *Br Med J*, **2**(6098), 1307–1314. ISSN 0007-1447, 1468-5833, doi:10.1136/bmj.2.6098.1307.

Rabe-Hesketh S, Pickles A, Skrondal A (2003). “Correcting for Covariate Measurement Error in Logistic Regression Using Nonparametric Maximum Likelihood Estimation.” *Statistical Modelling*, **3**(3), 215–232. ISSN 1471-082X, doi:10.1191/1471082X03st056oa.

Skrondal A, Rabe-Hesketh S (2004). *Generalized Latent Variable Modeling*, Interdisciplinary Statistics Series. Chapman and Hall/CRC, Boca Raton, Florida.

See Also

Other datasets: [cognition](#), [epilep](#), [hsced](#), [latent_covariates](#), [latent_covariates_long](#), [lifespan](#), [mresp](#), [mresp_hsced](#)

epilep

Epilepsy Data

Description

Longitudinal epilepsy data from Leppik et al. (1987). This documentation is based on Chapter 11.3 of Skrondal and Rabe-Hesketh (2004), where the dataset is used.

Usage

epilep

Format

epilep **A data frame with 236 rows and 7 columns::**

subj Subject ID.

y Number of seizures.

treat Dummy variable for treatment group.

visit Time at visit.

v4 Dummy for visit 4.

lage Logarithm of age.

lbas Logarithm of a quarter of the number of seizures in the eight weeks preceding entry into the trial.

Source

<http://www.gllamm.org/books/readme.html#11.3>

References

Leppik IE, Dreifuss FE, Porter R, Bowman T, Santilli N, Jacobs M, Crosby C, Cloyd J, Stackman J, Graves N, Sutula T, Welty T, Vickery J, Brundage R, Gates J, Gumnit RJ, Gutierrez A (1987). "A Controlled Study of Progabide in Partial Seizures: Methodology and Results." *Neurology*, **37**(6), 963–963. ISSN 0028-3878, 1526-632X, doi:[10.1212/WNL.37.6.963](https://doi.org/10.1212/WNL.37.6.963).

Skrondal A, Rabe-Hesketh S (2004). *Generalized Latent Variable Modeling*, Interdisciplinary Statistics Series. Chapman and Hall/CRC, Boca Raton, Florida.

See Also

Other datasets: [cognition](#), [diet](#), [hsced](#), [latent_covariates](#), [latent_covariates_long](#), [lifespan](#), [mresp](#), [mresp_hsced](#)

`extract_optim_parameters.galamm`*Extract parameters from fitted model for use as initial values*

Description

This function extracts parameter values from a fitted model object in a form that can be directly provided as initial values for a new model fit.

Usage

```
## S3 method for class 'galamm'  
extract_optim_parameters(object)
```

Arguments

`object` Object of class `galamm` returned from `galamm`.

Value

A list object containing the following elements:

- `theta` Numerical vector of variance components, i.e., entries of the lower Cholesky form of the covariance matrix of random effects.
- `beta` Fixed regression coefficients.
- `lambda` Factor loadings.
- `weights` Weights for heteroscedastic residuals.

See Also

Other optimization functions: `galamm_control()`

Examples

```
# Fit linear mixed model with heteroscedastic residuals  
mod <- galamm(  
  formula = y ~ x + (1 | id),  
  weights = ~ (1 | item),  
  data = hsced  
)  
  
# Extract parameters  
start <- extract_optim_parameters(mod)  
  
# Fit again using the Nelder-Mead algorithm, using start as initial values:  
mod_nm <- galamm(  
  formula = y ~ x + (1 | id),  
  weights = ~ (1 | item),
```

```
data = hsced,  
start = start,  
control = galamm_control(method = "Nelder-Mead")  
)
```

factor_loadings.galammm

Extract factor loadings from galamm object

Description

Extract factor loadings from galamm object

Usage

```
## S3 method for class 'galamm'  
factor_loadings(object)
```

Arguments

object Object of class galamm returned from [galamm](#).

Details

This function has been named `factor_loadings` rather than just `loadings` to avoid conflict with `stats::loadings`.

Value

A matrix containing the estimated factor loadings with corresponding standard deviations.

Author(s)

The example for this function comes from `PLmixed`, with authors Nicholas Rockwood and Minjeong Jeon (Rockwood and Jeon 2019).

See Also

[fixef.galammm\(\)](#) for fixed regression coefficients, [confint.galammm\(\)](#) for confidence intervals, and [coef.galammm\(\)](#) for coefficients more generally.

Other details of model fit: [VarCorr\(\)](#), [coef.galammm\(\)](#), [confint.galammm\(\)](#), [deviance.galammm\(\)](#), [family.galammm\(\)](#), [fitted.galammm\(\)](#), [fixef\(\)](#), [formula.galammm\(\)](#), [llikAIC\(\)](#), [logLik.galammm\(\)](#), [nobs.galammm\(\)](#), [predict.galammm\(\)](#), [print.VarCorr.galammm\(\)](#), [ranef.galammm\(\)](#), [residuals.galammm\(\)](#), [sigma.galammm\(\)](#), [vcov.galammm\(\)](#)

Examples

```
# Logistic mixed model with factor loadings, example from PLmixed
data("IRTsim", package = "PLmixed")

# Reduce data size for the example to run faster
IRTsub <- IRTsim[IRTsim$item < 4, ]
IRTsub <- IRTsub[sample(nrow(IRTsub), 300), ]
IRTsub$item <- factor(IRTsub$item)

# Fix loading for first item to 1, and estimate the two others freely
loading_matrix <- matrix(c(1, NA, NA), ncol = 1)

# Estimate model
mod <- galamm(y ~ item + (0 + ability | sid) + (0 + ability | school),
  data = IRTsub, family = binomial, load.var = "item",
  factor = "ability", lambda = loading_matrix
)

# Show estimated factor loadings, with standard errors
factor_loadings(mod)
```

family.galammm

Extract family or families from fitted galamm

Description

This function returns a list of families for an object of class `galamm`, returned from `galamm`.

Usage

```
## S3 method for class 'galamm'
family(object, ...)
```

Arguments

`object` An object of class `galamm` returned from `galamm`.
`...` Optional arguments passed on to other methods. Currently not used.

Value

A list of family objects.

See Also

[galamm\(\)](#)

Other details of model fit: [VarCorr\(\)](#), [coef.galammm\(\)](#), [confint.galammm\(\)](#), [deviance.galammm\(\)](#), [factor_loadings.galammm\(\)](#), [fitted.galammm\(\)](#), [fixef\(\)](#), [formula.galammm\(\)](#), [llikAIC\(\)](#), [logLik.galammm\(\)](#), [nobs.galammm\(\)](#), [predict.galammm\(\)](#), [print.VarCorr.galammm\(\)](#), [ranef.galammm\(\)](#), [residuals.galammm\(\)](#), [sigma.galammm\(\)](#), [vcov.galammm\(\)](#)

Examples

```
# Mixed response model
loading_matrix <- matrix(c(1, NA), ncol = 1)
families <- c(gaussian, binomial)
family_mapping <- ifelse(mresp$itemgroup == "a", 1, 2)

mixed_resp <- galamm(
  formula = y ~ x + (0 + level | id),
  data = mresp,
  family = families,
  family_mapping = family_mapping,
  load.var = "itemgroup",
  lambda = loading_matrix,
  factor = "level"
)

# This model has two family objects
family(mixed_resp)
```

fitted.galamm

Extract model fitted values

Description

Extracts fitted values from a model including random effects.

Usage

```
## S3 method for class 'galamm'
fitted(object, ...)
```

Arguments

object An object of class `galamm` returned from `galamm`.
 ... Optional arguments passed on to other methods. Currently not used.

Value

A numerical vector with fit values for each row in the input data.

See Also

Other details of model fit: `VarCorr()`, `coef.galamm()`, `confint.galamm()`, `deviance.galamm()`, `factor_loadings.galamm()`, `family.galamm()`, `fixef()`, `formula.galamm()`, `llikAIC()`, `logLik.galamm()`, `nobs.galamm()`, `predict.galamm()`, `print.VarCorr.galamm()`, `ranef.galamm()`, `residuals.galamm()`, `sigma.galamm()`, `vcov.galamm()`

Examples

```
# Linear mixed model with heteroscedastic residuals
mod <- galamm(
  formula = y ~ x + (1 | id),
  weights = ~ (1 | item),
  data = hsced
)

# Extract fitted values and plot against x
plot(hsced$x, fitted(mod))
```

fixef

Extract fixed effects from galamm objects

Description

Extract the fixed regression coefficients.

Usage

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

Arguments

`object` An object of class `galamm`, returned from `galamm`.
`...` Optional arguments passed on to other methods. Currently not used.

Value

A named numeric vector containing the requested fixed effects.

See Also

`ranef.galamm()` for random effects, `coef.galamm()` for coefficients more generally, and `confint.galamm()` for confidence intervals.

Other details of model fit: `VarCorr()`, `coef.galamm()`, `confint.galamm()`, `deviance.galamm()`, `factor_loadings.galamm()`, `family.galamm()`, `fitted.galamm()`, `formula.galamm()`, `l1kAIC()`, `logLik.galamm()`, `nobs.galamm()`, `predict.galamm()`, `print.VarCorr.galamm()`, `ranef.galamm()`, `residuals.galamm()`, `sigma.galamm()`, `vcov.galamm()`

Examples

```
# Poisson GLMM
count_mod <- galamm(
  formula = y ~ lbas * treat + lage + v4 + (1 | subj),
  data = epilep, family = poisson
)

# Extract fixed effects
fixef(count_mod)
```

formula.galammm	<i>Extract formula from fitted galamm object</i>
-----------------	--

Description

Extract formula from fitted galamm object

Usage

```
## S3 method for class 'galamm'
formula(x, ...)
```

Arguments

x Object of class galamm returned from [galamm](#).

... Optional arguments passed on to other methods. Currently not used.

Value

The formula used to fit the model.

See Also

Other details of model fit: [VarCorr\(\)](#), [coef.galammm\(\)](#), [confint.galammm\(\)](#), [deviance.galammm\(\)](#), [factor_loadings.galammm\(\)](#), [family.galammm\(\)](#), [fitted.galammm\(\)](#), [fixef\(\)](#), [llikAIC\(\)](#), [logLik.galammm\(\)](#), [nobs.galammm\(\)](#), [predict.galammm\(\)](#), [print.VarCorr.galammm\(\)](#), [ranef.galammm\(\)](#), [residuals.galammm\(\)](#), [sigma.galammm\(\)](#), [vcov.galammm\(\)](#)

Examples

```
# Mixed response model -----
# The mresp dataset contains a mix of binomial and Gaussian responses.
# We need to estimate a factor loading which scales the two response types.
loading_matrix <- matrix(c(1, NA), ncol = 1)
```

```

# Define mapping to families.
families <- c(gaussian, binomial)
family_mapping <- ifelse(mresp$itemgroup == "a", 1, 2)

# Fit the model
mod <- galamm(
  formula = y ~ x + (0 + level | id),
  data = mresp,
  family = families,
  family_mapping = family_mapping,
  factor = "level",
  load.var = "itemgroup",
  lambda = loading_matrix
)

# Formula
formula(mod)

```

galamm

Fit a generalized additive latent and mixed model

Description

This function fits a generalized additive latent and mixed model (GALAMMs), as described in Sørensen et al. (2023). The building blocks of these models are generalized additive mixed models (GAMMs) (Wood 2017), of which generalized linear mixed models (Breslow and Clayton 1993; Harville 1977; Henderson 1975; Laird and Ware 1982) are special cases. GALAMMs extend upon GAMMs by allowing factor structures, as commonly used to model hypothesized latent traits underlying observed measurements. In this sense, GALAMMs are an extension of generalized linear latent and mixed models (GLLAMMs) (Skrondal and Rabe-Hesketh 2004; Rabe-Hesketh et al. 2004) which allows semiparametric estimation. The implemented algorithm used to compute model estimates is described in Sørensen et al. (2023), and is an extension of the algorithm used for fitting generalized linear mixed models by the `lme4` package (Bates et al. 2015). The syntax used to define factor structures is based on that used by the `PLmixed` package, which is detailed in Rockwood and Jeon (2019).

Usage

```

galamm(
  formula,
  weights = NULL,
  data,
  family = gaussian,
  family_mapping = rep(1, nrow(data)),
  load.var = NULL,
  lambda = NULL,

```



```

    factor = NULL,
    factor_interactions = NULL,
    na.action = getOption("na.action"),
    start = NULL,
    control = galamm_control()
  )

```

Arguments

formula	A formula specifying the model. Smooth terms are defined in the style of the <code>mgcv</code> and <code>gamm4</code> packages, see (Wood 2017) for an introduction. Random effects are specified using <code>lme4</code> syntax, which is described in detail in (Bates et al. 2015). Factor loadings will also be part of the model formula, and is based on the syntax of the <code>PLmixed</code> package (Rockwood and Jeon 2019).
weights	An optional formula object specifying an expression for the residual variance. Defaults to <code>NULL</code> , corresponding to homoscedastic errors. The formula is defined in <code>lme4</code> style; see vignettes and examples for details.
data	A <code>data.frame</code> containing all the variables specified by the model formula, with the exception of factor loadings.
family	A a list or character vector containing one or more model families. For each element in <code>family</code> there should be a corresponding element in <code>family_mapping</code> specifying which elements of the response are conditionally distributed according to the given family. Currently family can be one of <code>gaussian</code> , <code>binomial</code> , and <code>poisson</code> , and only canonical link functions are supported. The family arguments can either be provided as character values, e.g., <code>c("gaussian", "poisson")</code> or <code>list("gaussian", "poisson")</code> , as function names, e.g., <code>c(gaussian, poisson)</code> or <code>list(gaussian, poisson)</code> , or as function calls, e.g., <code>list(gaussian(), poisson())</code> . In the latter case, they must be provided in a list, and not as a vector. Mixing the different ways of describing the family also works, e.g., <code>list("gaussian", poisson())</code> , but in this case they must be provided in a list. When provided as character values, the argument is case sensitive.
family_mapping	Optional vector mapping from the elements of <code>family</code> to rows of <code>data</code> . Defaults to <code>rep(1, nrow(data))</code> , which means that all observations are distributed according to the first element of <code>family</code> . The length of <code>family_mapping</code> must be identical to the number of observations, <code>nrow(data)</code> .
load.var	Optional character specifying the name of the variable in <code>data</code> identifying what the factors load onto. Default to <code>NULL</code> , which means that there are no loading variables. Argument is case sensitive.
lambda	Optional factor loading matrix. Numerical values indicate that the given value is fixed, while <code>NA</code> means that the entry is a parameter to be estimated. Numerical values can only take the values 0 or <ol style="list-style-type: none"> 1. The number of columns of <code>lambda</code> must be identical to the number of elements in <code>factor</code>. Defaults to <code>NULL</code>, which means that there is no factor loading matrix. If <code>lambda</code> is provided as a vector, it will be converted to a matrix with a single column.
factor	Optional character vector whose j th entry corresponds to the j th column of the corresponding matrix in <code>lambda</code> . The number of elements in <code>factor</code> must be

equal to the number of columns in `lambda`. Defaults to `NULL`, which means that there are no factor loadings. Argument is case sensitive.

<code>factor_interactions</code>	Optional list of length equal to the number of columns in <code>lambda</code> . Each list element should be a formula object containing the write-hand side of a regression model, of the form $\sim x + z$. Defaults to <code>NULL</code> , which means that no factor interactions are used.
<code>na.action</code>	Character of length one specifying a function which indicates what should happen when the data contains NAs. The default is set to the <code>na.action</code> setting of <code>options</code> , which can be seen with <code>options("na.action")</code> . The other alternatives are <code>"na.fail"</code> or <code>"na.exclude"</code> , which means that the function fails if there are NAs in data.
<code>start</code>	Optional named list of starting values for parameters. Possible names of list elements are <code>"theta"</code> , <code>"beta"</code> , <code>"lambda"</code> , and <code>"weights"</code> , all of which should be numerical vectors with starting values. Default to <code>NULL</code> , which means that some relatively sensible defaults are used. Names of parameters must be given in all lower case.
<code>control</code>	Optional control object for the optimization procedure of class <code>galamm_control</code> resulting from calling <code>galamm_control</code> . Defaults to <code>NULL</code> , which means that the defaults of <code>galamm_control</code> are used.

Value

A model object of class `galamm`, containing the following elements:

- `call` the matched call used when fitting the model.
- `random_effects` a list containing the following two elements:
 - `b` random effects in original parametrization.
 - `u` random effects standardized to have identity covariance matrix.
- `model` a list with various elements related to the model setup and fit:
 - `deviance` deviance of final model.
 - `deviance_residuals` deviance residuals of the final model.
 - `df` degrees of freedom of model.
 - `family` a list of one or more family objects, as specified in the `family` arguments to `galamm`.
 - `factor_interactions` List of formulas specifying interactions between latent and observed variables, as provided to the argument `factor_interactions` to `galamm`. If not provided, it is `NULL`.
 - `fit` a numeric vector with fitted values.
 - `fit_population` a numeric vector with fitted values excluding random effects.
 - `hessian` Hessian matrix of final model, i.e., the second derivative of the log-likelihood with respect to all model parameters.
 - `lmod` Linear model object returned by `lme4::lFormula`, which is used internally for setting up the models.
 - `loglik` Log-likelihood of final model.

- n Number of observations.
- pearson_residual Pearson residuals of final model.
- reduced_hessian Logical specifying whether the full Hessian matrix was computed, or a Hessian matrix with derivatives only with respect to beta and lambda.
- response A numeric vector containing the response values used when fitting the model.
- weights_object Object with weights used in model fitting. Is NULL when no weights were used.
- parameters A list object with model parameters and related information:
 - beta_inds Integer vector specifying the indices of fixed regression coefficients among the estimated model parameters.
 - dispersion_parameter One or more dispersion parameters of the final model.
 - lambda_dummy Dummy matrix of factor loadings, which shows the structure of the loading matrix that was supplied in the lambda arguments.
 - lambda_inds Integer vector specifying the indices of factor loadings among the estimated model parameters.
 - lambda_interaction_inds Integer vector specifying the indices of regression coefficients for interactions between latent and observed variables.
 - parameter_estimates Numeric vector of final parameter estimates.
 - parameter_names Names of all parameters estimates.
 - theta_inds Integer vector specifying the indices of variance components among the estimated model parameters. Technically these are the entries of the Cholesky decomposition of the covariance matrix.
 - weights_inds Integer vector specifying the indices of estimated weights (used in heteroscedastic Gaussian models) among the estimated model parameters.
- gam List containing information about smooth terms in the model. If no smooth terms are contained in the model, then it is a list of length zero.

References

- Bates DM, Mächler M, Bolker B, Walker S (2015). “Fitting Linear Mixed-Effects Models Using Lme4.” *Journal of Statistical Software*, **67**(1), 1–48. ISSN 1548-7660, doi:10.18637/jss.v067.i01.
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Skrondal A, Rabe-Hesketh S (2004). *Generalized Latent Variable Modeling*, Interdisciplinary Statistics Series. Chapman and Hall/CRC, Boca Raton, Florida.

Sørensen Ø, Fjell AM, Walhovd KB (2023). “Longitudinal Modeling of Age-Dependent Latent Traits with Generalized Additive Latent and Mixed Models.” *Psychometrika*, **88**(2), 456–486. ISSN 1860-0980, doi:10.1007/s1133602309910z.

Wood SN (2017). *Generalized Additive Models: An Introduction with R*, 2 edition. Chapman and Hall/CRC.

See Also

Other modeling functions: `s()`, `t2()`

Examples

```
# Mixed response model -----
# The mresp dataset contains a mix of binomial and Gaussian responses.
# We need to estimate a factor loading which scales the two response types.
loading_matrix <- matrix(c(1, NA), ncol = 1)

# Define mapping to families.
families <- c(gaussian, binomial)
family_mapping <- ifelse(mresp$itemgroup == "a", 1, 2)

# Fit the model
mod <- galamm(
  formula = y ~ x + (0 + level | id),
  data = mresp,
  family = families,
  family_mapping = family_mapping,
  factor = "level",
  load.var = "itemgroup",
  lambda = loading_matrix
)

# Summary information
summary(mod)

# Heteroscedastic model -----
```

```

# Residuals allowed to differ according to the item variable
# We also set the initial value of the random intercept standard deviation
# to 1
mod <- galamm(
  formula = y ~ x + (1 | id), weights = ~ (1 | item),
  data = hsced, start = list(theta = 1)
)
summary(mod)

# Generalized additive mixed model with factor structures -----

# The cognition dataset contains simulated measurements of three latent
# time-dependent processes, corresponding to individuals' abilities in
# cognitive domains. We focus here on the first domain, and take a single
# random timepoint per person:
dat <- subset(cognition, domain == 1)
dat <- split(dat, f = dat$id)
dat <- lapply(dat, function(x) x[x$timepoint %in% sample(x$timepoint, 1), ])
dat <- do.call(rbind, dat)
dat$item <- factor(dat$item)

# At each timepoint there are three items measuring ability in the cognitive
# domain. We fix the factor loading for the first measurement to one, and
# estimate the remaining two. This is specified in the loading matrix.
loading_matrix <- matrix(c(1, NA, NA), ncol = 1)

# We can now estimate the model.
mod <- galamm(
  formula = y ~ 0 + item + sl(x, factor = "loading") +
    (0 + loading | id),
  data = dat,
  load.var = "item",
  lambda = loading_matrix,
  factor = "loading"
)

# We can plot the estimated smooth term
plot_smooth(mod, shade = TRUE)

# Interaction between observed and latent covariates -----
# Define the loading matrix
lambda <- matrix(c(1, NA, NA), ncol = 1)

# Define the regression functions, one for each row in the loading matrix
factor_interactions <- list(~1, ~1, ~x)

# Fit the model
mod <- galamm(
  formula = y ~ type + x:response + (0 + loading | id),
  data = latent_covariates,
  load.var = "type",
  lambda = lambda,

```

```

    factor = "loading",
    factor_interactions = factor_interactions
  )

# The summary output now include an interaction between the latent variable
# and x, for predicting the third element in "type"
summary(mod)

```

galamm_control	<i>Control values for galamm fit</i>
----------------	--------------------------------------

Description

This function can be called for controlling the optimization procedure used when fitting GALAMMs using [galamm](#).

Usage

```

galamm_control(
  optim_control = list(),
  method = c("L-BFGS-B", "Nelder-Mead"),
  maxit_conditional_modes = 10,
  pirls_tol_abs = 0.01,
  reduced_hessian = FALSE
)

```

Arguments

- | | |
|-------------------------|--|
| optim_control | List containing optimization parameters. If method = "L-BFGS-B" it is passed on to <code>stats::optim</code> 's control argument and if method = "Nelder-Mead", it is passed on to <code>lme4::Nelder_Mead</code> 's control argument. If not otherwise specified, and method = "L-BFGS-B", the following arguments are set to non-default values: <code>fnscale = -1</code> and <code>lmm = 20</code> . |
| method | Character string defining the algorithm to be used for maximizing the marginal log-likelihood. The default is "L-BFGS-B", which uses the limited memory Broyden-Fletcher-Goldfarb-Shanno algorithm with box constrained as implemented in <code>stats::optim</code> . The other options is "Nelder-Mead", which calls the Nelder-Mead algorithm with box constraints implemented in <code>lme4::Nelder_Mead</code> . The argument is case sensitive. |
| maxit_conditional_modes | Maximum number of iterations in penalized iteratively reweighted least squares algorithm. Ignored if family = "gaussian" for all observations, since then a single step gives the exact answer. |
| pirls_tol_abs | Absolute convergence criterion for penalized iteratively reweighted least squares algorithm. Defaults to 0.01, which means that when the reduction in marginal likelihood between two iterations is below 0.01, the iterations stop. |

reduced_hessian

Logical value. Defaults to TRUE, which means that the full Hessian matrix at the maximum marginal likelihood solution is computed. If FALSE, a reduced Hessian matrix with second order partial derivatives with respect to fixed regression coefficients and factor loadings. The latter can help if the full Hessian is not positive definite.

Value

Object of class `galamm_control`, which typically will be provided as an argument to `galamm`.

References

- Bates DM, Mächler M, Bolker B, Walker S (2015). “Fitting Linear Mixed-Effects Models Using lme4.” *Journal of Statistical Software*, **67**(1), 1–48. ISSN 1548-7660, doi:10.18637/jss.v067.i01.
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See Also

`galamm()`

Other optimization functions: `extract_optim_parameters.galamm()`

Examples

```
# Define control object with quite a high degree of verbosity (trace = 6)
# and using the last 20 BFGS updates to estimate the Hessian in L-BFGS-B.
control <- galamm_control(optim_control = list(trace = 6, lmm = 20))
```

hsced

Example Data with Heteroscedastic Residuals

Description

Simulated dataset with residual standard deviation that varies between items.

Usage

hsced

Format

hsced **A data frame with 1200 rows and 5 columns::**

id Subject ID.

age Timepoint.

item Item indicator.

x Explanatory variable

y Outcome.

References

There are no references for Rd macro \insertAllCites on this help page.

See Also

Other datasets: [cognition](#), [diet](#), [epilep](#), [latent_covariates](#), [latent_covariates_long](#), [lifespan](#), [mresp](#), [mresp_hsced](#)

latent_covariates

Simulated Data with Latent and Observed Covariates Interaction

Description

Simulated dataset for use in examples and testing with a latent covariate interacting with an observed covariate.

Usage

latent_covariates

Format

latent_covariates **A data frame with 600 rows and 5 columns::**

id Subject ID.

type Type of observation in the y variable. If it equals "measurement1" or "measurement2" then the observation is a measurement of the latent variable. If it equals "response", then the observation is the actual response.

x Explanatory variable.

y Observed response. Note, this includes both the actual response, and the measurements of the latent variable, since mathematically they are all treated as responses.

response Dummy variable indicating whether the given row is a response or not.

See Also

Other datasets: [cognition](#), [diet](#), [epilep](#), [hsced](#), [latent_covariates_long](#), [lifespan](#), [mresp](#), [mresp_hsced](#)

latent_covariates_long

Simulated Longitudinal Data with Latent and Observed Covariates Interaction

Description

Simulated dataset for use in examples and testing with a latent covariate interacting with an observed covariate. In this data, each response has been measured six times for each subject.

Usage

```
latent_covariates_long
```

Format

latent_covariates_long **A data frame with 800 rows and 5:**

columns:

id Subject ID.

type Type of observation in the y variable. If it equals "measurement1" or "measurement2" then the observation is a measurement of the latent variable. If it equals "response", then the observation is the actual response.

x Explanatory variable.

y Observed response. Note, this includes both the actual response, and the measurements of the latent variable, since mathematically they are all treated as responses.

response Dummy variable indicating whether the given row is a response or not.

See Also

Other datasets: [cognition](#), [diet](#), [epilep](#), [hsced](#), [latent_covariates](#), [lifespan](#), [mresp](#), [mresp_hsced](#)

lifespan	<i>Simulated Dataset with Lifespan Trajectories of Three Cognitive Domains</i>
----------	--

Description

This dataset is simulated based on the data used in Section 4.1 of Sørensen et al. (2023).

Usage

lifespan

Format

lifespan **A data frame with 54,457 rows and 7 columns::**

id Subject ID.

domain Cognitive domain being measured. One of "epmem" for episodic memory, "wmem" for working memory and "execfun" for executive function/speed.

timepoint Integer indicating the timepoint number.

age Age of participant at the timepoint.

test The particular test at this observation.

y Response. For "epmem" and "wmem" this is the number of successes in 16 trials. For "execfun" it is the time in seconds to complete the task.

retest Integer indicating whether the participant has taken the test at a previous timepoint.

domainepmem Dummy variable for domain=="epmem".

domainwmem Dummy variable for domain=="wmem".

domainexecfun Dummy variable for domain=="execfun".

References

Sørensen Ø, Fjell AM, Walhovd KB (2023). "Longitudinal Modeling of Age-Dependent Latent Traits with Generalized Additive Latent and Mixed Models." *Psychometrika*, **88**(2), 456–486. ISSN 1860-0980, doi:[10.1007/s1133602309910z](https://doi.org/10.1007/s1133602309910z).

See Also

Other datasets: [cognition](#), [diet](#), [epilep](#), [hsced](#), [latent_covariates](#), [latent_covariates_long](#), [mresp](#), [mresp_hsced](#)

`logLik.galamm`*Extract Log-Likelihood of galamm Object*

Description

Extract Log-Likelihood of galamm Object

Usage

```
## S3 method for class 'galamm'  
logLik(object, ...)
```

Arguments

<code>object</code>	Object
<code>...</code>	Other arguments

Value

Object of class `logLik`

See Also

[deviance.galamm\(\)](#) for a function returning deviance and [logLik\(\)](#) for the generic function.

Other details of model fit: [VarCorr\(\)](#), [coef.galamm\(\)](#), [confint.galamm\(\)](#), [deviance.galamm\(\)](#), [factor_loadings.galamm\(\)](#), [family.galamm\(\)](#), [fitted.galamm\(\)](#), [fixef\(\)](#), [formula.galamm\(\)](#), [llikAIC\(\)](#), [nobs.galamm\(\)](#), [predict.galamm\(\)](#), [print.VarCorr.galamm\(\)](#), [ranef.galamm\(\)](#), [residuals.galamm\(\)](#), [sigma.galamm\(\)](#), [vcov.galamm\(\)](#)

Examples

```
# Linear mixed model with heteroscedastic residuals  
mod <- galamm(  
  formula = y ~ x + (1 | id),  
  weights = ~ (1 | item),  
  data = hsced  
)  
  
# Extract log likelihood  
logLik(mod)
```

mresp	<i>Simulated Mixed Response Data</i>
-------	--------------------------------------

Description

Very basic mixed response dataset with one set of normally distributed responses and one set of binomially distributed responses.

Usage

mresp

Format

mresp **A data frame with 4000 rows and 5 columns::**

id Subject ID.

x Predictor variable.

y Response.

itemgroup Factor variable which equals "a" for the normally distributed responses and "b" for the binomially distributed response (with 1 trial).

See Also

Other datasets: [cognition](#), [diet](#), [epilep](#), [hsced](#), [latent_covariates](#), [latent_covariates_long](#), [lifespan](#), [mresp_hsced](#)

mresp_hsced	<i>Simulated Mixed Response Data with Heteroscedastic Residuals</i>
-------------	---

Description

Mixed response dataset with one set of normally distributed responses and one set of binomially distributed responses. The normally distributed response follow two different residual standard deviations.

Usage

mresp_hsced

Format

mresp **A data frame with 4000 rows and 5 columns::**

id Subject ID.

x Predictor variable.

y Response.

itemgroup Factor variable which equals "a" for the normally distributed responses and "b" for the binomially distributed response (with 1 trial).

grp Grouping variable denoting which of the two residual standard deviations apply. Only relevant for the normally distributed responses.

isgauss Dummy variable indicating whether the observation on the given line is normally (Gaussian) distributed or not.

See Also

Other datasets: [cognition](#), [diet](#), [epilep](#), [hsced](#), [latent_covariates](#), [latent_covariates_long](#), [lifespan](#), [mresp](#)

nobs.galammm

Extract the Number of Observations from a galamm Fit

Description

Extract the Number of Observations from a galamm Fit

Usage

```
## S3 method for class 'galamm'
nobs(object, ...)
```

Arguments

object An object of class galamm returned from [galamm](#).

... Optional arguments passed on to other methods. Currently not used.

Value

A number

See Also

Other details of model fit: [VarCorr\(\)](#), [coef.galammm\(\)](#), [confint.galammm\(\)](#), [deviance.galammm\(\)](#), [factor_loadings.galammm\(\)](#), [family.galammm\(\)](#), [fitted.galammm\(\)](#), [fixef\(\)](#), [formula.galammm\(\)](#), [llikAIC\(\)](#), [logLik.galammm\(\)](#), [predict.galammm\(\)](#), [print.VarCorr.galammm\(\)](#), [ranef.galammm\(\)](#), [residuals.galammm\(\)](#), [sigma.galammm\(\)](#), [vcov.galammm\(\)](#)

Examples

```
# Example model from lme4
data(sleepstudy, package = "lme4")
fm1 <- galamm(Reaction ~ Days + (Days | Subject), data = sleepstudy)

# There are 180 observations, which matches the number of rows in sleepstudy
nobs(fm1)
nrow(sleepstudy)
```

plot.galamm

Diagnostic plots for galamm objects

Description

Diagnostic plots for galamm objects

Usage

```
## S3 method for class 'galamm'
plot(x, ...)
```

Arguments

`x` An object of class `galamm` returned from `galamm`.
`...` Optional arguments passed on to the plot function.

Value

A plot is displayed.

See Also

`residuals.galamm()` for extracting residuals and `plot()` for the generic function.

Other summary functions: `anova.galamm()`, `plot_smooth.galamm()`, `print.summary.galamm()`, `summary.galamm()`

Examples

```
# Linear mixed model example from lme4
data("sleepstudy", package = "lme4")
mod <- galamm(Reaction ~ Days + (Days | Subject), data = sleepstudy)

# Diagnostic plot
plot(mod)
```

plot_smooth.galamm *Plot smooth terms for galamm fits*

Description

Plots smooth terms of a fitted galamm object. This function is a thin wrapper around `mgcv::plot.gam` (Wood 2017).

Usage

```
## S3 method for class 'galamm'
plot_smooth(object, ...)
```

Arguments

`object` Object of class galamm returned from `galamm`.
`...` Other optional arguments, passed on to `mgcv::plot.gam`.

Value

A plot is displayed on the screen.

References

Wood SN (2017). *Generalized Additive Models: An Introduction with R*, 2 edition. Chapman and Hall/CRC.

See Also

Other summary functions: `anova.galamm()`, `plot.galamm()`, `print.summary.galamm()`, `summary.galamm()`

Examples

```
# Generalized additive mixed model with factor structures -----

# The cognition dataset contains simulated measurements of three latent
# time-dependent processes, corresponding to individuals' abilities in
# cognitive domains. We focus here on the first domain, and take a single
# random timepoint per person:
dat <- subset(cognition, domain == 1)
dat <- split(dat, f = dat$id)
dat <- lapply(dat, function(x) x[x$timepoint %in% sample(x$timepoint, 1), ])
dat <- do.call(rbind, dat)
dat$item <- factor(dat$item)

# At each timepoint there are three items measuring ability in the cognitive
# domain. We fix the factor loading for the first measurement to one, and
# estimate the remaining two. This is specified in the loading matrix.
loading_matrix <- matrix(c(1, NA, NA), ncol = 1)
```

```

# We can now estimate the model.
mod <- galamm(
  formula = y ~ 0 + item + sl(x, factor = "loading") +
    (0 + loading | id),
  data = dat,
  load.var = "item",
  lambda = loading_matrix,
  factor = "loading"
)

# We can plot the estimated smooth term
plot_smooth(mod, shade = TRUE)

# We can turn off the rug at the bottom
plot_smooth(mod, shade = TRUE, rug = FALSE)

```

predict.galammm

Predictions from a model at new data values

Description

Predictions are given at the population level, i.e., with random effects set to zero. For fitted models including random effects, see [fitted.galammm](#). For mixed response models, only predictions on the scale of the linear predictors is supported.

Usage

```

## S3 method for class 'galamm'
predict(object, newdata = NULL, type = c("link", "response"), ...)

```

Arguments

object	An object of class <code>galamm</code> returned from galamm .
newdata	Data from for which to evaluate predictions, in a <code>data.frame</code> . Defaults to "NULL", which means that the predictions are evaluate at the data used to fit the model.
type	Character argument specifying the type of prediction object to be returned. Case sensitive.
...	Optional arguments passed on to other methods. Currently used for models with smooth terms, for which these arguments are forwarded to <code>mgcv::predict.gam</code> .

Value

A numeric vector of predicted values.

See Also

`fitted.galamm()` for model fits, `residuals.galamm()` for residuals, and `predict()` for the generic function.

Other details of model fit: `VarCorr()`, `coef.galamm()`, `confint.galamm()`, `deviance.galamm()`, `factor_loadings.galamm()`, `family.galamm()`, `fitted.galamm()`, `fixef()`, `formula.galamm()`, `llikAIC()`, `logLik.galamm()`, `nobs.galamm()`, `print.VarCorr.galamm()`, `ranef.galamm()`, `residuals.galamm()`, `sigma.galamm()`, `vcov.galamm()`

Examples

```
# Poisson GLMM
count_mod <- galamm(
  formula = y ~ lbas * treat + lage + v4 + (1 | subj),
  data = epilep, family = poisson
)

# Plot response versus link:
plot(
  predict(count_mod, type = "link"),
  predict(count_mod, type = "response")
)

# Predict on a new dataset
nd <- data.frame(lbas = c(.3, .2), treat = c(0, 1), lage = 0.2, v4 = -.2)
predict(count_mod, newdata = nd)
predict(count_mod, newdata = nd, type = "response")
```

`print.summary.galamm` *Print method for summary GALAMM fits*

Description

Print method for summary GALAMM fits

Usage

```
## S3 method for class 'summary.galamm'
print(x, digits = max(3, getOption("digits") - 3), ...)
```

Arguments

<code>x</code>	An object of class <code>summary.galamm</code> returned from <code>summary.galamm</code> .
<code>digits</code>	Number of digits to present in outputs.
<code>...</code>	Further arguments passed on to other methods. Currently used by <code>stats::printCoefmat</code> for printing approximate significance of smooth terms.

Value

Summary printed to screen. Invisibly returns the argument `x`.

Author(s)

Some of the code for producing summary information has been derived from the summary methods of `mgcv` (author: Simon Wood) and `lme4` (Bates et al. 2015) (authors: Douglas M. Bates, Martin Maechler, Ben Bolker, and Steve Walker).

References

There are no references for Rd macro `\insertAllCites` on this help page.

See Also

[summary.galamm\(\)](#) for the summary function and [print\(\)](#) for the generic function.

Other summary functions: [anova.galamm\(\)](#), [plot.galamm\(\)](#), [plot_smooth.galamm\(\)](#), [summary.galamm\(\)](#)

Examples

```
# Linear mixed model with heteroscedastic residuals
mod <- galamm(
  formula = y ~ x + (1 | id),
  weights = ~ (1 | item),
  data = hsced
)

summary(mod)
```

`print.VarCorr.galamm` *Print method for variance-covariance objects*

Description

Print method for variance-covariance objects

Usage

```
## S3 method for class 'VarCorr.galamm'
print(
  x,
  digits = max(3, getOption("digits") - 2),
  comp = c("Std.Dev.", "Variance"),
  corr = any(comp == "Std.Dev."),
  ...
)
```

Arguments

x	An object of class <code>c("VarCorr.galamm", "VarCorr.merMod")</code> , returned from VarCorr.galamm .
digits	Optional arguments specifying number of digits to use when printing.
comp	Character vector of length 1 or 2 specifying which variance components to print. Case sensitive. Can take one of the values "Std.Dev." and "Variance".
corr	Logical value indicating whether covariances or correlations should be printed.
...	Optional arguments passed on to other methods. Currently not used.

Value

The variance-covariance information is printed to the console and the argument `x` is silently returned.

Author(s)

This function is derived from `lme4:::print.VarCorr.merMod` written by Douglas M. Bates, Martin Maechler, Ben Bolker, and Steve Walker.

References

Bates DM, Mächler M, Bolker B, Walker S (2015). "Fitting Linear Mixed-Effects Models Using Lme4." *Journal of Statistical Software*, **67**(1), 1–48. ISSN 1548-7660, doi:[10.18637/jss.v067.i01](https://doi.org/10.18637/jss.v067.i01).

See Also

[VarCorr.galamm\(\)](#) for the function creating the variance-covariance objects.

Other details of model fit: [VarCorr\(\)](#), [coef.galamm\(\)](#), [confint.galamm\(\)](#), [deviance.galamm\(\)](#), [factor_loadings.galamm\(\)](#), [family.galamm\(\)](#), [fitted.galamm\(\)](#), [fixef\(\)](#), [formula.galamm\(\)](#), [llikAIC\(\)](#), [logLik.galamm\(\)](#), [nobs.galamm\(\)](#), [predict.galamm\(\)](#), [ranef.galamm\(\)](#), [residuals.galamm\(\)](#), [sigma.galamm\(\)](#), [vcov.galamm\(\)](#)

Examples

```
# Linear mixed model with heteroscedastic residuals
mod <- galamm(
  formula = y ~ x + (1 | id),
  weights = ~ (1 | item),
  data = hsced
)

# Extract information on variance and covariance
VarCorr(mod)
```

ranef.galammm	<i>Extract random effects from galamm object.</i>
---------------	---

Description

Extract random effects from galamm object.

Usage

```
## S3 method for class 'galamm'  
ranef(object, ...)
```

Arguments

object	An object of class galamm, returned from galamm .
...	Optional parameters passed on to other methods. Currently not used.

Value

An object of class ranef.galammm, containing the requested random effects.

Author(s)

This function is derived from `lme4::ranef.merMod`, written by Douglas Bates, Martin Maechler, Ben Bolker, Steve Walker.

References

Bates DM, Mächler M, Bolker B, Walker S (2015). “Fitting Linear Mixed-Effects Models Using Lme4.” *Journal of Statistical Software*, **67**(1), 1–48. ISSN 1548-7660, doi:[10.18637/jss.v067.i01](https://doi.org/10.18637/jss.v067.i01).

See Also

[fixef.galammm\(\)](#) for fixed effects and [coef.galammm\(\)](#) for coefficients more generally.

Other details of model fit: [VarCorr\(\)](#), [coef.galammm\(\)](#), [confint.galammm\(\)](#), [deviance.galammm\(\)](#), [factor_loadings.galammm\(\)](#), [family.galammm\(\)](#), [fitted.galammm\(\)](#), [fixef\(\)](#), [formula.galammm\(\)](#), [llikAIC\(\)](#), [logLik.galammm\(\)](#), [nobs.galammm\(\)](#), [predict.galammm\(\)](#), [print.VarCorr.galammm\(\)](#), [residuals.galammm\(\)](#), [sigma.galammm\(\)](#), [vcov.galammm\(\)](#)

Examples

```
# Poisson GLMM  
count_mod <- galamm(  
  formula = y ~ lbas * treat + lage + v4 + (1 | subj),  
  data = epilep, family = poisson  
)  
  
# Extract random effects
```

```
ranef(count_mod)
```

```
residuals.galamm      Residuals of galamm objects
```

Description

Residuals of galamm objects

Usage

```
## S3 method for class 'galamm'
residuals(object, type = c("pearson", "deviance"), ...)
```

Arguments

object	An object of class galamm returned from galamm .
type	Character of length one describing the type of residuals to be returned. One of "pearson" and "deviance". Argument is case sensitive.
...	Optional arguments passed on to other methods. Currently not used.

Value

Numeric vector of residual values.

See Also

[fitted.galamm\(\)](#) for model fitted values, [predict.galamm\(\)](#) for model predictions, and [plot.galamm\(\)](#) for diagnostic plots. The generic function is [residuals\(\)](#).

Other details of model fit: [VarCorr\(\)](#), [coef.galamm\(\)](#), [confint.galamm\(\)](#), [deviance.galamm\(\)](#), [factor_loadings.galamm\(\)](#), [family.galamm\(\)](#), [fitted.galamm\(\)](#), [fixef\(\)](#), [formula.galamm\(\)](#), [llikAIC\(\)](#), [logLik.galamm\(\)](#), [nobs.galamm\(\)](#), [predict.galamm\(\)](#), [print.VarCorr.galamm\(\)](#), [ranef.galamm\(\)](#), [sigma.galamm\(\)](#), [vcov.galamm\(\)](#)

Examples

```
# Poisson GLMM
count_mod <- galamm(
  formula = y ~ lbas * treat + lage + v4 + (1 | subj),
  data = epilep, family = poisson
)

# Extract residuals
residuals(count_mod)
```

s

Set up smooth term with factor loading

Description

This is a very thin wrapper around `mgcv::s`. It enables the specification of loading variables for smooth terms. The last letter "l", which stands for "loading", has been added to avoid namespace conflicts with `mgcv` and `gamm4`.

Usage

```
s1(..., factor = NULL)
```

Arguments

`...` Arguments passed on to `mgcv::s`.
`factor` Optional character argument specifying the loading variable. Case sensitive.

Details

The documentation of the function `mgcv::s` should be consulted for details on how to properly set up smooth terms. In particular, note that these terms distinguish between ordered and unordered factor terms in the `by` variable, which can be provided in `...` and is forwarded to `mgcv::s`.

Value

An object of class `xx.smooth.spec`, where `xx` is a basis identifying code given by the `bs` argument of `s`. It differs from the smooth returned by `mgcv::s` in that it has an additional attribute named `"factor"` which specifies any factor loading which this smooth term should be multiplied with in order to produce the observed outcome.

References

Wood SN (2003). "Thin Plate Regression Splines." *Journal of the Royal Statistical Society. Series B (Statistical Methodology)*, **65**(1), 95–114. ISSN 1369-7412.

Wood SN (2017). *Generalized Additive Models: An Introduction with R*, 2 edition. Chapman and Hall/CRC.

See Also

Other modeling functions: `galamm()`, `t2()`

Examples

```
# Linear mixed model with factor structures
dat <- subset(cognition, domain == 1 & timepoint == 1)
loading_matrix <- matrix(c(1, NA, NA), ncol = 1)

# Model with four thin-plate regression splines as basis functions
mod <- galamm(
  formula = y ~ 0 + item + sl(x, k = 4, factor = "loading"),
  data = dat,
  load.var = "item",
  lambda = loading_matrix,
  factor = "loading"
)

# Model with four cubic regression splines as basis functions
mod <- galamm(
  formula = y ~ 0 + item +
    sl(x, bs = "cr", k = 4, factor = "loading"),
  data = dat,
  load.var = "item",
  lambda = loading_matrix,
  factor = "loading"
)
```

sigma.galamm

Extract square root of dispersion parameter from galamm object

Description

Extracts the square root of the dispersion parameter(s) from an object of class `galamm`, returned from [galamm](#). In the case of conditionally Gaussian responses, this is the residual standard deviation. When there are multiple dispersion parameters, e.g., with mixed response type models, the square root of all of them are returned in a numeric vector.

Usage

```
## S3 method for class 'galamm'
sigma(object, ...)
```

Arguments

<code>object</code>	An object of class <code>galamm</code> , returned from galamm .
<code>...</code>	Optional parameters passed on to other methods. Currently not used.

Value

The square root of one or more dispersion parameters.

See Also[galamm\(\)](#)

Other details of model fit: [VarCorr\(\)](#), [coef.galamm\(\)](#), [confint.galamm\(\)](#), [deviance.galamm\(\)](#), [factor_loadings.galamm\(\)](#), [family.galamm\(\)](#), [fitted.galamm\(\)](#), [fixef\(\)](#), [formula.galamm\(\)](#), [llikAIC\(\)](#), [logLik.galamm\(\)](#), [nobs.galamm\(\)](#), [predict.galamm\(\)](#), [print.VarCorr.galamm\(\)](#), [ranef.galamm\(\)](#), [residuals.galamm\(\)](#), [vcov.galamm\(\)](#)

Examples

```
# Linear mixed model with heteroscedastic residuals
mod <- galamm(
  formula = y ~ x + (1 | id),
  weights = ~ (1 | item),
  data = hsced
)

# Extract residual standard deviation.
sigma(mod)

# The residual standard deviation applies to the base case. The variance
# function shown in the model output shows the estimated multiplier for
# various grouping levels:
summary(mod)
```

summary.galamm

*Summarizing GALAMM fits***Description**

Summary method for class "galamm".

Usage

```
## S3 method for class 'galamm'
summary(object, ...)
```

Arguments

object An object of class galamm returned from [galamm](#).
... Further arguments passed on to other methods. Currently not used.

Value

A list of summary statistics of the fitted model of class `summary.galamm`, containing the following elements:

- AICtab a table of model fit measures, returned by [llikAIC](#).

- call the matched call used when fitting the model.
- `fixef` a matrix with fixed effect estimated, returned by `fixef`.
- `gam` List containing information about smooth terms in the model. If no smooth terms are contained in the model, then it is a list of length zero.
- `model` a list with various elements related to the model setup and fit. See `?galamm` for details.
- `parameters` A list object with model parameters and related information. See `?galamm` for details.
- `Lambda` An object containing the estimated factor loadings. Returned from `factor_loadings.galamm`. If there are no estimated factor loadings, then this object is `NULL`.
- `random_effects` a list containing the random effects. See `?galamm` for details.
- `VarCorr` An object of class `VarCorr.galamm`, returned from `VarCorr.galamm`.
- `weights` An object containing information about estimated variance functions, when there are heteroscedastic residuals. Otherwise the object is `NULL`.

Author(s)

Some of the code for producing summary information has been derived from the summary methods of `mgcv` (author: Simon Wood) and `lme4` (Bates et al. 2015) (authors: Douglas M. Bates, Martin Maechler, Ben Bolker, and Steve Walker).

See Also

`print.summary.galamm()` for the print method and `summary()` for the generic.

Other summary functions: `anova.galamm()`, `plot.galamm()`, `plot_smooth.galamm()`, `print.summary.galamm()`

Examples

```
# Linear mixed model with heteroscedastic residuals
mod <- galamm(
  formula = y ~ x + (1 | id),
  weights = ~ (1 | item),
  data = hsced
)

summary(mod)
```

Description

This is a very thin wrapper around `mgcv::t2`. It enables the specification of loading variables for smooth terms. The last letter "l", which stands for "loading", has been added to avoid namespace conflicts with `mgcv` and `gamm4`.

Usage

```
t2l(..., factor = NULL)
```

Arguments

... Arguments passed on to `mgcv::t2`.

factor Optional character of length one specifying the loading variable. Case sensitive.

Details

The documentation of the function `mgcv::t2` should be consulted for details on how to properly set up smooth terms. In particular, note that these terms distinguish between ordered and unordered factor terms in the by variable, which can be provided in ... and is forwarded to `mgcv::t2`.

Value

An object of class `xx.smooth.spec`, where `xx` is a basis identifying code given by the `bs` argument of `t2`. It differs from the smooth returned by `mgcv::s` in that it has an additional attribute named "factor" which specifies any factor loading which this smooth term should be multiplied with in order to produce the observed outcome.

References

Wood SN (2003). "Thin Plate Regression Splines." *Journal of the Royal Statistical Society. Series B (Statistical Methodology)*, **65**(1), 95–114. ISSN 1369-7412.

Wood SN (2017). *Generalized Additive Models: An Introduction with R*, 2 edition. Chapman and Hall/CRC.

See Also

Other modeling functions: [galamm\(\)](#), [s\(\)](#)

Examples

```
# Linear mixed model with factor structures
dat <- subset(cognition, domain == 1 & timepoint == 1)
loading_matrix <- matrix(c(1, NA, NA), ncol = 1)

# Model with four cubic regression splines as basis functions
mod <- galamm(
  formula = y ~ 0 + item + t2l(x, k = 4, factor = "loading"),
  data = dat,
  load.var = "item",
  lambda = loading_matrix,
  factor = "loading"
)

# Model with four thin-plate regression splines as basis functions
mod <- galamm(
  formula = y ~ 0 + item +
```

```

    t2l(x, bs = "tp", k = 4, factor = "loading"),
    data = dat,
    load.var = "item",
    lambda = loading_matrix,
    factor = "loading"
  )

```

 VarCorr

Extract variance and correlation components from model

Description

Extract variance and correlation components from model

Usage

```

## S3 method for class 'galamm'
VarCorr(x, sigma = 1, ...)

```

Arguments

<code>x</code>	An object of class <code>galamm</code> returned from <code>galamm</code> .
<code>sigma</code>	Numeric value used to multiply the standard deviations. Defaults to 1.
<code>...</code>	Other arguments passed onto other methods. Currently not used.

Value

An object of class `c("VarCorr.galamm", "VarCorr.merMod")`.

See Also

`print.VarCorr.galamm()` for the print function.

Other details of model fit: `coef.galamm()`, `confint.galamm()`, `deviance.galamm()`, `factor_loadings.galamm()`, `family.galamm()`, `fitted.galamm()`, `fixef()`, `formula.galamm()`, `llikAIC()`, `logLik.galamm()`, `nobs.galamm()`, `predict.galamm()`, `print.VarCorr.galamm()`, `ranef.galamm()`, `residuals.galamm()`, `sigma.galamm()`, `vcov.galamm()`

Examples

```

# Linear mixed model with heteroscedastic residuals
mod <- galamm(
  formula = y ~ x + (1 | id),
  weights = ~ (1 | item),
  data = hsced
)

# Extract information on variance and covariance

```

```

VarCorr(mod)

# Convert to data frame
# (this invokes lme4's function as.data.frame.VarCorr.merMod)
as.data.frame(VarCorr(mod))

```

vcov.galammm

Calculate variance-covariance matrix for GALAMM fit

Description

Calculate variance-covariance matrix for GALAMM fit

Usage

```

## S3 method for class 'galamm'
vcov(object, parm = "beta", ...)

```

Arguments

object	Object of class <code>galamm</code> returned from <code>galamm</code> .
parm	The parameters for which the variance-covariance matrix should be calculated. Character vector with one or more of the elements "theta", "beta", "lambda", and "weights". Can also be an integer vector. When given as a character, it must be in only lowercase letters.
...	Further arguments passed on to other methods. Currently not used.

Value

A variance-covariance matrix.

See Also

`confint.galammm()` for the method computing confidence intervals. See `vcov()` for the generic function.

Other details of model fit: `VarCorr()`, `coef.galammm()`, `confint.galammm()`, `deviance.galammm()`, `factor_loadings.galammm()`, `family.galammm()`, `fitted.galammm()`, `fixef()`, `formula.galammm()`, `llikAIC()`, `logLik.galammm()`, `nobs.galammm()`, `predict.galammm()`, `print.VarCorr.galammm()`, `ranef.galammm()`, `residuals.galammm()`, `sigma.galammm()`

Examples

```
# Linear mixed model with heteroscedastic residuals
mod <- galamm(
  formula = y ~ x + (1 | id),
  weights = ~ (1 | item),
  data = hsced
)

# Extract covariance matrix for fixed regression coefficients
vcov(mod, parm = "beta")

# and then for weights, which gives us the variance.
vcov(mod, parm = "weights")
```

Index

- * **datasets**
 - cognition, 5
 - diet, 8
 - epilep, 9
 - hsced, 24
 - latent_covariates, 24
 - latent_covariates_long, 25
 - lifespan, 26
 - mresp, 28
 - mresp_hsced, 28
- * **details of model fit**
 - coef.galamm, 4
 - confint.galamm, 6
 - deviance.galamm, 7
 - factor_loadings.galamm, 11
 - family.galamm, 12
 - fitted.galamm, 13
 - fixef, 14
 - formula.galamm, 15
 - logLik.galamm, 27
 - nobs.galamm, 29
 - predict.galamm, 32
 - print.VarCorr.galamm, 34
 - ranef.galamm, 36
 - residuals.galamm, 37
 - sigma.galamm, 39
 - VarCorr, 43
 - vcov.galamm, 44
- * **modeling functions**
 - galamm, 16
 - s, 38
 - t2, 41
- * **optimization functions**
 - extract_optim_parameters.galamm, 10
 - galamm_control, 22
- * **summary functions**
 - anova.galamm, 3
 - plot.galamm, 30
 - plot_smooth.galamm, 31
 - print.summary.galamm, 33
 - summary.galamm, 40
- anova(), 3
- anova.galamm, 3, 30, 31, 34, 41
- coef(), 4
- coef.galamm, 4, 6, 7, 11–15, 27, 29, 33, 35–37, 40, 43, 44
- coef.galamm(), 6, 11, 14, 36
- cognition, 5, 8, 9, 24–26, 28, 29
- confint(), 6
- confint.galamm, 4, 6, 7, 11–15, 27, 29, 33, 35–37, 40, 43, 44
- confint.galamm(), 11, 14, 44
- deviance(), 7
- deviance.galamm, 4, 6, 7, 11–15, 27, 29, 33, 35–37, 40, 43, 44
- deviance.galamm(), 27
- diet, 5, 8, 9, 24–26, 28, 29
- epilep, 5, 8, 9, 24–26, 28, 29
- extract_optim_parameters
 - (extract_optim_parameters.galamm), 10
- extract_optim_parameters.galamm, 10, 23
- factor_loadings
 - (factor_loadings.galamm), 11
- factor_loadings.galamm, 4, 6, 7, 11, 12–15, 27, 29, 33, 35–37, 40, 41, 43, 44
- family.galamm, 4, 6, 7, 11, 12, 13–15, 27, 29, 33, 35–37, 40, 43, 44
- fitted.galamm, 4, 6, 7, 11, 12, 13, 14, 15, 27, 29, 32, 33, 35–37, 40, 43, 44
- fitted.galamm(), 33, 37
- fixef, 4, 6, 7, 11–13, 14, 15, 27, 29, 33, 35–37, 40, 41, 43, 44
- fixef.galamm(), 4, 6, 11, 36

- formula.galamm*, 4, 6, 7, 11–14, 15, 27, 29, 33, 35–37, 40, 43, 44
galamm, 3, 4, 6, 7, 10–15, 16, 22, 23, 29–32, 36–40, 42–44
galamm(), 12, 23, 40
galamm_control, 10, 18, 22
hsced, 5, 8, 9, 24, 25, 26, 28, 29
latent_covariates, 5, 8, 9, 24, 24, 25, 26, 28, 29
latent_covariates_long, 5, 8, 9, 24, 25, 25, 26, 28, 29
lifespan, 5, 8, 9, 24, 25, 26, 28, 29
llikAIC, 4, 6, 7, 11–15, 27, 29, 33, 35–37, 40, 43, 44
logLik(), 27
logLik.galamm, 4, 6, 7, 11–15, 27, 29, 33, 35–37, 40, 43, 44
logLik.galamm(), 7
mresp, 5, 8, 9, 24–26, 28, 29
mresp_hsced, 5, 8, 9, 24–26, 28, 28
nobs.galamm, 4, 6, 7, 11–15, 27, 29, 33, 35–37, 40, 43, 44

plot(), 30
plot.galamm, 3, 30, 31, 34, 41
plot.galamm(), 37
plot_smooth(plot_smooth.galamm), 31
plot_smooth.galamm, 3, 30, 31, 34, 41
predict(), 33
predict.galamm, 4, 6, 7, 11–15, 27, 29, 32, 35–37, 40, 43, 44
predict.galamm(), 37
print(), 34
print.summary.galamm, 3, 30, 31, 33, 41
print.summary.galamm(), 41
print.VarCorr.galamm, 4, 6, 7, 11–15, 27, 29, 33, 34, 36, 37, 40, 43, 44
print.VarCorr.galamm(), 43

ranef(ranef.galamm), 36
ranef.galamm, 4, 6, 7, 11–15, 27, 29, 33, 35, 36, 37, 40, 43, 44
ranef.galamm(), 4, 14
residuals(), 37

residuals.galamm, 4, 6, 7, 11–15, 27, 29, 33, 35, 36, 37, 40, 43, 44
residuals.galamm(), 30, 33

s, 20, 38, 42
sigma.galamm, 4, 6, 7, 11–15, 27, 29, 33, 35–37, 39, 43, 44
sl(s), 38
summary(), 41
summary.galamm, 3, 30, 31, 33, 34, 40
summary.galamm(), 3, 34

t2, 20, 38, 41
t2l(t2), 41

VarCorr, 4, 6, 7, 11–15, 27, 29, 33, 35–37, 40, 43, 44
VarCorr.galamm, 35, 41
VarCorr.galamm(), 35
vcov(), 44
vcov.galamm, 4, 6, 7, 11–15, 27, 29, 33, 35–37, 40, 43, 44
vcov.galamm(), 6