

Package ‘gvcn.cat’

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Type Package

Title Regularized Categorical Effects/Categorical Effect
Modifiers/Continuous/Smooth Effects in GLMs

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Author Margret Oelker

Maintainer Margret Oelker <Margret.Oelker@stat.uni-muenchen.de>

Description Generalized structured regression models with regularized categorical effects, categorical effect modifiers, continuous effects and smooth effects.

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cat_control

*Auxiliary Function for gvcml.cat***Description**

Auxiliary function for `gvcml.cat`. Modifies the algorithm's internal parameters.

Usage

```
cat_control(center = FALSE, standardize = FALSE, accuracy = 2, digits = 4,
g = 0.5, epsilon = 10^(-5), maxi = 250, c = 10^(-5), gama = 20, steps = 25,
nu = 1, tuning.criterion = "GCV", K = 5, cv.refit = FALSE,
lambda.upper=50, lambda.lower=0, lambda.accuracy=.01, scaled.lik=FALSE,
adapted.weights=FALSE, adapted.weights.adj = FALSE, adapted.weights.ridge =
FALSE, assured.intercept=TRUE,
level.control = FALSE, case.control = FALSE, pairwise = TRUE,
grouped.cat.diffs = FALSE, bootstrap = 0, start.ml = FALSE, L0.log = TRUE,
subjspec.gr = FALSE, high = NULL, ...)
```

Arguments

center	logical; if TRUE, all metric covariates are centered by their empirical mean
standardize	logical; if TRUE, the design matrix is standardized by its (weighted) empirical variances
accuracy	integer; number of digits being compared when setting coefficients equal/to zero
digits	integer; number of digits for estimates
g	step length parameter for the PIRLS-algorithm; out of)0,1(
epsilon	small, positive, real constant; the PIRLS-algorithm is terminated when the (scaled, absolute) difference of the coefficients of the current iteration and the coefficients of the previous iteration is smaller than epsilon
maxi	integer; maximal number of iterations in the fitting algorithm
c	small, positive, real constant; needed for the approximation of the absolute value function in the PIRLS-algorithm
gama	positive number; tuning parameter for the approximation of the L0 norm
steps	integer; tuning parameter for path-plotting; minimal number of estimates employed for path-plotting
nu	optional weighting parameter
tuning.criterion	loss criterion for cross-validation; one out of "GCV" (generalized cross validation criterion), "deviance" (K-fold cross-validation with the predictive deviance as criterion)
K	integer; number of folds for cross-validation
cv.refit	logical; if TRUE, cross-validation is based on a refit of the selected coefficients

lambda.upper	integer; upper bound for cross-validation of lambda
lambda.lower	integer; lower bound for cross-validation of lambda
lambda.accuracy	numeric; how accurate shall lambda be cross-validated?; minimal absolute difference between two candidates for lambda
scaled.lik	if TRUE, the likelihood in the objective function is scaled by 1/n
adapted.weights	logical; if TRUE, penalty terms are weighted adaptively, that is by inverse ML-estimates; set to FALSE, if ML-estimates do not exist/are too close to zero; only for specials v, p, grouped, SCAD, elastic
adapted.weights.adj	logical; if TRUE, adapted weights of several categorical covariates are scaled such that they are comparable
adapted.weights.ridge	logical; if TRUE, adapted weights are based on an estimate that is slightly penalized by a Ridge penalty
assured.intercept	logical; shall a constant intercept remain in the model in any case?
level.control	logical; if TRUE, the penalty terms are adjusted for different number of penalty terms per covariate
case.control	logical; if TRUE, the penalty terms are adjusted for the number of observations on each level of a categorical covariate
pairwise	experimental option; disabled if TRUE
grouped.cat.diff	experimental option; disabled if FALSE
bootstrap	experimental option; disabled if 0
start.ml	logical; if TRUE, the initial value is the ML-estimate
L0.log	experimental option; disabled if TRUE
subjspec.gr	experimental option; disabled if FALSE
high	experimental option; disabled if NULL
...	further arguments passed to or from other methods

Value

Returns a list containing the (checked) input arguments.

See Also

Function [gvcm.cat](#)

gvcm.cat	<i>Regularized Categorical Effects/Categorical Effect Modifiers/Continuous/Smooth effects in GLMs</i>
----------	---

Description

The function fits generalized linear models with regularized categorical effects, categorical effect modifiers, continuous effects and smooth effects. The model is specified by giving a symbolic description of the linear predictor and a description of the error distribution. Estimation employs different regularization and model selection strategies. These strategies are either a penalty or a forward selection strategy employing AIC/BIC. For non-differentiable penalties, a local quadratic approximation is employed, see Oelker and Tutz (2013).

Usage

```
gvcm.cat(formula, data, family = gaussian, method = c("lqa", "AIC", "BIC"),
tuning = list(lambda=TRUE, specific=FALSE, phi=0.5, grouped.fused=0.5,
elastic=0.5, vs=0.5, spl=0.5), weights, offset, start, control,
model = FALSE, x = FALSE, y = FALSE, plot=FALSE, ...)
```

```
pest(x, y, indices, family = gaussian,
tuning = list(lambda=TRUE, specific=FALSE, phi=0.5, grouped.fused=0.5,
elastic=0.5, vs=0.5, spl=0.5), weights, offset, start = NULL,
control = cat_control(), plot=FALSE, ...)
```

```
abc(x, y, indices, family = gaussian, tuning = c("AIC", "BIC"),
weights, offset, start, control = cat_control(), plot=FALSE, ...)
```

Arguments

formula	an object of class formula : a symbolic description of the model to be fitted. See details
data	a data frame, containing the variables in the model
family	a family object describing the error distribution and link function to be used in the model; this can be a character string naming a family function, a family function or the result of a call to a family function, see family for details; currently only gaussian, binomial, poisson, Gamma are working
method	fitting method; one out of "lqa", "AIC" or "BIC"; method "lqa" induces penalized estimation; it employs a PIRLS-algorithm (see Fan and Li, 2001; Oelker and Tutz, 2013). Methods "AIC" and "BIC" employ a forward selection strategy
tuning	a list; tuning parameters for penalized estimation; lambda is the scalar, overall penalty parameter; if lambda is a vector of values, these values are cross-validated; if lambda = TRUE, lambda is cross-validated on log scale between lambda.lower and lambda.upper; see cat_control . If lambda is a vector with the same length as elements in the formula and if specific equals a vector of proper length, the entries of specific are interpreted as specific tuning

	parameters for each entry of the formula. <code>phi</code> , <code>grouped.fused</code> , <code>elastic</code> , <code>vs</code> and <code>sp1</code> are parameters that weigh the terms of some penalties; must be out of interval $(0,1)$; the default 0.5 corresponds to equal weights
<code>weights</code>	an optional weight vector (for the observations)
<code>offset</code>	an optional offset
<code>start</code>	initial values for the PIRLS algorithm for method <code>lqa</code>
<code>control</code>	a list of parameters for controlling the fitting process; if empty, set to <code>cat_control()</code> ; see cat_control
<code>model</code>	for functions <code>gvcm.cat</code> : a logical value indicating whether the employed model frame shall be returned or not
<code>x, y</code>	for function <code>gvcm.cat</code> : logical values indicating whether the response vector and the model matrix used in the fitting process shall be returned or not; for functions <code>pest</code> and <code>abc</code> : <code>y</code> must be a response vector, <code>x</code> a proper coded design matrix
<code>plot</code>	logical; if TRUE, estimates needed to plot coefficient paths are computed
<code>indices</code>	for <code>pest</code> and <code>abc</code> only: the to be used index argument; see function index
<code>...</code>	further arguments passed to or from other methods

Details

A typical [formula](#) has the form $\text{response} \sim 1 + \text{terms}$; where `response` is the response vector and `terms` is a series of terms which specifies a linear predictor. There are some special terms for regularized terms:

- `v(x, u, n="L1", bj=TRUE)`: varying coefficients enter the [formula](#) as `v(x,u)` where `u` denotes the categorical effect modifier and `x` the modified covariate. A varying intercept is denoted by `v(1,u)`. Varying coefficients with categorical effect modifiers are penalized as described in Oelker et. al. 2012. The argument `bj` and the element `phi` in argument `tuning` allow for the described weights.
- `p(u, n="L1")`: ordinal/nominal covariates `u` given as `p(u)` are penalized as described in Gertheiss and Tutz (2010). For numeric covariates, `p(u)` indicates a pure Lasso penalty.
- `grouped(u, ...)`: penalizes a group of covariates with the grouped Lasso penalty of Yuan and Lin (2006); so far, working for categorical covariates only
- `sp(x, knots=20, n="L2")`: implements a continuous `x` covariate non-parametrically as $f(x)$; $f(x)$ is represented by centered evaluations of basis functions (cubic B-splines with number of knots = `knots`); for `n="L2"`, the curvature of $f(x)$ is penalized by a Ridge penalty; see Eilers and Marx (1996)
- `SCAD(u)`: penalizes a covariate `u` with the SCAD penalty by Fan and Li (2001); for categorical covariates `u`, differences of coefficients are penalized by a SCAD penalty, see Gertheiss and Tutz (2010)
- `elastic(u)`: penalizes a covariate `u` with the elastic net penalty by Zou and Hastie (2005); for categorical covariates `u`, differences of coefficients are penalized by the elastic net penalty, see Gertheiss and Tutz (2010)

If the `formula` contains no (varying) intercept, `gvcm.cat` assumes a constant intercept. There is no way to avoid an intercept.

For specials `p` and `v`, there is the special argument `n`: if `n="L1"`, the absolute values in the penalty are replaced by squares of the same terms; if `n="L2"`, the absolute values in the penalty are replaced by quadratic, Ridge-type terms; if `n="L0"`, the absolute values in the penalty are replaced by an indicator for non-zero entries of the same terms.

For methods "AIC" and "BIC", the coefficients are not penalized but selected by a forward selection strategy whenever it makes sense; for special `v(x, u)`, the selection strategy is described in Oelker et. al. 2012; the approach for the other specials corresponds to this idea.

For binomial families the response can also be a success/failure rate or a two-column matrix with the columns giving the numbers of successes and failures.

Function `pest` computes penalized estimates, that is, it implements method "lqa" (PIRLS-algorithm).

Function `abc` implements the forward selection strategy employing AIC/BIC.

Categorical effect modifiers and penalized categorical covariates are dummy coded as required by the penalty. If `x` in `v(x, u)` is binary, it is effect coded (first category refers to -1). Other covariates are coded like given by `getOption`.

There is a summary function: `summary.gvcm.cat`

Value

`gvcm.cat` returns an object of class "gvcm.cat" which inherits from class "glm" which inherits from class "lm". An object of class "gvcm.cat" contains:

<code>coefficients</code>	named vector of coefficients
<code>coefficients.reduced</code>	reduced vector of coefficients; selected coefficients/differences of coefficients are set to zero
<code>coefficients.refitted</code>	refitted vector of coefficients; i.e. maximum likelihood estimate of that model containing selected covariates only; same length as <code>coefficients.reduced</code>
<code>coefficients.oml</code>	maximum likelihood estimate of the full model
<code>residuals</code>	deviance residuals
<code>fitted.values</code>	fitted mean values
<code>rank</code>	degrees of freedom model; for <code>method="lqa"</code> estimated by the trace of the generalized head matrix; for methods "AIC", "BIC" estimated like default in <code>glm.fit</code>
<code>family</code>	the <code>family</code> object used
<code>linear.predictors</code>	linear fit on link scale
<code>deviance</code>	scaled deviance
<code>aic</code>	a version of Akaike's Information Criterion; minus twice the maximized log-likelihood plus twice the rank. For binomial and Poisson families the dispersion is fixed at one. For a gaussian family the dispersion is estimated from the residual deviance, and the number of parameters is the rank plus one.
<code>null.deviance</code>	the deviance for the null model, comparable with deviance; the null model includes a non-varying intercept only

iter	number of iterations
weights	working weights of the final iteration
df.residual	the residual degrees of freedom/degrees of freedom error; computed like rank
df.null	the residual degrees of freedom for the null model
converged	logical; fulfills the PIRLS-algorithm the given convergence conditions?
boundary	logical; is the fitted value on the boundary of the attainable values?
offset	the offset vector used
control	the value of the control argument used
contrasts	the contrasts used
na.action	information returned by <code>model.frame</code> on the special handling of NAs; currently always <code>na.omit</code>
plot	in principle, a list containing two matrixes needed for different types of plots: if input option <code>plot=TRUE</code> , the first matrix contains estimates needed to plot coefficient paths; if <code>lambda</code> was cross-validated, the second matrix contains the cross-validation scores
tuning	a list, employed tuning parameters; if <code>lambda</code> was cross-validated, the optimal value is returned
indices	used index argument; see function <code>index</code>
number.selectable.parameters	number of coefficients that could be selected
number.removed.parameters	number of actual removed coefficients
x.reduction	a matrix; transforms model frame <code>x</code> into its reduced version; e.g. needed for refitting
beta.reduction	a matrix; transforms the coefficients into its reduced version
call	the matched call
formula	the <code>formula</code> supplied
terms	the <code>terms</code> object used
data	the data argument
x, y	if requested, the model matrix/the response vector
model	if requested, the model frame
xlevels	a record of the levels of the factors used in fitting
bootstrap.errors	experimental
method	same as input argument <code>method</code>

In addition, non-empty fits will have components `qr`, `R` and `effects` relating to the final weighted linear fit.

Note

Please note that the functions `gvcn.cat`, `pest` and the fitting procedure for penalized estimation `gvcn.cat.fit` are organized like the functions `glm/glm.fit` whenever possible. This was done to avoid mistakes and to provide a well-known structure.

Author(s)

Margret-Ruth Oelker (<margret.oelker@stat.uni-muenchen.de>)

References

- Eilers, P. H. C. and B. D. Marx (1996). Flexible smoothing with b-splines and penalties. *Statist. Sci.* 11 (2), 89-121.
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- Zou, H. and T. Hastie (2005). Regularization and variable selection via the Elastic Net. *R. Stat. Soc. Ser. B Stat. Methodol.* 67 (2), 301-320.

See Also

Functions [index](#), [cat_control](#), [plot.gvcm.cat](#), [predict.gvcm.cat](#), [simulation](#)

Examples

```
## example for function simulation()
covariates <- list(x1=list("unif", c(0,2)),
                  x2=list("unif", c(0,2)),
                  x3=list("unif", c(0,2)),
                  u=list("multinom",c(0.3,0.4,0.3), "nominal")
                  )
true.f <- y ~ 1 + v(x1,u) + x2
true.coefs <- c(0.2, 0.3,.7,.7, -.5)
data <- simulation(400, covariates, NULL, true.f, true.coefs , binomial(), seed=456)
## example for function gvcm.cat()
f <- y ~ v(1,u) + v(x1,u) + v(x2,u)
m1 <- gvcm.cat(f, data, binomial(), plot=TRUE, control=cat_control(lambda.upper=19))
summary(m1)
## example for function predict.gvcm.cat
newdata <- simulation(200, covariates, NULL, true.f, true.coefs , binomial(), seed=789)
prediction <- predict.gvcm.cat(m1, newdata)
## example for function plot.gvcm.cat
plot(m1)
plot(m1, type="score")
plot(m1, type="coefs")
```

gvc <code>m.cat</code> -internal	<i>Internal Function of gvc<code>m.cat</code>()</i>
----------------------------------	---

Description

For internal use only.

See Also

Function [gvc`m.cat`](#)

gvc <code>m.cat.flex</code>	<i>Regularized Effects with Flexible Smoothing Parameters</i>
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Description

The function fits the same models with the same approximation as in [gvc`m.cat`](#) but the choice of the tuning parameter `lambda` for the penalty differs: instead of weighting the penalty terms and choosing on global tuning parameter based on (generalized) cross-validation methods that again rely on the converged model, [gvc`m.cat.flex`](#) estimates several penalty parameteres `lambda_i` by linking the local quadratic approximation of [gvc`m.cat`](#) with the fantastic methods implemented in the package [mgcv](#). This is why the arguments of [gvc`m.cat`](#) and [gvc`m.cat.flex`](#) differ. [gvc`m.cat.flex`](#) is not as well-developed as [gvc`m.cat`](#).

Usage

```
gvcm.cat.flex(whichCoefs, intercept = TRUE, data, family = gaussian(), method = "REML",
  tuning = NULL, indexNrCoefs, indexPenNorm, indexPenA, indexPenWeight,
  control = list(c=1e-05, epsilon=1e-07, gama=35, maxi=1500, nu=.5))
```

Arguments

whichCoefs	vector with covariates (as characters)
intercept	logical
data	a data frame, with named and coded covariates
family	a family object describing the error distribution and link function to be used in the model; see family for details; everyl family that is compatible with gam is working
method	see gam
tuning	for function gam : argument <code>sp</code>
indexNrCoefs	vector with number of coefficients per covariate
indexPenNorm	vector with norm of the employed penalty (as.character)
indexPenA	list with the penalty matrices <code>A_j</code> for each covariate <code>j</code>
indexPenWeight	list, possible weights for the penalty terms (each entry is a vector)
control	a list of parameters for controlling the fitting process; must be NULL or contain all named elements

Details

The local quadratic approximation are linked to the methods of [mgcv](#) by alternating the update of the penalty and the update of the PIRLS algorithm/estimating the tuning parameters `lambda_i` via [mgcv](#). Therefore, [gvcm.cat.flex](#) can be slow (but will be faster than [gvcm.cat](#) for the most part).

Value

A [gamObject](#).

See Also

Function [gvcm.cat](#).

Examples

```
## Not run:
# compare gvcm.cat.flex and gvcm.cat for Lasso-type penalties:
n <- 100
ncov <- 7
set.seed(123)
X <- matrix(rnorm(n*ncov, sd=5), ncol=ncov)
coefs <- rpois(ncov + 1, 2)
y <- cbind(1, X)
data <- as.data.frame(cbind(y, X))
names(data) <- c("y", paste("x", 1:ncov, sep=""))

m1 <- gvcm.cat.flex(
  whichCoefs = paste("x", 1:ncov, sep=""),
  data=data,
  indexNrCoefs=rep(1, ncov),
  indexPenNorm=rep("L1", ncov),
  indexPenA=list(1,1,1,1,1,1,1),
  indexPenWeight=list(1,1,1,1,1,1,1)
)

m2 <- gvcm.cat(y ~ 1 + p(x1) + p(x2) + p(x3) + p(x4) + p(x5) + p(x6) + p(x7),
  data=data, tuning=list(lambda=m1$sp, specific=TRUE), start=rep(1, 8))

rbind(m1$coefficients, m2$coefficients)

# Lasso-type fusion penalty with gvcm.cat.flex
n <- 100
ncat <- 8
set.seed(567)
X <- t(rmultinom(n, 1, rep(1/ncat, ncat)))[, -1]
coefs <- c(rpois(1, 2), sort(rpois(ncat-1, 1)))

y <- cbind(1, X)
data <- as.data.frame(y)
data$x1 <- X
names(data) <- c("y", "x1")
```

```

A <- a(1:(ncat-1), ncat-2)

m3 <- gvcm.cat.flex(
  whichCoefs = c("x1"),
  data = data,
  indexNrCoefs = c(ncat-1),
  indexPenNorm = c("L1"),
  indexPenA = list(A),
  indexPenWeight = list(rep(1, ncol(A))),
  tuning = 100 # fixed and large - in order to demonstrate the fusion of the coefficients
)
m3$coefficients

## End(Not run)

```

index	<i>Functions to Build Design Matrices and Indices for Function gvcm.cat()</i>
-------	---

Description

design() builds design matrices for function [gvcm.cat](#); index() computes indices with information about the terms of the formula.

Usage

```

design(formula, data)

index(dsgn, data = data, formula = formula)

```

Arguments

formula	an object of class "formula"; see gvcm.cat
data	a data frame; see gvcm.cat
dsgn	value of function design()

Details

Function index returns a matrix with one indicator vector per row. The columns refer to the elements of the formula (same order). The indicator/indices are:

- index1 : gives the number of coefficients belonging to each term in the formula. An entry is 1 if the according term is metric, it equals the number of the coded variable's categories, if the variable is a factor. If a continuous variable is modified by a factor u the entry equals the number of u's categories
- index2 : indicates varying coefficients. An entry is 0 if the according coefficient is not varying, it is -1 if the according coefficient is nominal, 1 if it is ordinal

- `index2b`: conforms to indicator b_j in Oelker et. al. 2012
- `index3` : indicates penalized covariates $p(u)$. An entry is 0 if the according covariate is not penalized, it is -1 if the according covariate is nominal, 1 if it is ordinal or metric
- `index4` : indicates penalized covariates $grouped(u)$. An entry is 0 if the according covariate is not penalized, it is -1 if the according covariate is nominal, 1 if it is ordinal or metric
- `index5` : experimental
- `index6` : indicates penalized covariates sp
- `index7` : indicates penalized covariates SCAD. An entry is 0 if the according covariate is not penalized, it is -1 if the according covariate is nominal, 1 if it is ordinal or metric
- `index8` : indicates penalized covariates `elastic`. An entry is 0 if the according covariate is not penalized, it is -1 if the according covariate is nominal, 1 if it is ordinal or metric
- `index9` : experimental

Value

<code>X</code>	the model matrix
<code>Terms</code>	the according terms.object
<code>m</code>	the model frame
<code>int</code>	either 0, indicating that the intercept is varying, or 1 indicating that the intercept is constant
<code>formula</code>	sorted version of the given formula, index vectors will refer to this formula
<code>a matrix</code>	value of function <code>index</code>

References

Oelker, M.-R., J. Gertheiss and G. Tutz (2012). Regularization and model melection with categorial predictors and effect modifiers in generalized linear models. *Department of Statistics at the University of Munich: Technical Report 122*.

See Also

Functions [pest](#), [abc](#)

Examples

```
## example for function simulation()
covariates <- list(x1=list("unif", c(0,2)),
                 x2=list("unif", c(0,2)),
                 x3=list("unif", c(0,2)),
                 u=list("multinom",c(0.3,0.4,0.3), "nominal")
                 )
true.f <- y ~ 1 + v(x1,u) + x2
true.coefs <- c(0.2, 0.3,.7,.7, -.5)
data <- simulation(400, covariates, NULL, true.f, true.coefs , binomial(), seed=456)
## example for function index()
f <- y ~ v(1,u) + v(x1,u) + v(x2,u)
```

```
dsgn <- design(f, data)
index(dsgn, data)
```

plot.gvcm.cat

Plot Method for gvcm.cat Objects

Description

Function to visualize a [gvcm.cat](#) object.

Usage

```
## S3 method for class 'gvcm.cat'
plot(x, accuracy = 2, type = "path", individual = FALSE,
      xlim, ylim, main = NULL, indent = 0, color = TRUE, xscale = "lambda",
      label = TRUE, intercept = TRUE, ...)
```

Arguments

x	a gvcm.cat object; for type="path", a gvcm.cat object with value plot unequal NA is required
accuracy	integer; number of digits being compared when setting coefficients equal/to zero for plotting
type	one out of "path", "score", "coefs"; defines the type of the plot
individual	logical; for type="path" and type="coefs" only; for type="path", it indicates whether the paths of all coefficients shall be plotted into one common figure (default) or in an individual figure per covariate; paths of single covariates can be selected by giving a vector containing the covariates (as characters and as given in the formula, e.g.: individual.paths=c("v(1,u)", "v(x1,u1)")) for type="coefs", the default is one plot per covariate. individual allows to select single covariates.
xlim	the x limits (x1, x2) of the plot
ylim	the y limits (y1, y2) of the plot
main	title of the plot
indent	numeric; if larger zero, coefficient names printed on top of each other are adjusted
color	logical; if FALSE, lines are gray and dotted/dashed
xscale	for type="path" only; if xscale="lambda", the x-axis is scaled as $1 - \lambda / \lambda_{max}$; if xscale="beta", the scale of the x-axis is the scaled L1 norm of the penalized coefficients.
label	omits additional information printed in the plot, if FALSE
intercept	for type="coefs" and type="path" only; if FALSE, for type="path", the path of the intercept is not plotted; if FALSE, for type="coefs", intercept is not added to smooth functions
...	further arguments passed to or from other methods

Details

Default option `type="path"` delivers a graphic with the coefficient paths between 0 (= maximal penalization) and 1 (= no penalization). Maximal penalization is defined by the minimal penalty parameter `lambda` that sets all penalized coefficients to zero (to constant relating to the intercept and assured `.intercept = TRUE`). Minimal penalization means no penalization at all, i.e. `lambda = 0`. Of course the minimal penalty parameter causing maximal penalization depends on how selection and clustering of coefficients is defined (see function [gvcm.cat](#) and [cat_control](#)). Coefficients belonging to one covariate are plotted in the same color, coefficients that are not modified are plotted as dashed lines. Paths are drawn by connecting steps estimates related to different values of `lambda`, see [cat_control](#).

Option `type="score"` plots the cross-validation score (depending on criterion in [cat_control](#)) as a function of penalty parameter `lambda` and marks the chosen penalty parameter as a dotted line. Option `type="coefs"` plots the penalized coefficients whenever possible. So far, there is no plot for methods "AIC" and "BIC".

Value

A plot.

See Also

Function [gvcm.cat](#)

Examples

```
## see example for function gvcm.cat
```

predict.gvcm.cat	<i>Predict Method for gvcm.cat Fits</i>
------------------	---

Description

Obtains predictions from a fitted `gvcm.cat` object.

Usage

```
## S3 method for class 'gvcm.cat'
predict(object, newdata, type = "link", ...)
```

Arguments

object	a fitted object of class gvcm.cat
newdata	a data frame in which to look for variables with which to predict
type	the type of prediction required. The default is on the scale of the linear predictors; the alternative "response" is on the scale of the response variable. Thus for a binomial model the default predictions are of log-odds (probabilities on logit scale) and <code>type = "response"</code> gives the predicted probabilities
...	further arguments passed to or from other methods

Details

Observations containing [NAs](#) are always omitted.

Value

<code>fit</code>	predictions
<code>fit.refitted</code>	predictions assuming refitted coefficients
<code>fit.oml</code>	predictions assuming maximum likelihood estimates
<code>na.action</code>	information returned by <code>model.frame</code> on the special handling of NAs ; currently always <code>na.omit</code>

See Also

Function [gvcm.cat](#)

Examples

```
## see example for function gvcm.cat
```

simulation	<i>Simulates data with categorical covariates</i>
------------	---

Description

Simulates data with categorical covariates/categorical effect modifiers

Usage

```
simulation(n, covariates, correlation = NULL, formula, coefficients,
family, sd = 1, seed = rpois(1, 2348) * rnorm(1))
```

Arguments

<code>n</code>	number of observations; must be large enough, so that all categories of all factor variables exist and therefore vector coefficients fits
<code>covariates</code>	description of the covariates and effect modifiers included in the model; format: <code>list(name of variable 1 = list("distribution", c(parameters), "level of measurement"</code>
<code>correlation</code>	optional matrix, specifies the correlation of Gaussian covariates
<code>formula</code>	formula like in gvcm.cat (all variables contained in <code>formula</code> must be defined in <code>covariates</code>)
<code>coefficients</code>	true parameter vector
<code>family</code>	a <code>family</code> object; currently only gaussian, binomial, poisson, Gamma
<code>sd</code>	if <code>family = gaussian</code> , standard deviation of response; if <code>family = Gamma</code> the rate parameter like in rgamma
<code>seed</code>	specifies the to be used seed

Details

Remarks on covariates:

- all parameterizations like default in [Distributions](#).
- possible distributions of covariates (required as characters), their parameters (required as vectors) and constraints (in parentheses):
 - beta : shape1 (>0), shape2 (>0)
 - exp : rate (>0)
 - gamma : shape (>0)
 - lnorm : mean , sd (>0)
 - multinom: vector of the categories' probabilities (all elements must be >0, sum over all elements must be 1)
 - norm : mean, sd (>0)
 - pois : lambda (>0)
 - unif : min, max
- level of measurement is only needed for distribution = "multinom", must be "nominal" or "ordinal".
- If any, the covariates' correlation is specified by argument `correlation`. Correlations are defined for Gaussian covariates only. Matrix correlation refers to these covariates according to the order they are listed in `covariates`. So that the dimensions of correlation must fit to the number of normal distributed variables in `covariates`.

Value

A data frame containing all specified covariates (even if they are not included in [formula](#)) and the response (named `y`)

See Also

Function [gvcm.cat](#)

Examples

```
## example function simulation
covariates <- list(x1=list("unif", c(0,2)),
                  x2=list("unif", c(0,2)),
                  x3=list("unif", c(0,2)),
                  u=list("multinom",c(0.3,0.4,0.3), "nominal")
                  )
true.f <- y ~ 1 + v(x1,u) + x2
true.coefs <- c(0.2, 0.3,.7,.7, -.5)
data <- simulation(400, covariates, NULL, true.f, true.coefs , binomial(), seed=456)
```

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