

# Package ‘holiglm’

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**Type** Package

**Title** Holistic Generalized Linear Models

**Version** 0.2.1

**Description**

Holistic generalized linear models (HGLMs) extend generalized linear models (GLMs) by enabling the possibility to add further constraints to the model. The 'holiglm' package simplifies estimating HGLMs using convex optimization.

**Depends** R (>= 3.5.0), ROI.plugin.ecos

**Imports** slam, checkmate, MASS, SuppDists, ROI (>= 0.3-0)

**Suggests** ROI.plugin.mosek, ROI.plugin.scs, tinytest (>= 1.0.0)

**License** GPL-3

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**Author** Benjamin Schwendinger [aut, cre],  
Florian Schwendinger [aut],  
Laura Vana [aut]

**Maintainer** Benjamin Schwendinger <benjaminschwe@gmail.com>

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 holiglml-package

*holiglml: Holistic Generalized Linear Models*


---

## Description

The holistic generalized linear models package simplifies estimating GLMs under constraints by making use of convex optimization.

## Author(s)

**Maintainer:** Benjamin Schwendinger <benjaminschwe@gmail.com>

Authors:

- Florian Schwendinger
- Laura Vana

---

active\_coefficients     *Obtain all Active Coefficients*

---

**Description**

The function returns a logical vector which is TRUE for all active (i.e., non-zero) coefficients in the fitted model and FALSE otherwise.

**Usage**

```
active_coefficients(object, ...)  
  
acoef(object, ...)
```

**Arguments**

object            an object inheriting from "hglm" or "hglm.fit" from which the active coefficients obtained from.  
...               optional arguments currently ignored.

**Value**

a logical vector giving the active coefficients.

**Examples**

```
dat <- rhglm(100, c(1, 2, -3, 4, 5, -6))  
fit <- hglm(y ~ ., constraints = k_max(3), data = dat)  
active_coefficients(fit)
```

---

agg\_binomial            *Aggregate Binomial Data*

---

**Description**

A simple function for aggregating binomial data, from a form where y contains only 0 and 1 and X could contain duplicated rows, into a format where y is the matrix of counted successes and failures and X does not contain duplicates. If X contains factor variables, the model matrix corresponding to X will be returned.

**Usage**

```
agg_binomial(formula, data, as_list = TRUE)
```

**Arguments**

formula	a formula object defining the aggregation.
data	a data.frame to be aggregated.
as_list	a logical giving if the return value should be a list. If FALSE the return value is a data.frame.

**Value**

A list (or data.frame) containing aggregated binomial data with counted successes and failures.

**Examples**

```
set.seed(12345)
data <- data.frame(y = rbinom(50, 1, 0.7),
                  a = factor(sample(c(1, 2), 50, TRUE)),
                  b = factor(sample(c(1, 2, 3), 50, TRUE)))
agg_binomial(y ~ ., data)
```

---

as.OP.hglm_model	<i>Convert to OP</i>
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---

**Description**

Convert an object of class hglm\_model to ROI::OP.

**Usage**

```
## S3 method for class 'hglm_model'
as.OP(x)
```

**Arguments**

x	an object inheriting from "hglm_model".
---	---

**Value**

A **ROI** object of class "OP".

---

bike

*Bike Sharing Dataset*

---

### Description

This data set contains the daily count of rented bikes from the the Capital Bikeshare system in Washington D.C., USA, for the years 2011 and 2012. The dataset is already prepared (correct types + factor encodings) for model building.

### Format

A data.frame of dimension 731 x 12 containing daily data related to related bikes.

**dteday** a date vector giving the date of the rental.

**season** a factor with levels 'spring', 'summer', 'fall' and 'winter'.

**year** a factor with levels '2011' and '2012'.

**mnth** a factor with levels 'Jan', 'Feb', 'Mar', 'Apr', 'May', 'Jun', 'Jul', 'Aug', 'Sep', 'Oct', 'Nov' and 'Dec'.

**holiday** a boolean vector indicating if the day is a holiday.

**weathersit** a factor with levels 'good', 'neutral', 'bad' and 'very bad' giving the weather situation.

**temp** a numeric vector containing max-normalized temperature in Celsius with 41 as maximum.

**atemp** a numeric vector containing max-normalized feeling temperature in Celsius with 50 as maximum.

**hum** a numeric vector containing max-normalized humidity with 100 as maximum.

**windspeed** a numeric vector containing max-normalized windspeed with 67 as maximum.

**cnt** an integer vector containing counts of rented bikes.

### Source

<https://www.ics.uci.edu/~mlearn/MLRepository.html>

### References

Fanaee-T, Hadi. (2013). Bike Sharing Dataset. UCI Machine Learning Repository.

### Examples

```
data("bike")
hglm(formula = cnt ~ ., data=bike, family="poisson")
```

---

cov_matrix	<i>Construct Covariance matrix</i>
------------	------------------------------------

---

**Description**

Utility function for constructing covariance matrices based on a simple triplet format ([simple\\_triplet\\_matrix](#)).

**Usage**

```
cov_matrix(k, i, j, v)
```

**Arguments**

k	an integer giving the number of rows and columns of the constructed covariance matrix.
i	an integer vector giving the row indices.
j	an integer vector giving the row indices.
v	a numeric vector giving the corresponding values.

**Value**

A dense matrix of covariances.

**Examples**

```
cov_matrix(5, c(1, 2), c(2, 3), c(0.8, 0.9))
```

---

group_equal	<i>Group Equal Constraint</i>
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---

**Description**

Forces all covariates in the specified group to have the same coefficient.

**Usage**

```
group_equal(vars)
```

**Arguments**

vars	a vector specifying the indices or names of the covariates to which the constraint shall be applied.
------	--

**Value**

A holistic generalized model constraint, object inheriting from class "hg1mc".

**See Also**

Other Constraint-Constructors: [group\\_inout\(\)](#), [group\\_sparsity\(\)](#), [include\(\)](#), [k\\_max\(\)](#), [linear\(\)](#), [lower\(\)](#), [pairwise\\_sign\\_coherence\(\)](#), [rho\\_max\(\)](#), [sign\\_coherence\(\)](#), [upper\(\)](#)

**Examples**

```
dat <- rhglm(100, c(1, 2, 3, 4, 5, 6))
constraints <- group_equal(vars = c("x1", "x3"))
hglm(y ~ ., constraints = constraints, data = dat)
```

---

group\_inout

*In-Out Constraint*


---

**Description**

Forces coefficients of the covariates in the specified group to be either all zero or all nonzero.

**Usage**

```
group_inout(vars)
```

**Arguments**

**vars** a vector specifying the indices or names of the covariates to which the constraint shall be applied.

**Value**

A holistic generalized model constraint, object inheriting from class "hglm".

**See Also**

Other Constraint-Constructors: [group\\_equal\(\)](#), [group\\_sparsity\(\)](#), [include\(\)](#), [k\\_max\(\)](#), [linear\(\)](#), [lower\(\)](#), [pairwise\\_sign\\_coherence\(\)](#), [rho\\_max\(\)](#), [sign\\_coherence\(\)](#), [upper\(\)](#)

**Examples**

```
dat <- rhglm(100, c(1, 2, 3, 4, 5, 6))
constraints <- group_inout(c("x1", "x2", "x3"))
hglm(y ~ ., constraints = constraints, data = dat)
```

---

group_sparsity	<i>Group Sparsity Constraint</i>
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---

### Description

Constraint which restricts the number of covariates selected from a specific group.

### Usage

```
group_sparsity(vars, k = 1L)
```

### Arguments

vars	a vector specifying the indices or names of the covariates to which the group constraint shall be applied.
k	an integer giving the maximum number of covariates to be included in the model from the specified group.

### Value

A holistic generalized model constraint, object inheriting from class "hglm".

### See Also

Other Constraint-Constructors: [group\\_equal\(\)](#), [group\\_inout\(\)](#), [include\(\)](#), [k\\_max\(\)](#), [linear\(\)](#), [lower\(\)](#), [pairwise\\_sign\\_coherence\(\)](#), [rho\\_max\(\)](#), [sign\\_coherence\(\)](#), [upper\(\)](#)

### Examples

```
dat <- rhglm(100, c(1, 2, 0, 4, 5, 0))
constraints <- group_sparsity(c("x1", "x2", "x5"), 1L)
hglm(y ~ ., constraints = constraints, data = dat)
```

---

hglm	<i>Fitting Holistic Generalized Linear Models</i>
------	---

---

### Description

Fit a generalized linear model under holistic constraints.



**Usage**

```
hglm(  
  formula,  
  family = gaussian(),  
  data,  
  constraints = NULL,  
  weights = NULL,  
  scaler = c("auto", "center_standardization", "center_minmax", "standardization",  
            "minmax", "off"),  
  scale_response = NULL,  
  big_m = 100,  
  solver = "auto",  
  control = list(),  
  dry_run = FALSE  
)
```

```
holiglm(  
  formula,  
  family = gaussian(),  
  data,  
  constraints = NULL,  
  weights = NULL,  
  scaler = c("auto", "center_standardization", "center_minmax", "standardization",  
            "minmax", "off"),  
  scale_response = NULL,  
  big_m = 100,  
  solver = "auto",  
  control = list(),  
  dry_run = FALSE  
)
```

```
hglm_seq(  
  k_seq,  
  formula,  
  family = gaussian(),  
  data,  
  constraints = NULL,  
  weights = NULL,  
  scaler = c("auto", "center_standardization", "center_minmax", "standardization",  
            "minmax", "off"),  
  big_m = 100,  
  solver = "auto",  
  control = list()  
)
```

**Arguments**

formula	an object of class "formula" giving the symbolic description of the model to be fitted.
family	a description of the error distribution and link function to be used in the model.
data	a <code>data.frame</code> or <code>matrix</code> giving the data for the estimation.
constraints	a list of 'HGLM' constraints stored in a list of class "lohglm". Use <code>NULL</code> to turn off constraints.
weights	an optional vector of 'prior weights' to be used for the estimation.
scaler	a character string giving the name of the scaling function (default is "auto") to be employed for the covariates. This typically does not need to be changed.
scale_response	a boolean whether the response shall be standardized or not. Can only be used with family <code>gaussian()</code> . Default is <code>TRUE</code> for family <code>gaussian()</code> and <code>FALSE</code> for other families.
big_m	an upper bound for the coefficients, needed for the big-M constraint. Required to inherit from "hglm". Currently constraints created by <code>group_sparsity()</code> , <code>group_inout()</code> , <code>include()</code> and <code>group_equal()</code> use the big-M value specified here.
solver	a character string giving the name of the solver to be used for the estimation.
control	a list of control parameters passed to <code>ROI_solve</code> .
dry_run	a logical; if <code>TRUE</code> the model is not fit but only constructed.
k_seq	an integer vector giving the values of <code>k_max</code> for which the model should be estimated.

**Value**

An object of class "hglm" inheriting from "glm".

**Examples**

```
dat <- rhglm(100, c(1, 2, -3, 4, 5, -6))
hglm(y ~ ., constraints = NULL, data = dat)
# estimation without constraints
hglm(y ~ ., constraints = NULL, data = dat)
# estimation with an upper bound on the number of coefficients to be selected
hglm(y ~ ., constraints = k_max(3), data = dat)
# estimation without intercept
hglm(y ~ . - 1, data = dat)
```



**Arguments**

model	a 'HGLM' model (object of class "hglm_model").
constraints	a list of 'HGLM' constraints stored in a list of class "lohglm".
big_m	an upper bound for the coefficients, needed for the big-M constraint. Required to inherit from "hglm". Currently constraints created by group_sparsity(), group_inout(), include() and group_equal() use the big-M set here.
solver	a character string giving the name of the solver to be used for the estimation.
control	a list of control parameters passed to ROI_solve.
dry_run	a logical if TRUE the model is not fit but only constructed.

**Value**

an object of class "hglm\_fit" inheriting from "glm".

**Examples**

```
dat <- rhglm(100, c(1, 2, -3, 4, 5, -6))
x <- model.matrix(y ~ ., data = dat)
model <- hglm_model(x, y = dat[["y"]])
fit <- hglm_fit(model, constraints = k_max(3))
```

---

hglm\_model

*Create a HGLM Model*


---

**Description**

Create a HGLM model object.

**Usage**

```
hglm_model(
  x,
  y,
  family = gaussian(),
  weights = NULL,
  frame = NULL,
  solver = "auto"
)
```

**Arguments**

x	a numeric matrix giving the design matrix.
y	a vector giving the response variables.
family	a description of the error distribution and link function to be used in the model.
weights	an optional vector of 'prior weights' to be used for the estimation.
frame	an optional model frame object.
solver	a character string giving the name of the solver to be used for the estimation.

**Value**

An object of class "hglm\_model".

**Examples**

```
dat <- rhglm(100, c(1, 2, -3, 4, 5, -6))
x <- model.matrix(y ~ ., data = dat)
hglm_model(x, y = dat[["y"]])
```

---

include

*Include Constraint*

---

**Description**

Ensures that all covariates specified by vars have nonzero coefficients.

**Usage**

```
include(vars)
```

**Arguments**

vars            an integer vector specifying the indices for covariates which have to be in the model.

**Value**

A holistic generalized model constraint, object inheriting from class "hglm".

**See Also**

Other Constraint-Constructors: [group\\_equal\(\)](#), [group\\_inout\(\)](#), [group\\_sparsity\(\)](#), [k\\_max\(\)](#), [linear\(\)](#), [lower\(\)](#), [pairwise\\_sign\\_coherence\(\)](#), [rho\\_max\(\)](#), [sign\\_coherence\(\)](#), [upper\(\)](#)

**Examples**

```
dat <- rhglm(100, c(1, 2, 3, 4, 5, 6))
constraints <- include(vars = c("x1", "x3"))
hglm(y ~ ., constraints = constraints, data = dat)
```

---

<i>k_max</i>	<i>Constraint on the Number of Covariates</i>
--------------	---

---

### Description

Constraint on the maximum number of covariates to be used in the model.

### Usage

```
k_max(k)
```

### Arguments

*k* an positive integer with  $k \leq k_{max}$  giving the maximum number of covariates to be used in the model.

### Value

A holistic generalized model constraint, object inheriting from class "hglm".

### Note

- If an intercept is used, the upper bound on  $k_{max} + 1$  is given by number of columns of the model matrix.
- If no intercept is used, the upper bound on  $k_{max}$  is given by number of columns of the model matrix.

### See Also

Other Constraint-Constructors: [group\\_equal\(\)](#), [group\\_inout\(\)](#), [group\\_sparsity\(\)](#), [include\(\)](#), [linear\(\)](#), [lower\(\)](#), [pairwise\\_sign\\_coherence\(\)](#), [rho\\_max\(\)](#), [sign\\_coherence\(\)](#), [upper\(\)](#)

### Examples

```
dat <- rhglm(100, c(1, 2, -3, 4, 5, -6))
hglm(y ~ ., constraints = k_max(3), data = dat)
```

---

linear	<i>Linear Constraint</i>
--------	--------------------------

---

**Description**

Linear Constraint

**Usage**

```
linear(L, dir, rhs, on_big_m = FALSE)
```

**Arguments**

L	a named vector or matrix defining the linear constraints on the coefficients of the covariates.
dir	a character vector giving the direction of the linear constraints.
rhs	a numeric vector giving the right hand side of the linear constraint.
on_big_m	a logical indicating if the constraint should be imposed on the big-M related binary variables.

**Value**

A holistic generalized model constraint, object inheriting from class "hglm".

**See Also**

Other Constraint-Constructors: [group\\_equal\(\)](#), [group\\_inout\(\)](#), [group\\_sparsity\(\)](#), [include\(\)](#), [k\\_max\(\)](#), [lower\(\)](#), [pairwise\\_sign\\_coherence\(\)](#), [rho\\_max\(\)](#), [sign\\_coherence\(\)](#), [upper\(\)](#)

**Examples**

```
# vector constraint
beta <- c(1, -2, 3)
dat <- rhglm(100, beta)
constraints <- c(linear(c(x1 = 2, x2 = 1), "==", 0), rho_max(1))
hglm(y ~ ., data = dat, constraints = constraints)

# matrix constraint
dat <- rhglm(100, c(1, -2, 3, 4, 5, 6, 7))
mat <- diag(2)
colnames(mat) <- c("x1", "x5")
constraints <- c(linear(mat, c("==", "=="), c(-1, 3)), rho_max(1))
hglm(y ~ ., data = dat, constraints = constraints)
```

---

lower

*Lower Bound*

---

### Description

Set a lower bound on the coefficients of specific covariates.

### Usage

```
lower(kvars)
```

### Arguments

**kvars** a named vector giving the lower bounds. The names should correspond to the names of the covariates in the model matrix.

### Value

A holistic generalized model constraint, object inheriting from class "hglm".

### See Also

Other Constraint-Constructors: [group\\_equal\(\)](#), [group\\_inout\(\)](#), [group\\_sparsity\(\)](#), [include\(\)](#), [k\\_max\(\)](#), [linear\(\)](#), [pairwise\\_sign\\_coherence\(\)](#), [rho\\_max\(\)](#), [sign\\_coherence\(\)](#), [upper\(\)](#)

### Examples

```
dat <- rhglm(100, c(1, 2, -3, 4, 5, -6))
constraints <- lower(c(x2 = 0, x5 = 1))
hglm(y ~ ., constraints = constraints, data = dat)

# non-negative least squares
dat <- rhglm(100, c(1, 2, -3, 4, 5, -6))
constraints <- lower(setNames(double(5), paste0("x", 1:5)))
hglm(y ~ ., constraints = constraints, data = dat)
```

---

pairwise\_sign\_coherence

*Pairwise Sign Coherence*

---

### Description

Ensures that coefficients of covariates which exhibit strong pairwise correlation have a coherent sign.



**Usage**

```
pairwise_sign_coherence(
  rho = 0.8,
  exclude = "(Intercept)",
  big_m = 100,
  eps = 1e-06,
  use = c("everything", "all.obs", "complete.obs", "na.or.complete",
         "pairwise.complete.obs"),
  method = c("pearson", "kendall", "spearman")
)
```

**Arguments**

rho	a value in the range [0,1] specifying the maximum allowed collinearity between pairs of covariates.
exclude	a character vector giving the names of the covariates to be excluded from the constraint (default is "(Intercept)").
big_m	a double giving the big-M parameter.
eps	a double giving the epsilon for the equal sign constraint. Since most numerical solvers can only handle constraints up to some epsilon, e.g., the constraint $Ax \geq b$ is typically transformed to $ Ax - b  \geq 0$ . By providing an $\text{eps} > 0$ and changing the constraint to $ Ax - b  \geq \text{eps}$ we can ensure $ Ax - b  > 0$ .
use	an optional character string giving a method for computing covariances in the presence of missing values. The parameter is passed to <code>cor</code> , therefore see <code>cor</code> for more information.
method	a character string indicating which correlation coefficient is to be computed. The parameter is passed to <code>cor</code> , therefore see <code>cor</code> for more information.

**Value**

A holistic generalized model constraint, object inheriting from class "hglm".

**References**

Carrizosa E, Olivares-Nadal AV, Ramirez-Cobo P (2020) <doi:10.2436/20.8080.02.95>. Integer constraints for enhancing interpretability in linear regression. SORT-Statistics and Operations Research Transactions.

**See Also**

Other Constraint-Constructors: `group_equal()`, `group_inout()`, `group_sparsity()`, `include()`, `k_max()`, `linear()`, `lower()`, `rho_max()`, `sign_coherence()`, `upper()`

---

`rhglm`*Random HGLM Data*

---

## Description

A simple data generator for testing and example purposes.

## Usage

```
rhglm(  
  n,  
  beta,  
  sigma = diag(length(beta) - 1L),  
  family = gaussian(),  
  truncate_mu = FALSE,  
  as_list = FALSE,  
  ...  
)
```

## Arguments

<code>n</code>	the number of observations to be created.
<code>beta</code>	a numeric vector giving the magnitude of the coefficients (the first element is assumed to be the intercept).
<code>sigma</code>	a positive-definite symmetric matrix giving the covariance structure of the covariates (passed to <code>MASS::mvrnorm</code> ).
<code>family</code>	the family of the inverse link.
<code>truncate_mu</code>	a logical giving if <code>mu</code> should be truncated if necessary.
<code>as_list</code>	a logical (default is <code>FALSE</code> ), if <code>TRUE</code> a <code>list</code> is returned otherwise a <code>data.frame</code> is returned.
<code>...</code>	additional optional parameters. The arguments are passed to the random variables generating function of the response.

## Value

A `data.frame` (or `list`) containing the generated data.

## Examples

```
rhglm(10, 1:5)
```

rho\_max

*Constraint on the Pairwise Correlation of Covariates***Description**

Constraint which ensures that only one covariate out of a pair of covariates with a correlation of at least rho will be included in the final model.

**Usage**

```
rho_max(
  rho = 0.8,
  exclude = "(Intercept)",
  use = c("everything", "all.obs", "complete.obs", "na.or.complete",
         "pairwise.complete.obs"),
  method = c("pearson", "kendall", "spearman")
)
```

**Arguments**

rho	a value in the range [0,1] specifying, the maximum allowed collinearity between pairs of covariates.
exclude	variables to be excluded form the pairwise correlation constraints (default is "(Intercept)").
use	an optional character string giving a method for computing co-variances in the presence of missing values. The parameter is passed to <a href="#">cor</a> , therefore see <a href="#">cor</a> for more information.
method	a character string indicating which correlation coefficient is to be computed. See <a href="#">cor</a> for more information.

**Value**

A holistic generalized model constraint, object inheriting from class "hglm".

**See Also**

Other Constraint-Constructors: [group\\_equal\(\)](#), [group\\_inout\(\)](#), [group\\_sparsity\(\)](#), [include\(\)](#), [k\\_max\(\)](#), [linear\(\)](#), [lower\(\)](#), [pairwise\\_sign\\_coherence\(\)](#), [sign\\_coherence\(\)](#), [upper\(\)](#)

**Examples**

```
beta <- 1:3
Sigma <- cov_matrix(k = length(beta) - 1L, 1, 2, 0.9)
dat <- rhglm(100, beta, sigma = Sigma)
hglm(y ~ ., constraints = rho_max(0.8), data = dat)
```

---

sign_coherence	<i>Sign Coherence Constraint</i>
----------------	----------------------------------

---

**Description**

Constraint which ensures that the coefficients of the specified covariates have a coherent sign.

**Usage**

```
sign_coherence(vars, big_m = 100, eps = 1e-06)
```

**Arguments**

vars	a character vector giving the names of the covariates the constraint should be applied to.
big_m	a double giving the big-M parameter.
eps	a double giving the epsilon used to ensure that the constraint holds.

**Value**

A holistic generalized model constraint, object inheriting from class "hglm".

**See Also**

Other Constraint-Constructors: [group\\_equal\(\)](#), [group\\_inout\(\)](#), [group\\_sparsity\(\)](#), [include\(\)](#), [k\\_max\(\)](#), [linear\(\)](#), [lower\(\)](#), [pairwise\\_sign\\_coherence\(\)](#), [rho\\_max\(\)](#), [upper\(\)](#)

**Examples**

```
dat <- rhglm(100, c(1, -2, 3, 4, 5, 6))
constraints <- sign_coherence(c("x1", "x3"))
hglm(y ~ ., constraints = constraints, data = dat)
```

---

solution.hglm	<i>Extract Solution</i>
---------------	-------------------------

---

**Description**

The solution of the underlying optimization problem, can be accessed via the method 'solution'.

**Usage**

```
## S3 method for class 'hglm'
solution(
  x,
  type = c("primal", "dual", "aux", "psd", "msg", "objval", "status", "status_code"),
  force = FALSE,
  ...
)
```

**Arguments**

`x` an object of type 'hglm'.

`type` a character giving the name of the solution to be extracted.

`force` a logical to control the return value in the case that the status code is equal to 1 (i.e. something went wrong). By default force is FALSE and a solution is only provided if the status code is equal to 0 (i.e. success). If force is TRUE the status code is ignored and solutions are returned also where the solver signaled an issue.

... further arguments passed to or from other methods.

**Value**

the extracted solution.

---

upper	<i>Upper Bound</i>
-------	--------------------

---

**Description**

Set a upper bound on the coefficient of specific covariates.

**Usage**

```
upper(kvars)
```

**Arguments**

`kvars` a named vector giving the upper bounds. The names should correspond to the names of the covariates in the model matrix.

**Value**

A holistic generalized model constraint, object inheriting from class "hglm".

**See Also**

Other Constraint-Constructors: [group\\_equal\(\)](#), [group\\_inout\(\)](#), [group\\_sparsity\(\)](#), [include\(\)](#), [k\\_max\(\)](#), [linear\(\)](#), [lower\(\)](#), [pairwise\\_sign\\_coherence\(\)](#), [rho\\_max\(\)](#), [sign\\_coherence\(\)](#)

**Examples**

```
dat <- rhglm(100, c(1, 2, -3, 4, 5, -6))
constraints <- upper(c(x1 = 0, x4 = 1))
hglm(y ~ ., constraints = constraints, data = dat)
```

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