

# Package ‘HGLD’

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**Type** Package

**Title** Fit the Hurdle Generalized Lambda Distribution

**Version** 0.1.0

**Author** Diego Marcondes

**URL** <http://github.com/dmarcondes/HGLD>

**BugReports** <https://github.com/dmarcondes/HGLD/issues>

**Maintainer** Diego Marcondes <diegormmarcondes@gmail.com>

**Description** Functions to perform the Hurdle Generalized Lambda Distribution Regression and to fit the Hurdle Generalized Lambda Distribution to data. The functions of this package are modifications of the functions of packages gamlss, GLDEX and GLDReg, so that most of the credit for the functions of this package must go to the authors and maintainers of these three packages.

**License** GPL-3

**Imports** ggplot2 (>= 2.2.1),  
gridExtra (>= 2.1.1),  
grid (>= 3.3),  
gamlss.dist (>= 5.0-2),  
cluster (>= 2.0.6)

**Depends** R (>= 3.3),  
gamlss (>= 5.0-2),  
GLDEX (>= 2.0.0.5),  
GLDreg (>= 1.0.7)

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 6.0.1

**Reference** Marcondes, D.; Peixoto, C.; Maia, A. C.; A Survey of a Hurdle Model for Heavy-Tailed Data Based on the Generalized Lambda Distribution. (2017) arxiv1712.02183

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dhgld

*The Hurdle Generalized Lambda Distribution Family*

## Description

Density of the Hurdle Generalized Lambda Distribution.

## Usage

```
dhgld(x, mixture = FALSE, lambda1, lambda2 = NULL, prob = NULL,
      param = "fmkl", inverse.eps = 1e-08, max.iterations = 500)
```

## Arguments

<code>x</code>	Vector of data.
<code>mixture</code>	Whether to give the density of a mixture of HGLDs.
<code>lambda1</code>	A vector of length 5 with the five parameters of the HGLD, or of the first HGLD if <code>mixture = TRUE</code> .
<code>lambda2</code>	A vector of length 4 with the four parameters of the second HGLD if <code>mixture = TRUE</code> .
<code>prob</code>	The cluster parameter (probability) for the mixture of HGLDs.
<code>param</code>	"fmkl" or "rs".
<code>inverse.eps</code>	Accuracy of calculation for the numerical determination of $F(x)$ , defaults to $1e-8$ .
<code>max.iterations</code>	Maximum number of iterations in the numerical determination of $F(x)$ , defaults to 500.

## Details

If the parametrization of the RS or FMKL HGLD is not acceptable, the function returns NA. This function is based on the [GLDEX](#) package.

## Value

The probability density of the continuous part of the HGLD.

## References

Marcondes, D.; Peixoto, C.; Maia, A. C.; A Survey of a Hurdle Model for Heavy-Tailed Data Based on the Generalized Lambda Distribution. (2017) *arxiv1712.02183*

Su, S.; Fitting Single and Mixture of Generalized Lambda Distributions to Data via Discretized and Maximum Likelihood Methods: GLDEX in R. (2007), Journal of Statistical Software: \*21\* 9.

## Examples

```
library(ggplot2)
{ggplot(data.frame(x = seq(-3,1,0.01)),aes(x = x)) +
  stat_function(fun = function(x) dhgld(x = x,
                                     lambda1 = c(0.48,0.509,-0.000369,-0.0002483,-0,00039,
                                     param = "rs")))}

#mixture
lambda1 <- c(0.230,0.3514,-0.4472,-0.374,-0.3108)
lambda2 <- c(9.624,-1.227,-0.629,-0.8515)
{ggplot(data.frame(x = seq(-10,20,0.25)),aes(x = x)) +
  stat_function(fun = function(x) dhgld(x = x,mixture = TRUE,lambda1 = lambda1,
  lambda2 = lambda2,prob = 0.47954,param = "rs"))}
```

---

diag.hgld

---

*Diagnostic for the Hurdle Generalized Lambda Distribution*


---

## Description

Diagnostic plots and measures of goodness-of-fit for a Hurdle Generalized Lambda Distribution.

## Usage

```
diag.hgld(fit, facet = FALSE, facet.breaks, facet.labels, facet.ncol,
  trace = TRUE, no.test = 1000, len = floor(0,9 * length(fit$data[fit$data
  != 0])), alpha = 0,05, plotKS = TRUE, KS = TRUE)
```

## Arguments

fit	An object of class <a href="#">fit.hgld</a> .
facet	Whether the plots must be faceted for better visualization.
facet.breaks	The breaks in which to facet the data. Must be the endpoints of the intervals.
facet.labels	The labels of the categories given by the facet breaks.
facet.ncol	The number of columns for the facet plot.
trace	Whether a progress bar must be printed in order to trace the algorithm.
no.test	Total number of KS tests required.
len	Number of data to sample at each KS test.
alpha	Significance level of KS test.
plotKS	Whether to plot the KS resample test result within each plot.
KS	Whether the resample KS test must be performed to the non-zero values.

## Details

The diagnostics techniques are applied to the non-zero data values. Returns the qq-plot and the quantile plot between the data and the theoretical fitted HGLD. Also returns a table comparing sample moments with theoretical moments. A Kolmogorov-Simornov resample test is performed and the percentage of the times that the null hypotheses, i.e., goodness-of-fit, is not rejected is displayed. All diagnostics are performed for both the RS and FMKL HGLD.

## Value

qqRS	<a href="#">ggplot</a> qq-plot for the RS HGLD.
qqFMKL	<a href="#">ggplot</a> qq-plot for the FMKL HGLD.
quantRS	<a href="#">ggplot</a> quantile plot for the RS HGLD.
quantFMKL	<a href="#">ggplot</a> quantile plot for the FMKL HGLD.
moments	Moments comparison for the GLD fitted to the non-zero data for both parametrizations. These are not the moments of the HGLD, but are instead the moments of the GLD fitted to the non-zero data values.
KS	Percentage of no rejection for the KS resample test.

## References

Marcondes, D.; Peixoto, C.; Maia, A. C.; A Survey of a Hurdle Model for Heavy-Tailed Data Based on the Generalized Lambda Distribution. (2017) *arxiv1712.02183*

Su, S. Fitting Single and Mixture of Generalized Lambda Distributions to Data via Discretized and Maximum Likelihood Methods: GLDEX in R. (2007), Journal of Statistical Software: \*21\* 9.

## Examples

```
set.seed(100)
data <- healthcare[sample(1:nrow(healthcare), 50), ]
fit <- fit.hgld(data$log_expense)
d <- diag.hgld(fit, facet = FALSE, plotKS = FALSE)

#mixture
set.seed(100)
data <- c(rcauchy(20, location = 10), rep(0, 10), rcauchy(20))
fit <- fit.hgld(data = data, mixture = TRUE)
d <- suppressWarnings(diag.hgld(fit))
```

---

fit.hgld

---

*Fit the Hurdle Generalized Lambda Distribution*


---

## Description

Fit the Hurdle Generalized Lambda Distribution to a dataset by the Numerical Maximum Likelihood Method.

## Usage

```
fit.hgld(data, mixture = FALSE, clustering.m = clara, threshold = NULL,
  leap1 = 3, leap2 = 3, fun1 = "runif.sobol", fun2 = "runif.sobol",
  rs.leap = 3, fmk1.leap = 3, rs.init = c(-1,5, 1,5),
  fmk1.init = c(-0,25, 1,5), FUN = "runif.sobol", no = 10000)
```

## Arguments

data	A vector of data.
mixture	Whether a mixture of HGLD must be fitted.
clustering.m	Clustering method used in classifying the dataset into two parts when fitting a mixture of HGLD. Valid arguments include clara, fanny and pam from the cluster library, or threshold, if the data must be divided by a threshold. Default is clara. Or a logical vector specifying how data should be split. See <a href="#">fun.auto.bimodal.pml</a> for more details.
threshold	The threshold to divide the data if <i>clustering.m</i> = "threshold".
leap1	Scrambling (0,1,2,3) for the sobol sequence for the first distribution fit when fitting a mixture of HGLD. See scrambling/leap argument for <a href="#">runif.sobol</a> , <a href="#">runif.halton</a> or <a href="#">QUnif</a> of the <a href="#">GLDEX</a> package.
leap2	Scrambling (0,1,2,3) for the sobol sequence for the second distribution fit when fitting a mixture of HGLD. See scrambling/leap argument for <a href="#">runif.sobol</a> , <a href="#">runif.halton</a> or <a href="#">QUnif</a> of the <a href="#">GLDEX</a> package.
fun1	A character string of either "runif.sobol" (default), "runif.halton" or "QUnif" for the first distribution fit when fitting a mixture of HGLD. See <a href="#">fun.auto.bimodal.pml</a> for more details.
fun2	A character string of either "runif.sobol" (default), "runif.halton" or "QUnif" for the second distribution fit when fitting a mixture of HGLD. See <a href="#">fun.auto.bimodal.pml</a> for more details.
rs.leap	Scrambling (0,1,2,3) for the sobol sequence for the RS generalized lambda distribution fit. See scrambling/leap argument for <a href="#">runif.sobol</a> , <a href="#">runif.halton</a> or <a href="#">QUnif</a> of the <a href="#">GLDEX</a> package. See <a href="#">fun.data.fit.ml</a> for more details.
fmk1.leap	Scrambling (0,1,2,3) for the sobol sequence for the FMKL generalized lambda distribution fit. See scrambling/leap argument for <a href="#">runif.sobol</a> , <a href="#">runif.halton</a> or <a href="#">QUnif</a> of the <a href="#">GLDEX</a> package. See <a href="#">fun.data.fit.ml</a> for more details.
rs.init	Initial values (lambda3 and lambda4) for the RS generalized lambda distribution. See <a href="#">fun.data.fit.ml</a> for more details.
fmk1.init	Initial values (lambda3 and lambda4) for the FMKL generalized lambda distribution. See <a href="#">fun.data.fit.ml</a> for more details.
FUN	A character string of either "runif.sobol" (default), "runif.halton" or "QUnif". See <a href="#">fun.data.fit.ml</a> for more details.
no	Number of initial random values to find the best initial values for optimization. See <a href="#">fun.data.fit.ml</a> for more details.

## Details

Given a dataset, estimate by the Numerical Maximum Likelihood Method the five parameters of the HGLD. Fit both the RS and the FMKL parametrizations. Also fit a mixture of HGLDs.

**Value**

par	The estimate of the HGLD five parameters for both the RS and FMKL parametrizations if <i>mixture</i> = <i>FALSE</i> . Otherwise present the estimation of ten parameters: the zero probability mass, the four parameters of each GLD and the clustering parameter p.
data	The data used in the fit.
mixture	Whether a mixture of HGLD was fitted.

**References**

Marcondes, D.; Peixoto, C.; Maia, A. C.; A Survey of a Hurdle Model for Heavy-Tailed Data Based on the Generalized Lambda Distribution. (2017) *arxiv1712.02183*

Su, S.; Fitting Single and Mixture of Generalized Lambda Distributions to Data via Discretized and Maximum Likelihood Methods: GLDEX in R.(2007), Journal of Statistical Software: \*21\* 9.

**Examples**

```
set.seed(100)
data <- healthcare[sample(1:nrow(healthcare), 30), ]
fit <- fit.hgld(data$log_expense)

#mixture
set.seed(100)
data <- c(rcauchy(20, location = 10), rep(0, 10), rcauchy(20))
fit2 <- fit.hgld(data = data, mixture = TRUE)
```

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healthcare	<i>Healthcare dataset</i>
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**Description**

Dataset containing the healthcare expense of 129,257 customers of a Brazilian healthcare company between 2006 and 2009.

**Usage**

```
healthcare
```

**Format**

An object of class `tbl_df` (inherits from `tbl`, `data.frame`) with 289456 rows and 8 columns.

**Details**

The expenses are in Reais (Brazilian currency) and were deflated to the January 2006 value. In order to fit the models of the paper in the references it is necessary to truncate the expanses at R\$ 100.

**Value**

ID	The ID of the customer.
sex	The sex of the customer.
age	The age of the customer on the considered year.
expense	The healthcare expense of the customer on the considered year.
log_expense	The logarithm of the healthcare expense of the customer on the considered year.
year	The considered year.
previous_expense	The healthcare expense of the customer on the previous year.
log_previous_expense	The logarithm of the healthcare expense of the customer on the previous year.

**References**

Marcondes, D.; Peixoto, C.; Maia, A. C.; A Survey of a Hurdle Model for Heavy-Tailed Data Based on the Generalized Lambda Distribution. (2017) *arxiv1712.02183*

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phgld	<i>The Hurdle Generalized Lambda Distribution Family</i>
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**Description**

Accumulated probability function of the Hurdle Generalized Lambda Distribution.

**Usage**

```
phgld(q, mixture = FALSE, lambda1, lambda2 = NULL, prob = NULL,
      param = "fmkl", inverse.eps = 1e-08, max.iterations = 500)
```

**Arguments**

q	Vector of quantiles.
mixture	Whether to give the accumulated probability function of a mixture of HGLDs.
lambda1	A vector of length 5 with the five parameters of the HGLD, or of the first HGLD if mixture = TRUE.
lambda2	A vector of length 4 with the four parameters of the second HGLD if mixture = TRUE.
prob	The cluster parameter for the mixture of HGLDs.
param	"fmkl" or "rs".
inverse.eps	Accuracy of calculation for the numerical determination of F(x), defaults to 1e-8.
max.iterations	Maximum number of iterations in the numerical determination of F(x), defaults to 500.

## Details

If the parametrization of the RS or FMKL GLD is not acceptable, the function returns NA. This function is based on the [GLDEX](#) package.

## Value

The acumulated probability function of the HGLD.

## References

Marcondes, D.; Peixoto, C.; Maia, A. C.; A Survey of a Hurdle Model for Heavy-Tailed Data Based on the Generalized Lambda Distribution. (2017) *arxiv1712.02183*

Su, S.; Fitting Single and Mixture of Generalized Lambda Distributions to Data via Discretized and Maximum Likelihood Methods: GLDEX in R. (2007), Journal of Statistical Software: \*21\* 9.

## Examples

```
library(ggplot2)
{ggplot(data.frame(x = seq(2,8,0.25)),aes(x = x)) +
 stat_function(fun = function(x) phgld(q = x,lambda1 = c(0.54,4.561,0.0191,0.00925,0.0217)
                                     param = "rs"))}

#mixture
lambda1 <- c(0.230,0.3514,-0.4472,-0.3748,-0.3108)
lambda2 <- c(9.624,-1.227,-0.6290,-0.8515)
{ggplot(data.frame(x = seq(-5,15,0.25)),aes(x = x)) +
 stat_function(fun = function(x) phgld(q = x,mixture = TRUE,lambda1 = lambda1,
                                     lambda2 = lambda2,prob = 0.479,param = "rs"))}
```

---

plot.fit.hgld

*Plot the Hurdle Generalized Lambda Distribution*

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## Description

Density plot of the fit given by an object of class [fit.hgld](#).

## Usage

```
## S3 method for class 'fit.hgld'
plot(x, histogram = TRUE, hcolor = "white",
     hfill = "black", bins = 50, dcolor = c("black", "red"),
     dtype = c("solid", "solid"), xlab = "Data", ylab = "Density", ...)
```

## Arguments

x	A <a href="#">fit.hgld</a> object.
histogram	Logical. Whether the estimated density must be superimposed by the data histogram.
hcolor	Color of the histogram.
hfill	Color to fill the histogram.



bins	Number of histogram bins.
dcolor	Color of the RS and FMKL density plot.
dtype	Type of the RS and FMKL density line.
xlab	Label of the x-axis.
ylab	Label of the y-axis.
...	Arguments to be passed to methods.

### Details

The density may be plotted by itself or superimposed by the data histogram. Plot only the non-zero data values.

### Value

RS	<a href="#">ggplot</a> plot of the fitted RS GLD.
FMKL	<a href="#">ggplot</a> plot of the fitted FMKL GLD.
RF	<a href="#">ggplot</a> plot of the fitted RS and FMKL GLD on the same plot.

### References

Marcondes, D.; Peixoto, C.; Maia, A. C.; A Survey of a Hurdle Model for Heavy-Tailed Data Based on the Generalized Lambda Distribution. (2017) *arxiv1712.02183*

### Examples

```
set.seed(100)
data <- healthcare[sample(1:nrow(healthcare), 50), ]
fit <- fit.hgld(data$log_expense)
plot(fit)

#mixture
set.seed(100)
data <- c(rcauchy(20, location = 10), rep(0, 10), rcauchy(20))
fit <- fit.hgld(data = data, mixture = TRUE)
plot(fit)
```

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plot.reg.hgld	<i>Predict density plot of a Hurdle Generalized Lambda Regression</i>
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### Description

Predict density plot of a Hurdle Generalized Lambda Regression.

### Usage

```
## S3 method for class 'reg.hgld'
plot(x, newvalues, name = row.names(newvalues),
     color = NULL, xlab = "Data",
     xlim = c(min(x$NZdata[[all.vars(x$loc.formula)[1]]]),
              max(x$NZdata[[all.vars(x$loc.formula)[1]]])), title = "", ...)
```

**Arguments**

<code>x</code>	An object of class <code>reg.hglD</code> .
<code>newvalues</code>	A data frame with the new values of the covariates. Column names must match the ones given in formulas <i>loc.formula</i> and <i>zero.formula</i> .
<code>name</code>	A vector with the names of the new values profiles. Default is the row names of the <code>newvalues</code> data frame.
<code>color</code>	The color of each density curve. Must have one color for each row of <code>newvalues</code> .
<code>xlab</code>	Label of the x-axis.
<code>xlim</code>	A vector with the limits of the x-axis.
<code>title</code>	Legend title.
<code>...</code>	Arguments to be passed to methods.

**Details**

Given an object of class `reg.hglD` and new values for the covariates, returns the density function of the fitted HGLD, given the covariates. The contrast on the *newvalues* data frame must be same of the *data* used in the `reg.hglD` object. All density curves are plotted in the same plot.

**Value**

`plot` `ggplot` density plot for the given new values

**References**

Marcondes, D.; Peixoto, C.; Maia, A. C.; A Survey of a Hurdle Model for Heavy-Tailed Data Based on the Generalized Lambda Distributions. (2017) *arxiv1712.02183*

**Examples**

```
set.seed(100)
tmp <- na.omit(healthcare)
data <- tmp[sample(1:nrow(tmp), 50), ]
formula <- log_expense ~ age + sex + log_previous_expense
reg <- suppressWarnings(reg.hglD(data = data, zero.formula = formula, loc.formula = formula,
                                full = FALSE, param = "rs"))
newvalues <- tmp[sample(1:nrow(tmp), 5), c(2, 3, 5, 8)]
plot(reg, newvalues)
```

qhglD

*The Hurdle Generalized Lambda Distribution Family***Description**

Quantile function of a Hurdle Generalized Lambda Distribution.

**Usage**

```
qhglD(p, mixture = FALSE, lambda1, lambda2 = NULL, prob = NULL,
      param = "fmkl", trace = FALSE, inverse.eps = 1e-08,
      max.iterations = 500)
```

**Arguments**

<code>p</code>	Vector of probabilities.
<code>mixture</code>	Whether to give the quantile function of a mixture of HGLDs.
<code>lambda1</code>	A vector of length 5 with the five parameters of the HGLD, or of the first HGLD if <code>mixture = TRUE</code> .
<code>lambda2</code>	A vector of length 4 with the four parameters of the second HGLD if <code>mixture = TRUE</code> .
<code>prob</code>	The cluster parameter for the mixture HGLD.
<code>param</code>	"fmkl" or "rs".
<code>trace</code>	Whether a progress bar must be printed in order to trace the algorithm.
<code>inverse.eps</code>	Accuracy of calculation for the numerical determination of $F(x)$ , defaults to $1e-8$ .
<code>max.iterations</code>	Maximum number of iterations in the numerical determination of $F(x)$ , defaults to 500.

**Details**

The HGLD must be non-negative, otherwise the quantile is only approximate. If the parametrization of the RS or FMKL GLD is not acceptable, the function returns NA. This function is based on the [GLDEX](#) package.

**Value**

The quantile function of a HGLD.

**References**

Marcondes, D.; Peixoto, C.; Maia, A. C.; A Survey of a Hurdle Model for Heavy-Tailed Data Based on the Generalized Lambda Distribution. (2017) *arxiv1712.02183*

Su, S.; Fitting Single and Mixture of Generalized Lambda Distributions to Data via Discretized and Maximum Likelihood Methods: GLDEX in R. (2007), Journal of Statistical Software: \*21\* 9.

**Examples**

```
qhglD(p = seq(0.05, 1, 0.05), lambda1 = c(0.540, 3.561, 0.019, 0.009, 0.022), param = "rs")

#mixture
qhglD(p = seq(0.05, 1, 0.05), mixture = TRUE, lambda1 = c(0.1, 8, 1, 3, 6), lambda2 = c(0.3, 10, 3, 6),
      prob = 0.5, param = "fmkl")
```

qhglld.reg

*Predict quantiles of a Hurdle Generalized Lambda regression***Description**

Quantile function of a Hurdle Generalized Lambda Regression.

**Usage**

```
qhglld.reg(p, reg, newvalues, l0 = TRUE, location = TRUE)
```

**Arguments**

p	Vector of probabilities.
reg	An object of class <a href="#">reg.hgld</a> .
newvalues	A data frame with the new values of the covariates. Column names must match the ones given in formulas <i>loc.formula</i> and <i>zero.formula</i> of the <a href="#">reg.hgld</a> object.
l0	Whether to return the lambda0 of each profile.
location	Whether to return the location of each profile.

**Details**

Given an object of class [reg.hgld](#), a percentile and new values for the covariates, it returns the quantile of the fitted HGLD, given the covariates. The fitted HGLD must be non-negative. The contrast on the *newvalues* data frame must be the same of the *data* used to fit the [reg.hgld](#) regression.

**Value**

The quantile of the new values, according to the fitted HGLD regression.

**References**

Marcondes, D.; Peixoto, C.; Maia, A. C.; A Survey of a Hurdle Model for Heavy-Tailed Data Based on the Generalized Lambda Distribution. (2017) *arxiv1712.02183*

**Examples**

```
set.seed(100)
tmp <- na.omit(healthcare)
data <- tmp[sample(1:nrow(tmp), 50), ]
formula <- log_expense ~ age + sex + log_previous_expense
reg <- suppressWarnings(reg.hgld(data = data, zero.formula = formula, loc.formula = formula,
                                full = FALSE, param = "rs"))
newvalues <- tmp[sample(1:nrow(tmp), 5), c(2, 3, 8)]
qhglld.reg(p = seq(0.05, 0.95, 0.1), reg, newvalues)
```

reg.hgld

*Fit a Hurdle Generalized Lambda Distribution Regression model***Description**

Fit a Hurdle Generalized Lambda Distribution Regression model to a dataset.

**Usage**

```
reg.hgld(data, zero.formula, loc.formula, full = FALSE, param = "fmkl",
         maxit = 20000, init = NULL, alpha = 0.05, n.simu = 1000,
         plotKS = TRUE, h.bins = 50)
```

**Arguments**

data	A dataset containing the variables of the model.
zero.formula	A symbolic expression of the model to be fitted to the hurdle parameter of the distribution.
loc.formula	A symbolic expression of the model to be fitted to the location of the distribution.
full	Whether a simulation method must be applied to derive a confidence interval for the location regression coefficients.
param	"fmkl" or "rs".
maxit	Maximum number of iterations for numerical optimization.
init	Choose a different set of initial values to start the optimization process. This can either be full set of parameters including GLD parameter estimates, or it can just be the coefficient estimates of the regression model.
alpha	Significance level of the Confidence Interval for the GLD regression.
n.simu	Number of times to repeat the simulation runs, defaults to 1000.
plotKS	Whether to plot the KS resample test result within each plot.
h.bins	Number of bins for the GLD Regression normalized quantiles residuals histogram.

**Details**

Given a dataset, estimate by the Numerical Maximum Likelihood method the regression coefficients of the model and the five parameters of the error GLD. The regression coefficients that model the location of the distribution are estimated by the functions [GLD.lm](#) and [GLD.lm.full](#) of package [GLDreg](#). The regression coefficients that model the hurdle parameter of the distribution are estimated by function [gamlss](#).

**Value**

coefficients	The estimated coefficients of the HGLD regression.
Zplot	A function that generates the four diagnostic plots of the logistic regression.
Zres.sumarry	Summary of the normalised quantile residuals of the logistic regression.
Zfit	Normalised quantile residuals versus fitted values for the logistic regression.
Zindex	Normalised quantile residuals versus index for the logistic regression.



rhgld

*The Hurdle Generalized Lambda Distribution Family***Description**

Sample from a Hurdle Generalized Lambda Distribution.

**Usage**

```
rhgld(n, mixture = FALSE, lambda1, lambda2 = NULL, prob = NULL,
      param = "fmkl")
```

**Arguments**

n	Number of observations to be generated.
mixture	Whether to sample from a mixture of HGLDs.
lambda1	A vector of length 5 with the five parameters of the HGLD, or of the first HGLD if mixture = TRUE.
lambda2	A vector of length 4 with the four parameters of the second HGLD if mixture = TRUE
prob	The cluster parameter for the mixture HGLD.
param	"fmkl" or "rs".

**Details**

If the parametrization of the RS or FMKL GLD is not acceptable, the function returns NA. This function is based on the [GLDEX](#) package.

**Value**

A sample of a HGLD.

**References**

Marcondes, D.; Peixoto, C.; Maia, A. C.; A Survey of a Hurdle Model for Heavy-Tailed Data Based on the Generalized Lambda Distribution. (2017) *arxiv1712.02183*

Su, S.; Fitting Single and Mixture of Generalized Lambda Distributions to Data via Discretized and Maximum Likelihood Methods: GLDEX in R. (2007), Journal of Statistical Software: \*21\* 9.

**Examples**

```
set.seed(100)
rhgld(n = 1000, lambda1 = c(0.05, 0, 1, 3, 6), param = "fmkl")
rhgld(n = 1000, mixture = TRUE, lambda1 = c(0.05, 0, 1, 3, 6), lambda2 = c(0.3, 2, 3, 6), prob = 0.5,
      param = "fmkl")
```

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