Package 'glogis'

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Title Fitting and Testing Generalized Logistic Distributions

Description Tools for the generalized logistic distribution (Type I,
    also known as skew-logistic distribution), encompassing
    basic distribution functions (p, q, d, r, score), maximum
    likelihood estimation, and structural change methods.

LazyLoad yes

Depends R (>= 2.10.0), zoo

Suggests strucchange, fxregime, lattice

Imports graphics, stats, sandwich

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NeedsCompilation no

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R topics documented:

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breakpoints.glogisfit   *Segmented Fitting of the Generalized Logistic Distribution*

Description

Fitting univariate generalized logisitc distributions (Type I: skew-logistic with location, scale, and shape parameters) to segments of time series data.

Usage

```
## S3 method for class 'glogisfit'
breakpoints(obj, h = 0.15, breaks = NULL, ic = "LWZ",
  hpc = "none", ...)

## S3 method for class 'breakpoints.glogisfit'
refit(object, ...)

## S3 method for class 'breakpoints.glogisfit'
coef(object, log = TRUE, ...)

## S3 method for class 'breakpoints.glogisfit'
fitted(object, type = c("mean", "variance", "skewness"), ...)

## S3 method for class 'breakpoints.glogisfit'
confint(object, parm = NULL, level = 0.95, breaks = NULL,
  meat. = NULL, ...)
```

Arguments

obj	an object of class [glogisfit](#).
h	numeric. Minimal segment size either given as fraction relative to the sample size or as an integer giving the minimal number of observations in each segment.
breaks	integer specifying the maximal number of breaks to be calculated. By default the maximal number allowed by h is used.
ic	character specifying the default information criterion that should be employed for selecting the number of breakpoints. Default is "LWZ" (Liu-Wu-Zidek criterion, a modified BIC). Instead the classic "BIC" can be used.
hpc	a character specifying the high performance computing support. Default is "none", can be set to "foreach".
object	an object of class breakpoints.glogisfit as returned by the [breakpoints](#) method.
log	logical option in coef method indicating whether scale and shape parameters should be reported in logs (default) or the original levels.
type	character specifying which moments of the segmented fitted distribution should be extracted.

parm	integer. Either parm or breaks may be set, see below.
level	numeric. The confidence level to be used.
meat.	function. A function for extracting the meat of a sandwich estimator from a fitted object. By default, the inverse of bread is used, i.e., a correctly specified model is assumed.
...	arguments passed to methods.

Details

To test whether sequences (typically time series) of observations follow the same generalized logistic distribution, the stability of the parameters can be tested. If there is evidence for parameter instability, breakpoints can be estimated to find segments with stable parameters.

The methods from the **strucchange** and **fxregime** packages are leveraged. For testing, the generalized M-fluctuation tests from **strucchange** can directly be employed using gefp. For breakpoint estimation, the methods documented here provide a user interface to some internal functionality from the **fxregime** packages. They employ the (unexported) workhorse function gbreakpoints which is modeled after breakpoints from the **strucchange** package but employing user-defined estimation methods.

Optional support for high performance computing is available in the breakpoints method based on the **foreach** package for the dynamic programming algorithm. If hpc = "foreach" is to be used, a parallel backend should be registered before. See breakpoints for more information.

Value

breakpoints.glogisfit returns an object of class "breakpoints.glogisfit" that inherits from "gbreakpointsfull".

References

Windberger T, Zeileis A (2014). Structural Breaks in Inflation Dynamics within the European Monetary Union. *Eastern European Economics*, **52**(3), 66–88.

Zeileis A, Shah A, Patnaik I (2010). Testing, Monitoring, and Dating Structural Changes in Exchange Rate Regimes. *Computational Statistics and Data Analysis*, **54**(6), 1696–1706. doi: 10.1016/j.csda.2009.12.005.

See Also

glogisfit, fxregimes, breakpoints

Examples

```
## artifical data with one structural change
set.seed(1071)
x <- c(rglogis(50, -1, scale = 0.5, shape = 3), rglogis(50, 1, scale = 0.5, shape = 1))
x <- zoo(x, yearmon(seq(2000, by = 1/12, length = 100)))

## full sample estimation
gf <- glogisfit(x)
```

```
if(require("strucchange")) {

## structural change testing
gf_scus <- gefp(gf, fit = NULL)
plot(gf_scus, aggregate = FALSE)
plot(gf_scus, functional = meanL2BB)
sctest(gf_scus)
sctest(gf_scus, functional = meanL2BB)


## breakpoint estimation
gf_bp <- breakpoints(gf)
plot(gf_bp)
summary(gf_bp)
breakdates(gf_bp)
coef(gf_bp)
confint(gf_bp)

## fitted model
plot(x)
lines(gf_bp)
lines(fitted(gf_bp, type = "mean"), col = 4)
lines(confint(gf_bp))

}
```

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glogis   *The Generalized Logistic Distribution (Type I: Skew-Logitic)*

Description

Density, distribution function, quantile function and random generation for the logistic distribution
with parameters `location` and `scale`.

Usage

```
dglogis(x, location = 0, scale = 1, shape = 1, log = FALSE)
pglogis(q, location = 0, scale = 1, shape = 1, lower.tail = TRUE, log.p = FALSE)
qglogis(p, location = 0, scale = 1, shape = 1, lower.tail = TRUE, log.p = FALSE)
rglogis(n, location = 0, scale = 1, shape = 1)
sglogis(x, location = 0, scale = 1, shape = 1)
```

Arguments

x, q	vector of quantiles.
p	vector of probabilities.
n	number of observations. If length(n) > 1, the length is taken to be the number required.

```
location, scale, shape
                 location, scale, and shape parameters (see below).
```

log, log.p   logical; if TRUE, probabilities p are given as log(p).

lower.tail   logical; if TRUE (default), probabilities are $P[X \leq x]$, otherwise, $P[X > x]$.

Details

If location, scale, or shape are omitted, they assume the default values of 0, 1, and 1, respectively.

The generalized logistic distribution with location $= \mu$, scale $= \sigma$, and shape $= \gamma$ has distribution function

$$F(x) = \frac{1}{(1 + e^{-(x-\mu)/\sigma})^{\gamma}}$$

.

The mean is given by location + (digamma(shape) - digamma(1)) * scale, the variance by (psigamma(shape, deriv = 1) + psigamma(1, deriv = 1)) * scale^2) and the skewness by (psigamma(shape, deriv = 2) - psigamma(1, deriv = 2)) / (psigamma(shape, deriv = 1) + psigamma(1, deriv = 1))^(3/2)).

[dpq]glogis are calculated by leveraging the [dpq]logis and adding the shape parameter. rglogis uses inversion.

Value

dglogis gives the probability density function, pglogis gives the cumulative distribution function, qglogis gives the quantile function, and rglogis generates random deviates. sglogis gives the score function (gradient of the log-density with respect to the parameter vector).

References

Johnson NL, Kotz S, Balakrishnan N (1995) *Continuous Univariate Distributions*, volume 2. John Wiley & Sons, New York.

Shao Q (2002). Maximum Likelihood Estimation for Generalised Logistic Distributions. *Communications in Statistics – Theory and Methods*, **31**(10), 1687–1700.

Windberger T, Zeileis A (2014). Structural Breaks in Inflation Dynamics within the European Monetary Union. *Eastern European Economics*, **52**(3), 66–88.

Examples

```
## PDF and CDF
par(mfrow = c(1, 2))
x <- -100:100/10
plot(x, dglogis(x, shape = 2), type = "l", col = 4, main = "PDF", ylab = "f(x)")
lines(x, dglogis(x, shape = 1))
lines(x, dglogis(x, shape = 0.5), col = 2)
legend("topleft", c("generalized (0, 1, 2)", "standard (0, 1, 1)",
  "generalized (0, 1, 0.5)"), lty = 1, col = c(4, 1, 2), bty = "n")
plot(x, pglogis(x, shape = 2), type = "l", col = 4, main = "CDF", ylab = "F(x)")
lines(x, pglogis(x, shape = 1))
lines(x, pglogis(x, shape = 0.5), col = 2)

## artifical empirical example
```

```
set.seed(2)
x <- rglogis(1000, -1, scale = 0.5, shape = 3)
gf <- glogisfit(x)
plot(gf)
summary(gf)
```

---

glogisfit   *Fitting the Generalized Logistic Distribution*

Description

Fit a univariate generalized logisitc distribution (Type I: skew-logistic with location, scale, and shape parameters) to a sample of observations.

Usage

```
glogisfit(x, ...)
## Default S3 method:
glogisfit(x, weights = NULL, start = NULL, fixed = c(NA, NA, NA),
  method = "BFGS", hessian = TRUE, ...)
## S3 method for class 'formula'
glogisfit(formula, data, subset, na.action, weights, x = TRUE, ...)

## S3 method for class 'glogisfit'
plot(x, main = "", xlab = NULL, fill = "lightgray",
  col = "blue", lwd = 1, lty = 1, xlim = NULL, ylim = NULL,
  legend = "topright", moments = FALSE, ...)

## S3 method for class 'glogisfit'
summary(object, log = TRUE, breaks = NULL, ...)
## S3 method for class 'glogisfit'
coef(object, log = TRUE, ...)
## S3 method for class 'glogisfit'
vcov(object, log = TRUE, ...)
```

Arguments

x	a vector of observation (may be a ts or zoo time series).
weights	optional numeric vector of weights.
start	optional vector of starting values. The parametrization has to be in terms of location, log(scale), log(shape) where the original parameters (without logs) are as in dglogis. Default is to use c(0, 0, 0) (i.e., standard logistic). For details see below.
fixed	specification of fixed parameters (see description of start). NA signals that the corresponding parameter should be estimated. A standard logistic distribution could thus be fitted via fixed = c(NA, NA, 0).

method	character string specifying optimization method, see [optim](#) for the available options. Further options can be passed to optim through ....
hessian	logical. Should the Hessian be used to compute the variance/covariance matrix? If FALSE, no covariances or standard errors will be available in subsequent computations.
formula	symbolic description of the model, currently only x ~ 1 is supported.
data, subset, na.action	
	arguments controlling formula processing via [model.frame](#).
main, xlab, fill, col, lwd, lty, xlim, ylim	
	standard graphical parameters, see [plot](#) and [par](#).
legend	logical or character specification where to place a legend. legend = FALSE suppresses the legend. See [legend](#) for the character specification.
moments	logical. If a legend is produced, it can either show the parameter estimates (moments = FALSE, default) or the implied moments of the distribution.
object	a fitted glogisfit object.
log	logical option in some extractor methods indicating whether scale and shape parameters should be reported in logs (default) or the original levels.
breaks	interval breaks for the chi-squared goodness-of-fit test. Either a numeric vector of two or more cutpoints or a single number (greater than or equal to 2) giving the number of intervals.
...	arguments passed to methods.

Details

glogisfit estimates the generalized logistic distribution (Type I: skew-logistic) as given by [dglogis](#). Optimization is performed numerically by [optim](#) using analytical gradients. For obtaining numerically more stable results the scale and shape parameters are specified in logs. Starting values are chosen as c(0, 0, 0), i.e., corresponding to a standard (symmetric) logistic distribution. If these fail, better starting values are obtained by running a Nelder-Mead optimization on the original problem (without logs) first.

A large list of standard extractor methods is supplied to conveniently compute with the fitted objects, including methods to the generic functions [print](#), [summary](#), [plot](#) (reusing [hist](#) and [lines](#)), [coef](#), [vcov](#), [logLik](#), [residuals](#), and [estfun](#) and [bread](#) (from the **sandwich** package).

The methods for coef, vcov, summary, and bread report computations pertaining to the scale/shape parameters in logs by default, but allow for switching back to the original levels (employing the delta method).

Visualization employs a histogramm of the original data along with lines for the estimated density.

Further structural change methods for "glogisfit" objects are described in [breakpoints.glogisfit](#).

Value

glogisfit returns an object of class "glogisfit", i.e., a list with components as follows.

coefficients	estimated parameters from the model (with scale/shape in logs, if included),
vcov	associated estimated covariance matrix,

loglik	log-likelihood of the fitted model,
df	number of estimated parameters,
n	number of observations,
nobs	number of observations with non-zero weights,
weights	the weights used (if any),
optim	output from the `optim` call for maximizing the log-likelihood,
method	the method argument passed to the `optim` call,
parameters	the full set of model parameters (location/scale/shape), including estimated and fixed parameters, all in original levels (without logs),
moments	associated mean/variance/skewness,
start	the starting values for the parameters passed to the `optim` call,
fixed	the original specification of fixed parameters,
call	the original function call,
x	the original data,
converged	logical indicating successful convergence of `optim`,
terms	the terms objects for the model (if the `formula` method was used).

References

Shao Q (2002). Maximum Likelihood Estimation for Generalised Logistic Distributions. *Communications in Statistics – Theory and Methods*, **31**(10), 1687–1700.

Windberger T, Zeileis A (2014). Structural Breaks in Inflation Dynamics within the European Monetary Union. *Eastern European Economics*, **52**(3), 66–88.

See Also

dglogis, dlogis, breakpoints.glogisfit

Examples

```
## simple artificial example
set.seed(2)
x <- rglogis(1000, -1, scale = 0.5, shape = 3)
gf <- glogisfit(x)
plot(gf)
summary(gf)

## query parameters and associated moments
coef(gf)
coef(gf, log = FALSE)
gf$parameters
gf$moments
```

HICP	*Harmonised Index of Consumer Prices (1990–2010, OECD)*

Description

Time series data with HICP (Harmonised Index of Consumer Prices) for 21 countries (plus EU) for 1990–2010 as provided by the OECD; and corresponding seasonally adjusted inflation ratios.

Usage

```
data("HICP")

data("hicps")
```

Format

Monthly multiple `"zooreg"` time series with `"yearmon"` index from Jan 1990 (`HICP`) or Feb 1990 (`hicps`) to Dec 2010 for 21 countries (plus EU).

Details

`HICP` contains the raw unadjusted Harmonised Index of Consumer Prices as provided by the OECD from which unadjusted inflation rates can be easily computed (see examples).

As the different countries have rather different seasonal patterns which vary over time (especially in the 2000s), they will typically require seasonal adjustment before modeling. Hence, a seasonally adjusted version of the inflation rate series is provided as `hicps`, where X-12-ARIMA (version 0.3) has been employed for adjusted. An alternative seasonal adjustment can be easily computed use [stl](#) (see examples).

Source

Organisation for Economic Co-operation and Development (OECD)

https://stats.oecd.org/

References

Wikipedia (2010). "Harmonised Index of Consumer Prices – Wikipedia, The Free Encyclopedia." https://en.wikipedia.org/wiki/Harmonised_Index_of_Consumer_Prices, accessed 2010-06-10.

Windberger T, Zeileis A (2014). Structural Breaks in Inflation Dynamics within the European Monetary Union. *Eastern European Economics*, **52**(3), 66–88.

Examples

```
## price series
data("HICP", package = "glogis")

## corresponding raw unadjusted inflation rates (in percent)
hicp <- 100 * diff(log(HICP))

## seasonal adjustment of inflation rates (via STL)
hicps1 <- do.call("merge", lapply(1:ncol(hicp), function(i) {
  z <- na.omit(hicp[,i])
  coredata(z) <- coredata(as.ts(z) - stl(as.ts(z), s.window = 13)$time.series[, "seasonal"])
  z
}))
colnames(hicps1) <- colnames(hicp)

## load X-12-ARIMA adjusted inflation rates
data("hicps", package = "glogis")

## compare graphically for one country (Austria)
plot(hicp[, "Austria"], lwd = 2, col = "lightgray")
lines(hicps1[, "Austria"], col = "red")
lines(hicps[, "Austria"], col = "blue")
legend("topleft", c("unadjusted", "STL", "X-12-ARIMA"), lty = c(1, 1, 1),
  col = c("lightgray", "red", "blue"), bty = "n")

## compare graphically across all countries (via lattice)
if(require("lattice")) {
trellis.par.set(theme = canonical.theme(color = FALSE))
xyplot(merge(hicp, hicps1, hicps),
  screen = names(hicp)[rep(1:ncol(hicp), 3)],
  col = c("lightgray", "red", "blue")[rep(1:3, each = ncol(hicp))],
  lwd = c(2, 1, 1)[rep(1:3, each = ncol(hicp))])
}
```

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