

Package ‘fGarch’

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Title Rmetrics - Autoregressive Conditional Heteroskedastic Modelling

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Depends R (>= 2.6.0), stats, graphics, methods, timeDate, timeSeries, fBasics (>= 2100.78)

Suggests RUnit, Matrix, fastICA, tcltk

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Description Environment for teaching “Financial Engineering and Computational Finance”

NOTE SEVERAL PARTS ARE STILL PRELIMINARY AND MAY BE CHANGED IN THE FUTURE. THIS TYPICALLY INCLUDES FUNCTION AND ARGUMENT NAMES, AS WELL AS DEFAULTS FOR ARGUMENTS AND RETURN VALUES.

LazyLoad yes

LazyData yes

License GPL (>= 2)

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fGarch-package	<i>GARCH Modelling Package</i>
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Description

Package of econometric functions for modelling GARCH processes.

Details

Package:	fGarch
Type:	Package
Version:	270.73
Date:	2008
License:	GPL (>= 2)
Copyright:	(c) 1999-2008 Diethelm Wuertz and Rmetrics Foundation
URL:	http://www.rmetrics.org

GARCH, Generalized Autoregressive Conditional Heteroskedastic, models have become important in the analysis of time series data, particularly in financial applications when the goal is to analyze and forecast volatility.

For this purpose, the family of GARCH functions offers functions for simulating, estimating and

forecasting various univariate GARCH-type time series models in the conditional variance and an ARMA specification in the conditional mean. The function `garchFit` is a numerical implementation of the maximum log-likelihood approach under different assumptions, Normal, Student-t, GED errors or their skewed versions. The parameter estimates are checked by several diagnostic analysis tools including graphical features and hypothesis tests. Functions to compute n-step ahead forecasts of both the conditional mean and variance are also available.

The number of GARCH models is immense, but the most influential models were the first. Beside the standard ARCH model introduced by Engle [1982] and the GARCH model introduced by Bollerslev [1986], the function `garchFit` also includes the more general class of asymmetric power ARCH models, named APARCH, introduced by Ding, Granger and Engle [1993]. The APARCH models include as special cases the TS-GARCH model of Taylor [1986] and Schwert [1989], the GJR-GARCH model of Glosten, Jaganathan, and Runkle [1993], the T-ARCH model of Zakoian [1993], the N-ARCH model of Higgins and Bera [1992], and the Log-ARCH model of Geweke [1986] and Pentula [1986].

There exist a collection of review articles by Bollerslev, Chou and Kroner [1992], Bera and Higgins [1993], Bollerslev, Engle and Nelson [1994], Engle [2001], Engle and Patton [2001], and Li, Ling and McAleer [2002] which give a good overview of the scope of the research.

Time Series Simulation

contains functions to simulate artificial GARCH and APARCH time series processes.

Functions:

<code>garchSpec</code>	Specifies an univariate GARCH time series model,
<code>garchSim</code>	Simulates a GARCH/APARCH process.

Parameter Estimation

contains functions to fit the parameters of GARCH and APARCH time series processes.

Functions:

<code>garchFit</code>	Fits the parameters of a GARCH process,
<code>residuals</code>	Extracts residuals from a fitted 'fGARCH' object,
<code>fitted</code>	Extracts fitted values from a fitted 'fGARCH' object,
<code>volatility</code>	Extracts conditional volatility from a fitted 'fGARCH' object,
<code>coef</code>	Extracts coefficients from a fitted 'fGARCH' object,
<code>formula</code>	Extracts formula expression from a fitted 'fGARCH' object.

Forecasting

contains functions to forecast mean and variance of GARCH and APARCH processes.

Functions:

<code>predict</code>	Forecasts from an object of class 'fGARCH'.
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Standardized Distribution Functions

contains functions to model standardized distribution functions.

Functions:

<code>[dpqr]norm</code>	Normal distribution function,
<code>[dpqr]snorm</code>	Skew Normal distribution function,
<code>[s]normFit</code>	Fits parameters of [skew] Normal distribution,
<code>[dpqr]ged</code>	Generalized Error distribution function,
<code>[dpqr]sged</code>	Skew Generalized Error distribution function,
<code>[s]gedFit</code>	Fits parameters of [skew] Generalized Error distribution,
<code>[dpqr]std</code>	standardized Student-t distribution function,
<code>[dpqr]sstd</code>	Skew standardized Student-t distribution function,
<code>[s]stdFit</code>	Fits parameters of [skew] Student-t distribution,
<code>absMoments</code>	Computes absolute Moments of these distribution.

OX Interface

NOTE: `garchOxFit` is no longer part of `fGarch` package. If you are interested to use, please contact us.

contains a Windows interface to OX.

The function `garchOxFit` interfaces a subset of the functionality of the G@ARCH 4.0 Package written in Ox. G@RCH 4.0 is one of the most sophisticated packages for modelling univariate GARCH processes including GARCH, EGARCH, GJR, APARCH, IGARCH, FIGARCH, FIEGARCH, FIAPARCH and HYGARCH models. Parameters can be estimated by approximate (Quasi-) maximum likelihood methods under four assumptions: normal, Student-t, GED or skewed Student-t errors.

Author(s)

Diethelm Wuertz and Rmetrics Core Team.

`absMoments`

Absolute Moments of GARCH Distributions

Description

Computes absolute Moments of the skew Normal, skew GED, and standardized skew Student-t distributions

Usage

```
absMoments(n, density = c("dnorm", "dged", "dstd"), ...)
```

Arguments

`density` a character string naming the symmetric density function.
`n` the number of absolute Moments.
`...` parameters passed to the density function.

Value

`absMoments` returns a numeric vector of length `n` with the values of the absolute moments of the selected density function.

Author(s)

Diethelm Wuertz for the Rmetrics R-port.

References

Fernandez C., Steel M.F.J. (2000); *On Bayesian Modelling of Fat Tails and Skewness*, Preprint, 31 pages.

Examples

```
## absMoment -
  absMoments(4, "dstd", nu = 4)
```

coef-methods *GARCH Coefficients Methods*

Description

Coefficients methods for GARCH Modelling.

Methods

object = "ANY" Generic function.
object = "fGARCH" Extractor function for coefficients from a fitted GARCH model.
object = "fGARCHSPEC" Extractor function for coefficients from a GARCH specification structure.

Note

`coef` is a generic function which extracts coefficients from objects returned by modeling functions.

Author(s)

Diethelm Wuertz for the Rmetrics R-port.

Examples

```
## garchSpec -
# Use default parameters beside alpha:
spec = garchSpec(model = list(alpha = c(0.05, 0.05)))
spec
coef(spec)

## garchSim -
# Simulate an univariate "timeSeries" series
x = garchSim(spec, n = 200)
x = x[,1]

## garchFit -
fit = garchFit(~ garch(1, 1), data = x)

## coef -
coef(fit)
```

fGARCH-class	<i>Class "fGARCH"</i>
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Description

The class fGARCH represents a model of an heteroskedastic time series process.

Objects from the Class

Objects can be created by calls of the function `garchFit`. This object is a parameter estimate of an empirical GARCH process.

Slots

call: Object of class "call": the call of the `garch` function.

formula: Object of class "formula": a formula object specifying mean and variance equation.

method: Object of class "character": a string denoting the optimization method, by default the returned string is "Max Log-Likelihood Estimation".

data: Object of class "list": a list with one entry named `x`, containing the data of the time series to be estimated, the same as given by the input argument `series`.

fit: Object of class "list": a list with the results from the parameter estimation. The entries of the list depend on the selected algorithm, see below.

residuals: Object of class "numeric": a numeric vector with the residual values.

fitted: Object of class "numeric": a numeric vector with the fitted values.

h.t: Object of class "numeric": a numeric vector with the conditional variances.

sigma.t: Object of class "numeric": a numeric vector with the conditional standard deviations.

title: Object of class "character": a title string.

description: Object of class "character": a string with a brief description.

Methods

- plot** signature(x = "fGARCH", y = "missing"): plots an object of class 'fGARCH'.
- show** signature(object = "fGARCH"): prints an object of class 'fGARCH'.
- summary** signature(object = "fGARCH"): summarizes an object of class 'fGARCH'.
- predict** signature(object = "fGARCH"): forecasts mean and volatility from an object of class 'fGARCH'.
- fitted** signature(object = "fGARCH"): extracts fitted values from an object of class 'fGARCH'.
- residuals** signature(object = "fGARCH"): extracts residuals from an object of class 'fGARCH'.
- volatility** signature(object = "fGARCH"): extracts conditional volatility from an object of class 'fGARCH'.
- coef** signature(object = "fGARCH"): extracts fitted coefficients from an object of class 'fGARCH'.
- formula** signature(x = "fGARCH"): extracts formula expression from an object of class 'fGARCH'.

Author(s)

Diethelm Wuertz and Rmetrics Core Team.

fGARCHSPEC-class *Class "fGARCHSPEC"*

Description

Specification Structure for an univariate GARCH time series model.

Objects from the Class

Objects can be created by calls of the function `garchSpec`. This object specifies the parameters of an empirical GARCH process.

Slots

- call**: Object of class "call": the call of the `garch` function.
- formula**: Object of class "formula": a list with two formula entries for the mean and variance equation.
- model**: Object of class "list": a list with the model parameters.
- presample**: Object of class "matrix": a numeric matrix with presample values.
- distribution**: Object of class "character": a character string with the name of the conditional distribution.
- rseed**: Object of class "numeric": an integer with the random number generator seed.

Methods

show signature(object = "fGARCHSPEC"): prints an object of class 'fGARCHSPEC'.

Note

With Rmetrics Version 2.6.1 the class has been renamed from "garchSpec" to "fGARCHSPEC".

Author(s)

Diethelm Wuertz for the Rmetrics R-port.

fitted-methods

Extract GARCH Model Fitted Values

Description

Extracts fitted values from a fitted GARCH object.

Details

The function extracts the @fitted value slot from an object of class "fGARCH" as returned by the function garchFit.

The class of the returned value depends on the input to the function garchFit who created the object. The returned value is always of the same class as the input object to the argument data in the function garchFit, i.e. if you fit a "timeSeries" object, you will get back from the function fitted also a "timeSeries" object, if you fit an object of class "zoo", you will get back again a "zoo" object. The same holds for a "numeric" vector, for a "data.frame", and for objects of class "ts", "mts".

In contrast, the slot itself returns independent of the class of the data input always a numeric vector, i.e. the function call rslot(object, "fitted") will return a numeric vector.

Methods

object = "ANY" Generic function.

object = "fGARCH" Extractor function for fitted values.

Note

fitted is a generic function which extracts fitted values from objects returned by modeling functions.

Author(s)

Diethelm Wuertz for the Rmetrics R-port.

Examples

```
## Swiss Pension fund Index -
x = as.timeSeries(data(LPP2005REC))

## garchFit -
# Fit LPP40 Bechmark:
fit = garchFit(LPP40 ~ garch(1, 1), data = 100*x, trace = FALSE)
fit

## fitted -
# Fitted values are now a "timeSeries" object:
fitted = fitted(fit)
head(fitted)
class(fitted)

## slot -
# The slot contains a numeric Vector:
fitted = slot(fit, "fitted")
head(fitted)
class(fitted)
```

formula-methods

Extract GARCH Model formula

Description

Extracts formula from a formula GARCH object.

Details

The function extracts the `@formula` expression slot from an object of class "fGARCH" as returned by the function `garchFit`.

Note, the returned formula has always a left hand side. If the argument `data` was an univariate time series and no name was specified to the series, then the left hand side has assigned the name of the data.set. In the multivariate case the rectangular `data` object must always have column names, otherwise the fitting will be stopped and you get the error message

The class of the returned value depends on the input to the function `garchFit` who created the object. The returned value is always of the same class as the input object to the argument `data` in the function `garchFit`, i.e. if you fit a "timeSeries" object, you will get back from the function `fitted` also a "timeSeries" object, if you fit an object of class "zoo", you will get back again a "zoo" object. The same holds for a "numeric" vector, for a "data.frame", and for objects of class "ts", "mts".

In contrast, the slot itself returns independent of the class of the data input always a numeric vector, i.e. the function call `rslot(object, "fitted")` will return a numeric vector.

Methods

object = "ANY" Generic function.

object = "fGARCH" Extractor function for formula expression.

Note

formula is a generic function which extracts the formula expression from objects returned by modeling functions.

Author(s)

Diethelm Wuertz for the Rmetrics R-port.

Examples

```
## garchFit -
fit = garchFit(~garch(1, 1), data = garchSim())

## formula -
formula(fit)

## A Bivariate series and mis-specified formula:
x = garchSim(n = 500)
y = garchSim(n = 500)
z = cbind(x, y)
colnames(z)
class(z)
## Not run:
garchFit(z ~garch(1, 1), data = z, trace = FALSE)

## End(Not run)
# Returns:
# Error in .garchArgsParser(formula = formula, data = data, trace = FALSE) :
#   Formula and data units do not match.

## Doubled column names in data set - formula can't fit:
colnames(z) <- c("x", "x")
z[1:6,]
## Not run:
garchFit(x ~garch(1, 1), data = z, trace = FALSE)

## End(Not run)
# Again the error will be noticed:
# Error in garchFit(x ~ garch(1, 1), data = z) :
#   Column names of data are not unique.

## Missing column names in data set - formula can't fit:
z.mat <- as.matrix(z)
colnames(z.mat) <- NULL
z.mat[1:6,]
## Not run:
```

```

garchFit(x ~ garch(1, 1), data = z.mat, trace = FALSE)

## End(Not run)
# Again the error will be noticed:
# Error in .garchArgsParser(formula = formula, data = data, trace = FALSE) :
#   Formula and data units do not match

```

garchFit

Univariate GARCH Time Series Fitting

Description

Estimates the parameters of an univariate ARMA-GARCH/APARCH process.

Usage

```

garchFit(formula = ~ garch(1, 1), data = dem2gbp,
  init.rec = c("mci", "uev"),
  delta = 2, skew = 1, shape = 4,
  cond.dist = c("norm", "snorm", "ged", "sged", "std", "sstd",
    "snig", "QMLE"),
  include.mean = TRUE, include.delta = NULL, include.skew = NULL,
  include.shape = NULL, leverage = NULL, trace = TRUE,

  algorithm = c("nlminb", "lbfgsb", "nlminb+nm", "lbfgsb+nm"),
  hessian = c("ropt", "rcd"), control = list(),
  title = NULL, description = NULL, ...)

garchKappa(cond.dist = c("norm", "ged", "std", "snorm", "sged", "sstd",
  "snig"), gamma = 0, delta = 2, skew = NA, shape = NA)

```

Arguments

algorithm	a string parameter that determines the algorithm used for maximum likelihood estimation.
cond.dist	a character string naming the desired conditional distribution. Valid values are "dnorm", "dged", "dstd", "dsnorn", "dsged", "dsstd" and "QMLE". The default value is the normal distribution. See Details for more information.
control	control parameters, the same as used for the functions from <code>nlminb</code> , and 'bfgs' and 'Nelder-Mead' from <code>optim</code> .
data	an optional <code>timeSeries</code> or data frame object containing the variables in the model. If not found in <code>data</code> , the variables are taken from <code>environment(formula)</code> , typically the environment from which <code>armaFit</code> is called. If <code>data</code> is an univariate series, then the series is converted into a numeric vector and the name of the response in the formula will be neglected.

<code>delta</code>	a numeric value, the exponent <code>delta</code> of the variance recursion. By default, this value will be fixed, otherwise the exponent will be estimated together with the other model parameters if <code>include.delta=FALSE</code> .
<code>description</code>	a character string which allows for a brief description.
<code>formula</code>	formula object describing the mean and variance equation of the ARMA-GARCH/APARCH model. A pure GARCH(1,1) model is selected when e.g. <code>formula=~garch(1,1)</code> . To specify for example an ARMA(2,1)-APARCH(1,1) use <code>formula = ~arma(2,1)+apaarch(1,</code>
<code>gamma</code>	APARCH leverage parameter entering into the formula for calculating the expectation value.
<code>hessian</code>	a string denoting how the Hessian matrix should be evaluated, either <code>hessian="rcd"</code> , or <code>"ropt"</code> , the default, <code>"rcd"</code> is a central difference approximation implemented in R and <code>"ropt"</code> use the internal R function <code>optimhess</code> .
<code>include.delta</code>	a logical flag which determines if the parameter for the recursion equation <code>delta</code> will be estimated or not. If <code>include.delta=FALSE</code> then the shape parameter will be kept fixed during the process of parameter optimization.
<code>include.mean</code>	this flag determines if the parameter for the mean will be estimated or not. If <code>include.mean=TRUE</code> this will be the case, otherwise the parameter will be kept fixed during the process of parameter optimization.
<code>include.shape</code>	a logical flag which determines if the parameter for the shape of the conditional distribution will be estimated or not. If <code>include.shape=FALSE</code> then the shape parameter will be kept fixed during the process of parameter optimization.
<code>include.skew</code>	a logical flag which determines if the parameter for the skewness of the conditional distribution will be estimated or not. If <code>include.skew=FALSE</code> then the skewness parameter will be kept fixed during the process of parameter optimization.
<code>init.rec</code>	a character string indicating the method how to initialize the mean and variance recursion relation.
<code>leverage</code>	a logical flag for APARCH models. Should the model be leveraged? By default <code>leverage=TRUE</code> .
<code>shape</code>	a numeric value, the shape parameter of the conditional distribution.
<code>skew</code>	a numeric value, the skewness parameter of the conditional distribution.
<code>title</code>	a character string which allows for a project title.
<code>trace</code>	a logical flag. Should the optimization process of fitting the model parameters be printed? By default <code>trace=TRUE</code> .
<code>...</code>	additional arguments to be passed.

Details

"QMLE" stands for Quasi-Maximum Likelihood Estimation, which assumes normal distribution and uses robust standard errors for inference. Bollerslev and Wooldridge (1992) proved that if the mean and the volatility equations are correctly specified, the QML estimates are consistent and asymptotically normally distributed. However, the estimates are not efficient and "the efficiency loss

can be marked under asymmetric ... distributions” (Bollerslev and Wooldridge (1992), p. 166). The robust variance-covariance matrix of the estimates equals the (Eicker-White) sandwich estimator, i.e.

$$V = H^{-1}G'GH^{-1},$$

where V denotes the variance-covariance matrix, H stands for the Hessian and G represents the matrix of contributions to the gradient, the elements of which are defined as

$$G_{t,i} = \frac{\partial l_t}{\partial \zeta_i},$$

where l_t is the log likelihood of the t -th observation and ζ_i is the i -th estimated parameter. See sections 10.3 and 10.4 in Davidson and MacKinnon (2004) for a more detailed description of the robust variance-covariance matrix.

Value

garchFit

returns a S4 object of class "fGARCH" with the following slots:

@call	the call of the <code>garch</code> function.
@formula	a list with two formula entries, one for the mean and the other one for the variance equation.
@method	a string denoting the optimization method, by default the returned string is "Max Log-Likelihood Estimation".
@data	a list with one entry named <code>x</code> , containing the data of the time series to be estimated, the same as given by the input argument <code>series</code> .
@fit	a list with the results from the parameter estimation. The entries of the list depend on the selected algorithm, see below.
@residuals	a numeric vector with the (raw, unstandardized) residual values.
@fitted	a numeric vector with the fitted values.
@h.t	a numeric vector with the conditional variances ($h_t = \sigma_t^2$).
@sigma.t	a numeric vector with the conditional standard deviation.
@title	a title string.
@description	a string with a brief description.

The entries of the @fit slot show the results from the optimization.

Author(s)

Diethelm Wuertz for the Rmetrics R-port,
 R Core Team for the 'optim' R-port,
 Douglas Bates and Deepayan Sarkar for the 'nlminb' R-port,
 Bell-Labs for the underlying PORT Library,
 Ladislav Luksan for the underlying Fortran SQP Routine,
 Zhu, Byrd, Lu-Chen and Nocedal for the underlying L-BFGS-B Routine.

References

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- Nocedal J., Wright S.J. (1999); *Numerical Optimization*, Springer, New York.

Examples

```
## UNIVARIATE TIME SERIES INPUT:
# In the univariate case the lhs formula has not to be specified ...

# A numeric Vector from default GARCH(1,1) - fix the seed:
N = 200
x.vec = as.vector(garchSim(garchSpec(rseed = 1985), n = N)[,1])
garchFit(~ garch(1,1), data = x.vec, trace = FALSE)

# An univariate timeSeries object with dummy dates:
x.timeSeries = dummyDailySeries(matrix(x.vec), units = "GARCH11")
garchFit(~ garch(1,1), data = x.timeSeries, trace = FALSE)

## Not run:
# An univariate zoo object:
x.zoo = zoo(as.vector(x.vec), order.by = as.Date(rownames(x.timeSeries)))
garchFit(~ garch(1,1), data = x.zoo, trace = FALSE)

## End(Not run)

# An univariate "ts" object:
x.ts = as.ts(x.vec)
garchFit(~ garch(1,1), data = x.ts, trace = FALSE)

## MULTIVARIATE TIME SERIES INPUT:
```

```

# For multivariate data inputs the lhs formula must be specified ...

# A numeric matrix binded with dummy random normal variates:
X.mat = cbind(GARCH11 = x.vec, R = rnorm(N))
garchFit(GARCH11 ~ garch(1,1), data = X.mat)

# A multivariate timeSeries object with dummy dates:
X.timeSeries = dummyDailySeries(X.mat, units = c("GARCH11", "R"))
garchFit(GARCH11 ~ garch(1,1), data = X.timeSeries)

## Not run:
# A multivariate zoo object:
X.zoo = zoo(X.mat, order.by = as.Date(rownames(x.timeSeries)))
garchFit(GARCH11 ~ garch(1,1), data = X.zoo)

## End(Not run)

# A multivariate "mts" object:
X.mts = as.ts(X.mat)
garchFit(GARCH11 ~ garch(1,1), data = X.mts)

## MODELING THE PERCENTUAL SPI/SBI SPREAD FROM LPP BENCHMARK:

X.timeSeries = as.timeSeries(data(LPP2005REC))
X.mat = as.matrix(x.timeSeries)
## Not run: X.zoo = zoo(X.mat, order.by = as.Date(rownames(X.mat)))
X.mts = ts(X.mat)
garchFit(100*(SPI - SBI) ~ garch(1,1), data = X.timeSeries)
# The remaining are not yet supported ...
# garchFit(100*(SPI - SBI) ~ garch(1,1), data = X.mat)
# garchFit(100*(SPI - SBI) ~ garch(1,1), data = X.zoo)
# garchFit(100*(SPI - SBI) ~ garch(1,1), data = X.mts)

## MODELING HIGH/LOW RETURN SPREADS FROM MSFT PRICE SERIES:

X.timeSeries = MSFT
garchFit(Open ~ garch(1,1), data = returns(X.timeSeries))
garchFit(100*(High-Low) ~ garch(1,1), data = returns(X.timeSeries))

```

garchFitControl *GARCH Fitting Algorithms and Control*

Description

Estimates the parameters of an univariate GARCH process.

Usage

```
garchFitControl(
```

```

llh = c("filter", "internal", "testing"),
nlminb.eval.max = 2000,
nlminb.iter.max = 1500,
nlminb.abs.tol = 1.0e-20,
nlminb.rel.tol = 1.0e-14,
nlminb.x.tol = 1.0e-14,
nlminb.step.min = 2.2e-14,
nlminb.scale = 1,
nlminb.fscale = FALSE,
nlminb.xscale = FALSE,
sqp.mit = 200,
sqp.mfv = 500,
sqp.met = 2,
sqp.mec = 2,
sqp.mer = 1,
sqp.mes = 4,
sqp.xmax = 1.0e3,
sqp.tolx = 1.0e-16,
sqp.tolc = 1.0e-6,
sqp.tolg = 1.0e-6,
sqp.told = 1.0e-6,
sqp.tols = 1.0e-4,
sqp.rpf = 1.0e-4,
lbfgsb.REPORT = 10,
lbfgsb.lmm = 20,
lbfgsb.pgtol = 1e-14,
lbfgsb.factr = 1,
lbfgsb.fnscale = FALSE,
lbfgsb.parscale = FALSE,
nm.ndeps = 1e-14,
nm.maxit = 10000,
nm.abstol = 1e-14,
nm.reltol = 1e-14,
nm.alpha = 1.0,
nm.beta = 0.5,
nm.gamma = 2.0,
nm.fnscale = FALSE,
nm.parscale = FALSE)

```

Arguments

llh `llh = c("filter", "internal", "testing")[1]`, defaults to "filter".
nlminb.eval.max Maximum number of evaluations of the objective function allowed, defaults to 200.
nlminb.iter.max Maximum number of iterations allowed, defaults to 150.
nlminb.abs.tol Absolute tolerance, defaults to 1e-20.

<code>nlminb.rel.tol</code>	Relative tolerance, defaults to 1e-10.
<code>nlminb.x.tol</code>	X tolerance, defaults to 1.5e-8.
<code>nlminb.fscale</code>	defaults to FALSE.
<code>nlminb.xscale</code>	defaults to FALSE.
<code>nlminb.step.min</code>	Minimum step size, defaults to 2.2e-14.
<code>nlminb.scale</code>	defaults to 1.
<code>sqp.mit</code>	maximum number of iterations, defaults to 200.
<code>sqp.mfv</code>	maximum number of function evaluations, defaults to 500.
<code>sqp.met</code>	specifies scaling strategy: <code>sqp.met=1</code> - no scaling <code>sqp.met=2</code> - preliminary scaling in 1st iteration (default) <code>sqp.met=3</code> - controlled scaling <code>sqp.met=4</code> - interval scaling <code>sqp.met=5</code> - permanent scaling in all iterations
<code>sqp.mec</code>	correction for negative curvature: <code>sqp.mec=1</code> - no correction <code>sqp.mec=2</code> - Powell correction (default)
<code>sqp.mer</code>	restarts after unsuccessful variable metric updates: <code>sqp.mer=0</code> - no restarts <code>sqp.mer=1</code> - standard restart
<code>sqp.mes</code>	interpolation method selection in a line search: <code>sqp.mes=1</code> - bisection <code>sqp.mes=2</code> - two point quadratic interpolation <code>sqp.mes=3</code> - three point quadratic interpolation <code>sqp.mes=4</code> - three point cubic interpolation (default)
<code>sqp.xmax</code>	maximum stepsize, defaults to 1.0e+3.
<code>sqp.tolx</code>	tolerance for the change of the coordinate vector, defaults to 1.0e-16.
<code>sqp.tolc</code>	tolerance for the constraint violation, defaults to 1.0e-6.
<code>sqp.tolg</code>	tolerance for the Lagrangian function gradient, defaults to 1.0e-6.
<code>sqp.told</code>	defaults to 1.0e-6.
<code>sqp.tols</code>	defaults to 1.0e-4.
<code>sqp.rpf</code>	value of the penalty coefficient, default to 1.0D-4. The default value may be relatively small. Therefore, larger value, say one, can sometimes be more suitable.
<code>lbfgsb.REPORT</code>	The frequency of reports for the "BFGS" and "L-BFGS-B" methods if <code>control\$trace</code> is positive. Defaults to every 10 iterations.
<code>lbfgsb.lmm</code>	is an integer giving the number of BFGS updates retained in the "L-BFGS-B" method, It defaults to 5.

<code>lbfgsb.factr</code>	controls the convergence of the "L-BFGS-B" method. Convergence occurs when the reduction in the objective is within this factor of the machine tolerance. Default is $1e7$, that is a tolerance of about $1.0e-8$.
<code>lbfgsb.pgtol</code>	helps control the convergence of the "L-BFGS-B" method. It is a tolerance on the projected gradient in the current search direction. This defaults to zero, when the check is suppressed.
<code>lbfgsb.fnscale</code>	defaults to FALSE.
<code>lbfgsb.parscale</code>	defaults to FALSE.
<code>nm.ndeps</code>	A vector of step sizes for the finite-difference approximation to the gradient, on <code>par/parscale</code> scale. Defaults to $1e-3$.
<code>nm.maxit</code>	The maximum number of iterations. Defaults to 100 for the derivative-based methods, and 500 for "Nelder-Mead". For "SANN" <code>maxit</code> gives the total number of function evaluations. There is no other stopping criterion. Defaults to 10000.
<code>nm.abstol</code>	The absolute convergence tolerance. Only useful for non-negative functions, as a tolerance for reaching zero.
<code>nm.reltol</code>	Relative convergence tolerance. The algorithm stops if it is unable to reduce the value by a factor of <code>reltol * (abs(val) + reltol)</code> at a step. Defaults to <code>sqrt(.Machine\$double.eps)</code> , typically about $1e-8$.
<code>nm.alpha</code> , <code>nm.beta</code> , <code>nm.gamma</code>	Scaling parameters for the "Nelder-Mead" method. <code>alpha</code> is the reflection factor (default 1.0), <code>beta</code> the contraction factor (0.5), and <code>gamma</code> the expansion factor (2.0).
<code>nm.fnscale</code>	An overall scaling to be applied to the value of <code>fn</code> and <code>gr</code> during optimization. If negative, turns the problem into a maximization problem. Optimization is performed on <code>fn(par)/fnscale</code> .
<code>nm.parscale</code>	A vector of scaling values for the parameters. Optimization is performed on <code>par/parscale</code> and these should be comparable in the sense that a unit change in any element produces about a unit change in the scaled value.

Value

returns a list.

Author(s)

Diethelm Wuertz for the Rmetrics R-port,
 R Core Team for the 'optim' R-port,
 Douglas Bates and Deepayan Sarkar for the 'nlminb' R-port,
 Bell-Labs for the underlying PORT Library,
 Ladislav Luksan for the underlying Fortran SQP Routine,
 Zhu, Byrd, Lu-Chen and Nocedal for the underlying L-BFGS-B Routine.

Examples

```
##
```

garchSim

*Univariate GARCH/APARCH Time Series Simulation***Description**

Simulates a univariate GARCH/APARCH time series model.

Usage

```
garchSim(spec = garchSpec(), n = 100, n.start = 100, extended = FALSE)
```

Arguments

extended	logical parameter if the output series should be a 3 columns <code>timeSeries</code> object with <code>garch</code> , <code>sigma</code> and <code>eps</code> data (<code>extended = TRUE</code>) or a univariate GARCH/APARCH time series (<code>extended = FALSE</code>)
spec	a specification object of class "fGARCHSPEC" as returned by the function <code>garchSpec</code> . The model parameters are taken from the <code>@model</code> slot, a list with the following entries: <code>omega</code> - the constant coefficient of the variance equation, by default 1e-6; <code>alpha</code> - the value or vector of autoregressive coefficients, by default 0.1, specifying a model of order 1; <code>beta</code> - the value or vector of variance coefficients, by default 0.8, specifying a model of order 1; The optional values for the linear part are: <code>mu</code> - the intercept value, by default 0 (it implies that the mean = $\mu/(1-\text{sum}(\text{ar}))$); <code>ar</code> - the autoregressive ARMA coefficients, by default 0; <code>ma</code> - the moving average ARMA coefficients, by default 0. The optional parameters for the conditional distributions are: <code>skew</code> - the skewness parameter (also named <code>xi</code>), by default 0.9, effective only for the "dsnrm", the "dsged", and the "dsstd" skewed conditional distributions; <code>shape</code> = the shape parameter (also named <code>nu</code>), by default 2 for the "dged" and "dsged", and by default 4 for the "dstd" and "dsstd" conditional distributions. See also below for further details.
n	length of output series, an integer value. An integer value, by default <code>n=100</code> .
n.start	length of "burn-in" period, by default 100.

Details

The function `garchSim` simulates an univariate GARCH or APARCH time series process as specified by the argument `model`.

The `model` is an object of class "fGARCHSPEC" as returned by the function `garchSpec`. The returned model specification comes with a slot `@model` which is a list of just the numeric parameter entries. These are recognized and extracted for use by the function `garchSim`.

By default the series will be returned as an object of class "ts" or as a "numeric" vector. Having time/date positions, e.g. from an empirical process the numeric vector can be easily transformed into other time series objects like "timeSeries" or "zoo". So one can estimate the parameters of a GARCH process from empirical data using the function `garchFit` and simulate than statistically equivalent GARCH processes with the same set of model parameters using the function `garchSim`.

The third entry in the argument `returnClass="mts"` allows to return a trivariate time series, where the first column contains the simulated "garch" process, the second column the conditional standard deviations "h", and the last column the innovations named "eps".

Note, the default model specifies Bollerslev's GARCH(1,1) model with normal distributed innovations.

Value

The function `garchSim` returns an objects of class "timeSeries" attributed by a list with entry `$garchSpec` giving the GARCH specification structure as returned by the function `garchSpec` and with information on conditional standard deviations and innovations. See details above.

Author(s)

Diethelm Wuertz for the Rmetrics R-port.

Examples

```
## garchSpec -
spec = garchSpec()
spec

## garchSim -
# Simulate a "timeSeries" object:
x = garchSim(spec, n = 50)
class(x)
print(x)

## More simulations ...

# Default GARCH(1,1) - uses default parameter settings
spec = garchSpec(model = list())
garchSim(spec, n = 10)

# ARCH(2) - use default omega and specify alpha, set beta=0!
spec = garchSpec(model = list(alpha = c(0.2, 0.4), beta = 0))
garchSim(spec, n = 10)

# AR(1)-ARCH(2) - use default mu, omega
spec = garchSpec(model = list(ar = 0.5, alpha = c(0.3, 0.4), beta = 0))
garchSim(spec, n = 10)

# AR([1,5])-GARCH(1,1) - use default garch values and subset ar[.]
spec = garchSpec(model = list(mu = 0.001, ar = c(0.5,0,0,0,0.1)))
garchSim(spec, n = 10)
```

```

# ARMA(1,2)-GARCH(1,1) - use default garch values
spec = garchSpec(model = list(ar = 0.5, ma = c(0.3, -0.3)))
garchSim(spec, n = 10)

# GARCH(1,1) - use default omega and specify alpha/beta
spec = garchSpec(model = list(alpha = 0.2, beta = 0.7))
garchSim(spec, n = 10)

# GARCH(1,1) - specify omega/alpha/beta
spec = garchSpec(model = list(omega = 1e-6, alpha = 0.1, beta = 0.8))
garchSim(spec, n = 10)

# GARCH(1,2) - use default omega and specify alpha[1]/beta[2]
spec = garchSpec(model = list(alpha = 0.1, beta = c(0.4, 0.4)))
garchSim(spec, n = 10)

# GARCH(2,1) - use default omega and specify alpha[2]/beta[1]
spec = garchSpec(model = list(alpha = c(0.12, 0.04), beta = 0.08))
garchSim(spec, n = 10)

# snorm-ARCH(1) - use defaults with skew Normal
spec = garchSpec(model = list(beta = 0, skew = 0.8), cond.dist = "snorm")
garchSim(spec, n = 10)

# sged-GARCH(1,1) - using defaults with skew GED
model = garchSpec(model = list(skew = 0.93, shape = 3), cond.dist = "sged")
garchSim(model, n = 10)

# Taylor Schwert GARCH(1,1) - this belongs to the family of APARCH Models
spec = garchSpec(model = list(delta = 1))
garchSim(spec, n = 10)

# AR(1)-t-APARCH(2, 1) - a little bit more complex specification ...
spec = garchSpec(model = list(mu = 1.0e-4, ar = 0.5, omega = 1.0e-6,
  alpha = c(0.10, 0.05), gamma = c(0, 0), beta = 0.8, delta = 1.8,
  shape = 4, skew = 0.85), cond.dist = "sstd")
garchSim(spec, n = 10)

```

garchSpec

Univariate GARCH Time Series Specification

Description

Specifies an univariate GARCH time series model.

Usage

```

garchSpec(model = list(), presample = NULL,
  cond.dist = c("norm", "ged", "std", "snorm", "sged", "sstd"),
  rseed = NULL)

```

Arguments

<code>cond.dist</code>	a character string naming the desired conditional distribution. Valid values are "norm", "ged", "std", "snorm", "sged", "sstd". The default value is the normal distribution.
<code>model</code>	<p>a list of GARCH model parameters:</p> <p>omega - the constant coefficient of the variance equation, by default 1e-6;</p> <p>alpha - the value or vector of autoregressive coefficients, by default 0.1, specifying a model of order 1;</p> <p>beta - the value or vector of variance coefficients, by default 0.8, specifying a model of order 1;</p> <p>The values for the linear part are:</p> <p>mu - the mean value, by default NULL;</p> <p>ar - the autoregressive ARMA coefficients, by default NULL;</p> <p>ma - the moving average ARMA coefficients, by default NULL.</p> <p>The parameters for the conditional distributions are:</p> <p>skew - the skewness parameter (also named "xi"), by default 0.9, effective only for the "dsnorm", the "dsged", and the "dsstd" skewed conditional distributions;</p> <p>shape - the shape parameter (also named "nu"), by default 2 for the "dged" and "dsged", and by default 4 for the "dstd" and "dsstd" conditional distributions.</p> <p>Note, the default <code>model=list()</code> specifies Bollerslev's GARCH(1,1) model with normal conditional distributed innovations.</p>
<code>presample</code>	a numeric three column matrix with start values for the series, for the innovations, and for the conditional variances. For an ARMA(m,n)-GARCH(p,q) process the number of rows must be at least $\max(m,n,p,q)+1$, longer presamples are cutted. Note, all presamples are initialized by a normal-GARCH(p,q) process.
<code>rseed</code>	single integer argument, the seed for the initialization of the random number generator for the innovations. Using the default value <code>rseed=NULL</code> then the random number generation will be started with <code>set.seed(0)</code> .

Details

The function `garchSpec` specifies a GARCH or APARCH time series process which we can use for simulating artificial GARCH and/or APARCH models. This is very useful for testing the GARCH parameter estimation results, since your model parameters are known and well specified.

For example specifying a subet AR(5[1,5])-GARCH(2,1) model with a standardized Student-t distribution with four degrees of freedom will return the following printed output:

```
garchSpec(model = list(ar = c(0.5,0,0,0,0.1), alpha =
  c(0.1, 0.1), beta = 0.75, shape = 4), cond.dist = "std")
```

Formula:

```
~ ar(5) + garch(2, 1)
```

Model:

```

ar:      0.5 0 0 0 0.1
omega: 1e-06
alpha: 0.1 0.1
beta: 0.75
Distribution:
  std
Distributional Parameter:
  nu = 4
Presample:
  time          z          h y
0          0 -0.3262334 2e-05 0
-1        -1  1.3297993 2e-05 0
-2        -2  1.2724293 2e-05 0
-3        -3  0.4146414 2e-05 0
-4        -4 -1.5399500 2e-05 0

```

The "Formula" describes the formula expression specifying the generating process, "Model" lists the associated model parameters, "Distribution" the type of the conditional distribution function in use, "Distributional Parameters" lists the distributional parameter (if any), and the "Presample" shows the presample input matrix.

If we have specified `presample=NULL` in the argument list, then the presample is generated automatically by default as norm-AR()-GARCH() process.

Value

The returned value is an object of class "fGARCHSPEC".

Author(s)

Diethelm Wuertz for the Rmetrics R-port.

Examples

```

## garchSpec -

# Normal Conditional Distribution:
spec = garchSpec()
spec

# Skewed Normal Conditional Distribution:
spec = garchSpec(model = list(skew = 0.8), cond.dist = "snorm")
spec

# Skewed GED Conditional Distribution:
spec = garchSpec(model = list(skew = 0.9, shape = 4.8), cond.dist = "sged")
spec

## More specifications ...

```

```

# Default GARCH(1,1) - uses default parameter settings
garchSpec(model = list())

# ARCH(2) - use default omega and specify alpha, set beta=0!
garchSpec(model = list(alpha = c(0.2, 0.4), beta = 0))

# AR(1)-ARCH(2) - use default mu, omega
garchSpec(model = list(ar = 0.5, alpha = c(0.3, 0.4), beta = 0))

# AR([1,5])-GARCH(1,1) - use default garch values and subset ar[.]
garchSpec(model = list(mu = 0.001, ar = c(0.5,0,0,0,0.1)))

# ARMA(1,2)-GARCH(1,1) - use default garch values
garchSpec(model = list(ar = 0.5, ma = c(0.3, -0.3)))

# GARCH(1,1) - use default omega and specify alpha/beta
garchSpec(model = list(alpha = 0.2, beta = 0.7))

# GARCH(1,1) - specify omega/alpha/beta
garchSpec(model = list(omega = 1e-6, alpha = 0.1, beta = 0.8))

# GARCH(1,2) - use default omega and specify alpha[1]/beta[2]
garchSpec(model = list(alpha = 0.1, beta = c(0.4, 0.4)))

# GARCH(2,1) - use default omega and specify alpha[2]/beta[1]
garchSpec(model = list(alpha = c(0.12, 0.04), beta = 0.08))

# snorm-ARCH(1) - use defaults with skew Normal
garchSpec(model = list(beta = 0, skew = 0.8), cond.dist = "snorm")

# sged-GARCH(1,1) - using defaults with skew GED
garchSpec(model = list(skew = 0.93, shape = 3), cond.dist = "sged")

# Taylor Schwert GARCH(1,1) - this belongs to the family of APARCH Models
garchSpec(model = list(delta = 1))

# AR(1)-t-APARCH(2, 1) - a little bit more complex specification ...
garchSpec(model = list(mu = 1.0e-4, ar = 0.5, omega = 1.0e-6,
  alpha = c(0.10, 0.05), gamma = c(0, 0), beta = 0.8, delta = 1.8,
  shape = 4, skew = 0.85), cond.dist = "sstd")

```

plot-methods

GARCH Plot Methods

Description

Plot methods for GARCH Modelling.

Usage

```
## S4 method for signature 'fGARCH,missing':
plot(x, which = "ask", ...)
```

Arguments

```
x          an object of class "fREG"
which      a character string denoting which plot should be displayed.
...       optional arguments to be passed.
```

Details

The generic function `plot` allows to display 13 graphs. These are the

- Time SeriesPlot
- Conditional Standard Deviation Plot
- Series Plot with 2 Conditional SD Superimposed
- Autocorrelation function Plot of Observations
- Autocorrelation function Plot of Squared Observations
- Cross Correlation Plot
- Residuals Plot
- Conditional Standard Deviations Plot
- Standardized Residuals Plot
- ACF Plot of Standardized Residuals
- ACF Plot of Squared Standardized Residuals
- Cross Correlation Plot between r^2 and r
- Quantile-Quantile Plot of Standardized Residuals

Methods

```
x = "ANY", y = "ANY" Generic function.
x = "fGARCH", y = "missing" Plot function for objects of class "fGARCH".
```

Author(s)

Diethelm Wuertz for the Rmetrics R-port.

Examples

```
## garchSim -
# Default Garch(1,1) Model:
x = garchSim(n = 200)
head(x)

## garchFit -
fit = garchFit(formula = ~ garch(1, 1), data = x, trace = FALSE)

## Batch Plot:
```

```

plot(fit, which = 3)

## Not run:
## Plot:
# Interactive Plot:
plot(fit)

## End(Not run)

```

predict-methods *GARCH Prediction Function*

Description

Predicts a time series from a fitted GARCH object.

Usage

```

## S4 method for signature 'fGARCH':
predict(object, n.ahead = 10, trace = FALSE, mse = c("cond", "uncond"), plot=FALSE, nx

```

Arguments

<code>n.ahead</code>	an integer value, denoting the number of steps to be forecasted, by default 10.
<code>object</code>	an object of class <code>fGARCH</code> as returned by the function <code>garchFit</code> .
<code>trace</code>	a logical flag. Should the prediction process be traced? By default <code>trace=FALSE</code> .
<code>mse</code>	If set to "cond", <code>meanError</code> is defined as the conditional mean errors $\sqrt{E_t[x_{t+h} - E_t(x_{t+h})]^2}$. If set to "uncond", it is defined as $\sqrt{E[x_{t+h} - E_t(x_{t+h})]^2}$.
<code>plot</code>	If set to TRUE, the confidence intervals are computed and plotted
<code>nx</code>	The number of observations to be plotted along with the predictions. The default is <code>round(n*0.25)</code> , where <code>n</code> is the sample size.
<code>crit_val</code>	The critical values for the confidence intervals when <code>plot</code> is set to TRUE. The intervals are defined as $\hat{x}_{t+h} + \text{crit_val}[2] * \text{meanError}$ and $\hat{x}_{t+h} + \text{crit_val}[1] * \text{meanError}$ if two critical values are provided and $\hat{x}_{t+h} \pm \text{crit_val} * \text{meanError}$ if only one is given. If you do not provide critical values, they will be computed automatically.
<code>conf</code>	The confidence level for the confidence intervals if <code>crit_val</code> is not provided. By default it is set to 0.95. The critical values are then computed using the conditional distribution that was chosen to create the object with <code>garchFit</code> using the same shape and skew parameters. If the conditional distribution was set to "QMLE", the critical values are computed using the empirical distribution of the standardized residuals.
<code>...</code>	additional arguments to be passed.

Value

returns a data frame with the following columns: "meanForecast", "meanError", and "standardDeviation".

The number of records equals the number of forecasting steps `n.ahead`.

Methods

object = "ANY" Generic function.

object = "fGARCH" Predict function for objects of class "fGARCH".

Author(s)

Diethelm Wuertz for the Rmetrics R-port.

Examples

```
## garchFit -
# Parameter Estimation of Default GARCH(1,1) Model:
set.seed(123)
fit = garchFit(~ garch(1, 1), data = garchSim(), trace = FALSE)
fit

## predict -
predict(fit, n.ahead = 10)
predict(fit, n.ahead = 10, mse="uncond")

## predict with plotting: critical values = +- 2

predict(fit, n.ahead = 10, plot=TRUE, crit_val=2)

## predict with plotting: automatic critical values
## for different conditional distributions

set.seed(321)
fit2 = garchFit(~ garch(1, 1), data = garchSim(), trace = FALSE, cond.dist="sged")

## 95% confidence level
predict(fit2, n.ahead=20, plot=TRUE)

set.seed(444)
fit3 = garchFit(~ garch(1, 1), data = garchSim(), trace = FALSE, cond.dist="QMLE")

## 90% confidence level and nx=100

predict(fit3, n.ahead=20, plot=TRUE, conf=.9, nx=100)
```

residuals-methods *Extract GARCH Model Residuals*

Description

Extracts residuals from a fitted GARCH object.

Usage

```
## S4 method for signature 'fGARCH':  
residuals(object, standardize = FALSE)
```

Arguments

object an object of class "fGARCH" as returned from the function `garchFit`.
standardize a logical flag, should the residuals be standardized?

Details

The function extracts the `@residuals` slot from an object of class "fGARCH" as returned by the function `garchFit`.

The class of the returned value depends on the input to the function `garchFit` who created the object. The returned value is always of the same class as the input object to the argument `data` in the function `garchFit`, i.e. if you fit a "timeSeries" object, you will get back from the function `fitted` also a "timeSeries" object, if you fit an object of class "zoo", you will get back again a "zoo" object. The same holds for a "numeric" vector, for a "data.frame", and for objects of class "ts", "mts".

In contrast, the slot itself returns independent of the class of the data input always a numeric vector, i.e. the function call `rslot(object, "fitted")` will return a numeric vector.

Methods

object = "ANY" Generic function

object = "fGARCH" Extractor function for residual from an object of class "fGARCH".

Note

`residuals` is a generic function which extracts residual values from objects returned by modeling functions.

Author(s)

Diethelm Wuertz for the Rmetrics R-port.

Examples

```
## Swiss Pension func Index -
x = as.timeSeries(data(LPP2005REC))

## garchFit
fit = garchFit(LPP40 ~ garch(1, 1), data = 100*x, trace = FALSE)
fit

## residuals -
res = residuals(fit)
head(res)
class(res)

## slot -
res = slot(fit, "residuals")
head(res)
```

sged

*Skew GED Distribution and Parameter Estimation***Description**

Functions to compute density, distribution function, quantile function and to generate random variates for the generalized error distribution. In addition maximum likelihood estimators are available to fit the parameters of the distribution.

The functions are:

[dpqr]ged	Symmetric GED Distribution,
[dpqr]sged	Skew GED Distribution,
gedFit	MLE parameter fit for a GED distribution,
sgedFit	MLE parameter fit for a skew GED distribution,
sgedSlider	Displays interactively skew GED distribution.

Usage

```
dged(x, mean = 0, sd = 1, nu = 2)
pged(q, mean = 0, sd = 1, nu = 2)
qged(p, mean = 0, sd = 1, nu = 2)
rged(n, mean = 0, sd = 1, nu = 2)

dsged(x, mean = 0, sd = 1, nu = 2, xi = 1.5)
psged(q, mean = 0, sd = 1, nu = 2, xi = 1.5)
qsged(p, mean = 0, sd = 1, nu = 2, xi = 1.5)
rsged(n, mean = 0, sd = 1, nu = 2, xi = 1.5)

gedFit(x, ...)
```

```
sgedFit(x, ...)
sgedSlider(type = c("dist", "rand"))
```

Arguments

<code>mean, sd, nu, xi</code>	location parameter <code>mean</code> , scale parameter <code>sd</code> , shape parameter <code>nu</code> , skewness parameter <code>xi</code> .
<code>n</code>	the number of observations.
<code>p</code>	a numeric vector of probabilities.
<code>type</code>	a character string denoting which interactive plot should be displayed. Either a distribution plot <code>type="dist"</code> , the default value, or a random variates plot, <code>type="rand"</code> .
<code>x, q</code>	a numeric vector of quantiles.
<code>...</code>	parameters parsed to the optimization function <code>nlm</code> .

Details

Parameter Estimation:

The function `nlm` is used to minimize the "negative" maximum log-likelihood function. `nlm` carries out a minimization using a Newton-type algorithm.

Value

`d*` returns the density, `p*` returns the distribution function, `q*` returns the quantile function, and `r*` generates random deviates, all values are numeric vectors.

`[s]gedFit` returns a list with the following components:

<code>par</code>	The best set of parameters found.
<code>objective</code>	The value of objective corresponding to <code>par</code> .
<code>convergence</code>	An integer code. 0 indicates successful convergence.
<code>message</code>	A character string giving any additional information returned by the optimizer, or NULL. For details, see PORT documentation.
<code>iterations</code>	Number of iterations performed.
<code>evaluations</code>	Number of objective function and gradient function evaluations.

Author(s)

Diethelm Wuertz for the Rmetrics R-port.

References

Nelson D.B. (1991); *Conditional Heteroscedasticity in Asset Returns: A New Approach*, *Econometrica*, 59, 347–370.

Fernandez C., Steel M.F.J. (2000); *On Bayesian Modelling of Fat Tails and Skewness*, Preprint, 31 pages.

Examples

```
## sged -
par(mfrow = c(2, 2))
set.seed(1953)
r = rsged(n = 1000)
plot(r, type = "l", main = "sged", col = "steelblue")

# Plot empirical density and compare with true density:
hist(r, n = 25, probability = TRUE, border = "white", col = "steelblue")
box()
x = seq(min(r), max(r), length = 201)
lines(x, dsGED(x), lwd = 2)

# Plot df and compare with true df:
plot(sort(r), (1:1000/1000), main = "Probability", col = "steelblue",
      ylab = "Probability")
lines(x, psGED(x), lwd = 2)

# Compute quantiles:
round(qsGED(psGED(q = seq(-1, 5, by = 1))), digits = 6)

## sgedFit -
sgedFit(r)

## Not run:
## sgedSlider -
if (require(tcltk)) {
  sgedSlider("dist")
  sgedSlider("rand")
}

## End(Not run)
```

Description

Show methods for GARCH Modelling.

Methods

object = "ANY" Generic function.

object = "fGARCH" Print function for objects of class "fGARCH".

object = "fGARCHSPEC" Print function for objects of class "fGARCH".

Author(s)

Diethelm Wuertz for the Rmetrics R-port.

Examples

```
## garchSpec -
spec = garchSpec()
print(spec)

## garchSim -
x = garchSim(spec, n = 500)

## garchFit -
fit = garchFit(~ garch(1, 1), data = x)
print(fit)
```

 snorm

Skew Normal Distribution and Parameter Estimation

Description

Functions to compute density, distribution function, quantile function and to generate random variates for the skew normal distribution. In addition maximum likelihood estimators are available to fit the parameters of the distribution.

The functions are:

[dpqr] snorm	Skew Normal Distribution,
normFit	MLE parameter fit for Normal distribution,
snormFit	MLE parameter fit for skew Normal distribution,
snormSlider	Displays interactively skew Normal distribution.

Usage

```
dsnorm(x, mean = 0, sd = 1, xi = 1.5)
psnorm(q, mean = 0, sd = 1, xi = 1.5)
qsnorm(p, mean = 0, sd = 1, xi = 1.5)
rsnorm(n, mean = 0, sd = 1, xi = 1.5)

normFit(x, ...)
snormFit(x, ...)
```

```
snormSlider(type = c("dist", "rand"))
```

Arguments

`mean`, `sd`, `xi` location parameter `mean`, scale parameter `sd`, skewness parameter `xi`.
`n` the number of observations.
`p` a numeric vector of probabilities.
`type` a character string denoting which interactive plot should be displayed. Either a distribution plot `type="dist"`, the default value, or a random variates plot, `type="rand"`.
`x`, `q` a numeric vector of quantiles.
`...` parameters parsed to the optimization function `nlm`.

Details

Symmetric Normal Distribution:

The functions for the normal distribution are part of R's base package. The functions for the symmetric Student-t distribution are rescaled in such a way that they have unit variance in contrast to the Student-t family `dt`, `pt`, `qt` and `rt` which are part of R's base package. The generalized error distribution functions are defined as described by Nelson (1991).

Skew Normal Distribution:

The skew normal distribution functions are defined as described by Fernandez and Steel (2000).
`cr`

Parameter Estimation:

The function `nlm` is used to minimize the "negative" maximum log-likelihood function. `nlm` carries out a minimization using a Newton-type algorithm.

Value

`d*` returns the density, `p*` returns the distribution function, `q*` returns the quantile function, and `r*` generates random deviates,
all values are numeric vectors.

`[s]normFit` returns a list with the following components:

`par` The best set of parameters found.
`objective` The value of objective corresponding to `par`.
`convergence` An integer code. 0 indicates successful convergence.
`message` A character string giving any additional information returned by the optimizer, or NULL. For details, see PORT documentation.
`iterations` Number of iterations performed.
`evaluations` Number of objective function and gradient function evaluations.

Author(s)

Diethelm Wuertz for the Rmetrics R-port.

References

Fernandez C., Steel M.F.J. (2000); *On Bayesian Modelling of Fat Tails and Skewness*, Preprint, 31 pages.

Examples

```
## snorm -

# Random Numbers:
par(mfrow = c(2, 2))
set.seed(1953)
r = rsnorm(n = 1000)
plot(r, type = "l", main = "snorm", col = "steelblue")

# Plot empirical density and compare with true density:
hist(r, n = 25, probability = TRUE, border = "white", col = "steelblue")
box()
x = seq(min(r), max(r), length = 201)
lines(x, dsnorm(x), lwd = 2)

# Plot df and compare with true df:
plot(sort(r), (1:1000/1000), main = "Probability", col = "steelblue",
      ylab = "Probability")
lines(x, psnorm(x), lwd = 2)

# Compute quantiles:
round(qsnorm(psnorm(q = seq(-1, 5, by = 1))), digits = 6)

## snormFit -
snormFit(r)

## Not run:
## snormSlider -
if (require(tcltk)) {
  snormSlider("dist")
  snormSlider("rand")
}

## End(Not run)
```

Description

Functions to compute density, distribution function, quantile function and to generate random variates for the standardized skew Student-t distribution. In addition maximum likelihood estimators are available to fit the parameters of the distribution.

The functions are:

<code>[dpqr]std</code>	Symmetric Student-t Distribution,
<code>[dpqr]sstd</code>	Skew Student-t Distribution,
<code>stdFit</code>	MLE parameter fit for a Student-t distribution,
<code>sstdFit</code>	MLE parameter fit for a skew Student-t distribution,
<code>sstdSlider</code>	Displays interactively skew GED distribution.

Usage

```
dstd(x, mean = 0, sd = 1, nu = 5)
pstd(q, mean = 0, sd = 1, nu = 5)
qstd(p, mean = 0, sd = 1, nu = 5)
rstd(n, mean = 0, sd = 1, nu = 5)

dsstd(x, mean = 0, sd = 1, nu = 5, xi = 1.5)
psstd(q, mean = 0, sd = 1, nu = 5, xi = 1.5)
qsstd(p, mean = 0, sd = 1, nu = 5, xi = 1.5)
rsstd(n, mean = 0, sd = 1, nu = 5, xi = 1.5)

stdFit(x, ...)
sstdFit(x, ...)

sstdSlider(type = c("dist", "rand"))
```

Arguments

<code>mean, sd, nu, xi</code>	location parameter <code>mean</code> , scale parameter <code>sd</code> , shape parameter <code>nu</code> , skewness parameter <code>xi</code> .
<code>n</code>	the number of observations.
<code>p</code>	a numeric vector of probabilities.
<code>type</code>	a character string denoting which interactive plot should be displayed. Either a distribution plot <code>type="dist"</code> , the default value, or a random variates plot, <code>type="rand"</code> .
<code>x, q</code>	a numeric vector of quantiles.
<code>...</code>	parameters parsed to the optimization function <code>nlm</code> .

Details

Parameter Estimation:

The function `nlminb` is used to minimize the "negative" maximum log-likelihood function. `nlminb` carries out a minimization using a Newton-type algorithm.

Value

`d*` returns the density, `p*` returns the distribution function, `q*` returns the quantile function, and `r*` generates random deviates, all values are numeric vectors.

`[s]stdFit` returns a list with the following components:

<code>par</code>	The best set of parameters found.
<code>objective</code>	The value of objective corresponding to <code>par</code> .
<code>convergence</code>	An integer code. 0 indicates successful convergence.
<code>message</code>	A character string giving any additional information returned by the optimizer, or NULL. For details, see PORT documentation.
<code>iterations</code>	Number of iterations performed.
<code>evaluations</code>	Number of objective function and gradient function evaluations.

Author(s)

Diethelm Wuertz for the Rmetrics R-port.

References

Fernandez C., Steel M.F.J. (2000); *On Bayesian Modelling of Fat Tails and Skewness*, Preprint, 31 pages.

Examples

```
## sstd -
par(mfrow = c(2, 2))
set.seed(1953)
r = rsstd(n = 1000)
plot(r, type = "l", main = "sstd", col = "steelblue")

# Plot empirical density and compare with true density:
hist(r, n = 25, probability = TRUE, border = "white", col = "steelblue")
box()
x = seq(min(r), max(r), length = 201)
lines(x, dsstd(x), lwd = 2)

# Plot df and compare with true df:
plot(sort(r), (1:1000/1000), main = "Probability", col = "steelblue",
      ylab = "Probability")
lines(x, psstd(x), lwd = 2)

# Compute quantiles:
round(qsstd(psstd(q = seq(-1, 5, by = 1))), digits = 6)
```

```
## sstdFit -
  sstdFit(r, print.level = 2)
```

```
summary-methods    GARCH Summary Methods
```

Description

Summary methods for GARCH Modelling.

Methods

object = "ANY" Generic function

object = "fGARCH" Summary function for objects of class "fGARCH".

How to read a diagnostic summary report?

The first five sections return the title, the call, the mean and variance formula, the conditional distribution and the type of standard errors:

```
Title:
  GARCH Modelling

Call:
  garchFit(~ garch(1, 1), data = garchSim(), trace = FALSE)

Mean and Variance Equation:
  ~arch(0)

Conditional Distribution:
  norm

Std. Errors:
  based on Hessian
```

The next three sections return the estimated coefficients, and an error analysis including standard errors, t values, and probabilities, as well as the log Likelihood values from optimization:

```
Coefficient(s):
      mu      omega      alpha1      beta1
-5.79788e-05  7.93017e-06  1.59456e-01  2.30772e-01

Error Analysis:
      Estimate  Std. Error  t value  Pr(>|t|)
mu      -5.798e-05  2.582e-04  -0.225  0.822
```

```

omega 7.930e-06 5.309e-06 1.494 0.135
alpha1 1.595e-01 1.026e-01 1.554 0.120
beta1 2.308e-01 4.203e-01 0.549 0.583

```

```

Log Likelihood:
-843.3991 normalized: -Inf

```

The next section provides results on standardized residuals tests, including statistic and p values, and on information criterion statistic including AIC, BIC, SIC, and HQIC:

Standardized Residuals Tests:

			Statistic	p-Value
Jarque-Bera Test	R	Chi ²	0.4172129	0.8117146
Shapiro-Wilk Test	R	W	0.9957817	0.8566985
Ljung-Box Test	R	Q(10)	13.05581	0.2205680
Ljung-Box Test	R	Q(15)	14.40879	0.4947788
Ljung-Box Test	R	Q(20)	38.15456	0.008478302
Ljung-Box Test	R ²	Q(10)	7.619134	0.6659837
Ljung-Box Test	R ²	Q(15)	13.89721	0.5333388
Ljung-Box Test	R ²	Q(20)	15.61716	0.7400728
LM Arch Test	R	TR ²	7.049963	0.8542942

Information Criterion Statistics:

AIC	BIC	SIC	HQIC
8.473991	8.539957	8.473212	8.500687

Author(s)

Diethelm Wuertz for the Rmetrics R-port.

Examples

```

## garchSim -
x = garchSim(n = 200)

## garchFit -
fit = garchFit(formula = x ~ garch(1, 1), data = x, trace = FALSE)
summary(fit)

```

TimeSeriesData

Time Series Data Sets

Description

Data sets used in the examples of the timeSeries packages.

 volatility-methods *Extract GARCH Model Volatility*

Description

Extracts volatility from a fitted GARCH object.

Usage

```
## S3 method for class 'fGARCH':
volatility(object, type = c("sigma", "h"), ...)
```

Arguments

<code>object</code>	an object of class <code>fGARCH</code> as returned from the function <code>garchFit()</code> .
<code>type</code>	a character string denoting if the conditional standard deviations "sigma" or the variances "h" should be returned.
<code>...</code>	additional arguments to be passed.

Details

The function extracts the `@volatility` from the slots `@sigma.t` or `@h.t` of an object of class "fGARCH" as returned by the function `garchFit`.

The class of the returned value depends on the input to the function `garchFit` who created the object. The returned value is always of the same class as the input object to the argument `data` in the function `garchFit`, i.e. if you fit a "timeSeries" object, you will get back from the function `fitted` also a "timeSeries" object, if you fit an object of class "zoo", you will get back again a "zoo" object. The same holds for a "numeric" vector, for a "data.frame", and for objects of class "ts", "mts".

In contrast, the slot itself returns independent of the class of the data input always a numeric vector, i.e. the function call `rslot(object, "fitted")` will return a numeric vector.

Methods

object = "ANY" Generic function.

object = "fGARCH" Extractor function for volatility or standard deviation from an object of class "fGARCH".

Note

`volatility` is a generic function which extracts volatility values from objects returned by modeling functions.

Author(s)

Diethelm Wuertz for the Rmetrics R-port.

Examples

```
## Swiss Pension func Index -
x = as.timeSeries(data(LPP2005REC))

## garchFit
fit = garchFit(LPP40 ~ garch(1, 1), data = 100*x, trace = FALSE)
fit

## volatility -
# Standard Deviation:
volatility = volatility(fit, type = "sigma")
head(volatility)
class(volatility)
# Variance:
volatility = volatility(fit, type = "h")
head(volatility)
class(volatility)

## slot -
volatility = slot(fit, "sigma.t")
head(volatility)
class(volatility)
volatility = slot(fit, "h.t")
head(volatility)
class(volatility)
```

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