

# Package ‘gets’

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**Type** Package

**Title** General-to-Specific (GETS) Modelling and Indicator Saturation Methods

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**Description** Automated General-to-Specific (GETS) modelling of the mean and variance of a regression, and indicator saturation methods for detecting and testing for structural breaks in the mean.

**License** GPL (>= 2)

**Depends** R (>= 3.3.0), zoo, parallel

**Suggests** lgarch, xtable

**BugReports** <https://github.com/gsucarrat/gets/issues>

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<http://www.sucarrat.net/R/gets>

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gets-package	<i>General-to-Specific (GETS) Modelling and Indicator Saturation Methods</i>
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## Description

Automated multi-path General-to-Specific (GETS) modelling of the mean and variance of a regression, and indicator saturation methods for detecting structural breaks in the mean. The mean can be specified as an autoregressive model with covariates (an 'AR-X' model), and the variance can be specified as a dynamic log-variance model with covariates (a 'log-ARCH-X' model). For the statistical details of the model, see Section 4 in Pretis, Reade and Sucarrat (2018).

The four main functions of the package are `arx`, `getsm`, `getsv` and `isat`. The first function, `arx`, estimates an AR-X model with (optionally) a log-ARCH-X specification on the log-variance. The second function, `getsm`, undertakes GETS model selection of the mean specification of an `arx` object. The third function, `getsv`, undertakes GETS model selection of the log-variance specification

of an `arx` object. The fourth function, `isat`, undertakes GETS model selection of an indicator saturated mean specification. Extraction functions (mainly S3 methods) are also available, together with additional auxiliary functions used by the main functions.

For the advanced user, the package provides facilities for user-defined GETS and ISAT methods. To this end, the main functions are `getsFun` and `isat`.

## Details

Package: gets  
Type: Package  
Version: 0.23  
Date: 2020-05-04  
License: GPL-2

The code originated in relation with G. Sucarrat and A. Escribano (2012). Subsequently, Felix Pretis and James Reade joined for the development of the `isat` code and related functions. For a recent introduction to the package, see Pretis, Reade and Sucarrat (2018).

## Author(s)

Felix Pretis, <http://www.felixpretis.org/>  
James Reade, <https://sites.google.com/site/jjamesreade/>  
Genaro Sucarrat, <http://www.sucarrat.net/>

Maintainer: Genaro Sucarrat

## References

- Jurgen, A. Doornik, Hendry, David F., and Pretis, Felix (2013): 'Step Indicator Saturation', Oxford Economics Discussion Paper, 658.
- Pretis, Felix, Reade, James and Sucarrat, Genaro (2018): 'Automated General-to-Specific (GETS) Regression Modeling and Indicator Saturation for Outliers and Structural Breaks'. *Journal of Statistical Software* 86, Number 3, pp. 1-44. DOI: <https://www.jstatsoft.org/article/view/v086i03>
- Carlos Santos, Hendry, David, F. and Johansen, Soren (2007): 'Automatic selection of indicators in a fully saturated regression'. *Computational Statistics*, vol 23:1, pp.317-335
- Sucarrat, Genaro and Escribano, Alvaro (2012): 'Automated Financial Model Selection: General-to-Specific Modelling of the Mean and Volatility Specifications', *Oxford Bulletin of Economics and Statistics* 74, Issue 5 (October), pp. 716-735

## See Also

[arx](#), [getsm](#), [getsv](#), [isat](#)

## Examples

```
##Simulate from an AR(1):  
set.seed(123)
```

```

y <- arima.sim(list(ar=0.4), 60)

##Estimate an AR(2) with intercept as mean specification
##and a log-ARCH(4) as log-volatility specification:
myModel <- arx(y, mc=TRUE, ar=1:2, arch=1:4)

##GETS modelling of the mean of myModel:
simpleMean <- getsm(myModel)

##GETS modelling of the log-variance of myModel:
simpleVar <- getsv(myModel)

##results:
print(simpleMean)
print(simpleVar)

##step indicator saturation of an iid normal series:
set.seed(123)
y <- rnorm(30)
isat(y)

```

---

arx

---

*Estimate an AR-X model with log-ARCH-X errors*


---

## Description

Estimation by OLS, two-step OLS if a variance specification is specified: In the first the mean specification (AR-X) is estimated, whereas in the second step the log-variance specification (log-ARCH-X) is estimated.

The AR-X mean specification can contain an intercept, AR-terms, lagged moving averages of the regressand and other conditioning covariates ( $X$ ). The log-variance specification can contain log-ARCH terms, asymmetry or 'leverage' terms, log(EqWMA) where EqWMA is a lagged equally weighted moving average of past squared residuals (a volatility proxy) and other conditioning covariates ( $X$ ).

## Usage

```

arx(y, mc=FALSE, ar=NULL, ewma=NULL, mxreg=NULL, vc=FALSE,
    arch=NULL, asym=NULL, log.ewma=NULL, vxreg=NULL, zero.adj=0.1,
    vc.adj=TRUE, vcov.type=c("ordinary", "white", "newey-west"),
    qstat.options=NULL, normality.JarqueB=FALSE, user.estimate=NULL,
    user.diagnostics=NULL, tol=1e-07, LAPACK=FALSE, plot=NULL)

```

## Arguments

y	numeric vector, time-series or <code>zoo</code> object. Missing values in the beginning and at the end of the series is allowed, as they are removed with the <code>na.trim</code> command
mc	logical. TRUE includes an intercept in the mean specification, whereas FALSE (default) does not

ar	either NULL (default) or an integer vector, say, <code>c(2,4)</code> or <code>1:4</code> . The AR-lags to include in the mean specification. If NULL, then no lags are included
ewma	either NULL (default) or a <code>list</code> with arguments sent to the <code>eqwma</code> function. In the latter case a lagged moving average of <code>y</code> is included as a regressor
mxreg	either NULL (default) or a numeric vector or matrix, say, a <code>zoo</code> object, of conditioning variables. Note that, if both <code>y</code> and <code>mxreg</code> are <code>zoo</code> objects, then their samples are chosen to match
vc	logical. TRUE includes an intercept in the log-variance specification, whereas FALSE (default) does not. If the log-variance specification contains any other item but the log-variance intercept, then <code>vc</code> is set to TRUE
arch	either NULL (default) or an integer vector, say, <code>c(1,3)</code> or <code>2:5</code> . The log-ARCH lags to include in the log-variance specification
asym	either NULL (default) or an integer vector, say, <code>c(1)</code> or <code>1:3</code> . The asymmetry (i.e. 'leverage') terms to include in the log-variance specification
log.ewma	either NULL (default) or a vector of the lengths of the volatility proxies, see <a href="#">leqwma</a>
vxreg	either NULL (default) or a numeric vector or matrix, say, a <code>zoo</code> object, of conditioning variables. If both <code>y</code> and <code>mxreg</code> are <code>zoo</code> objects, then their samples are chosen to match
zero.adj	numeric value between 0 and 1. The quantile adjustment for zero values. The default 0.1 means the zero residuals are replaced by the 10 percent quantile of the absolute residuals before taking the logarithm
vc.adj	logical. If TRUE (default), then the log-variance intercept is adjusted by the estimate of $E[\ln(z^2)]$ , where $z$ is the standardised error. This adjustment is needed for the conditional scale to be equal to the conditional standard deviation. If FALSE, then the log-variance intercept is not adjusted
vcov.type	character vector, "ordinary" (default), "white" or "newey-west". If "ordinary", then the ordinary variance-covariance matrix is used for inference. If "white", then the White (1980) heteroscedasticity-robust matrix is used. If "newey-west", then the Newey and West (1987) heteroscedasticity and autocorrelation-robust matrix is used
qstat.options	NULL (default) or an integer vector of length two, say, <code>c(1,1)</code> . The first value sets the lag-order of the AR diagnostic test, whereas the second value sets the lag-order of the ARCH diagnostic test. If NULL, then the two values of the vector are set automatically
normality.JarqueB	FALSE (default) or TRUE. If TRUE, then the results of the Jarque and Bera (1980) test for non-normality in the residuals are included in the estimation results.
user.estimator	NULL (default) or a <code>list</code> with one entry, <code>name</code> , containing the name of the user-defined estimator
user.diagnostics	NULL (default) or a <code>list</code> with two entries, <code>name</code> and <code>pval</code> , see the <code>user.fun</code> argument in <a href="#">diagnostics</a>

tol	numeric value (default = $1e-07$ ). The tolerance for detecting linear dependencies in the columns of the regressors (see <code>qr</code> function). Only used if LAPACK is FALSE (default) and <code>user.estimator</code> is NULL.
LAPACK	logical. If TRUE, then use LAPACK. If FALSE (default), then use LINPACK (see <code>qr</code> function). Only used if <code>user.estimator</code> is NULL.
plot	NULL or logical. If TRUE, then the fitted values and the residuals are plotted. If NULL (default), then the value set by <code>options</code> determines whether a plot is produced or not.

### Details

See Pretis, Reade and Sucarrat (2018)

### Value

A list of class 'arx'

### Author(s)

Genaro Sucarrat, <http://www.sucarrat.net/>

### References

- C. Jarque and A. Bera (1980): 'Efficient Tests for Normality, Homoscedasticity and Serial Independence'. *Economics Letters* 6, pp. 255-259
- Pretis, Felix, Reade, James and Sucarrat, Genaro (2018): 'Automated General-to-Specific (GETS) Regression Modeling and Indicator Saturation for Outliers and Structural Breaks'. *Journal of Statistical Software* 86, Number 3, pp. 1-44
- Halbert White (1980): 'A Heteroskedasticity-Consistent Covariance Matrix Estimator and a Direct Test for Heteroskedasticity', *Econometrica* 48, pp. 817-838
- Whitney K. Newey and Kenned D. West (1987): 'A Simple, Positive Semi-Definite, Heteroskedasticity and Autocorrelation Consistent Covariance Matrix', *Econometrica* 55, pp. 703-708

### See Also

Extraction functions (mostly S3 methods): `coef.arx`, `ES`, `fitted.arx`, `plot.arx`, `print.arx`, `recursive`, `residuals.arx`, `sigma.arx`, `rsquared`, `summary.arx`, `VaR` and `vcov.arx`

Related functions: `getsm`, `getsv`, `isat`

### Examples

```
##Simulate from an AR(1):
set.seed(123)
y <- arima.sim(list(ar=0.4), 70)

##estimate an AR(2) with intercept:
```

```

arx(y, mc=TRUE, ar=1:2)

##Simulate four independent Gaussian regressors:
xregs <- matrix(rnorm(4*70), 70, 4)

##estimate an AR(2) with intercept and four conditioning
##regressors in the mean:
arx(y, mc=TRUE, ar=1:2, mxreg=xregs)

##estimate a log-variance specification with a log-ARCH(4)
##structure:
arx(y, arch=1:4)

##estimate a log-variance specification with a log-ARCH(4)
##structure and an asymmetry/leverage term:
arx(y, arch=1:4, asym=1)

##estimate a log-variance specification with a log-ARCH(4)
##structure, an asymmetry or leverage term, a 10-period log(EWMA) as
##volatility proxy, and the log of the squareds of the conditioning
##regressors in the log-variance specification:
arx(y, arch=1:4, asym=1, log.ewma=list(length=10), vxreg=log(xregs^2))

##estimate an AR(2) with intercept and four conditioning regressors
##in the mean, and a log-variance specification with a log-ARCH(4)
##structure, an asymmetry or leverage term, a 10-period log(EWMA) as
##volatility proxy, and the log of the squareds of the conditioning
##regressors in the log-variance specification:
arx(y, mc=TRUE, ar=1:2, mxreg=xregs, arch=1:4, asym=1,
    log.ewma=list(length=10), vxreg=log(xregs^2))

```

---

biascorr	<i>Bias-correction of coefficients following general-to-specific model selection</i>
----------	--

---

## Description

Takes a vector of coefficients (valid for orthogonal variables), their standard errors, the significance level the variables were selected at, and the sample size, to return bias-corrected coefficient estimates to account for the bias induced by model selection.

## Usage

```
biascorr(b, b.se, p.alpha, T)
```

## Arguments

b	a Kx1 vector of coefficients.
b.se	a Kx1 vector of standard errors of the coefficients in 'b'.

p.alpha	numeric value between 0 and 1, the significance level at which selection was conducted.
T	integer, the sample size of the original model selection regression.

### Details

The function computes the bias-corrected estimates of coefficients in regression models post general-to-specific model selection using the approach by Hendry and Krolzig (2005). The results are valid for orthogonal regressors only. Bias correction can be applied to the coefficient path in `isat` models where the only additional covariate besides indicators is an intercept - see Pretis (2015).

### Value

Returns a  $K \times 3$  matrix, where the first column lists the original coefficients, the second column the one-step corrected coefficients, and the third column the two-step bias-corrected coefficients.

### Author(s)

Felix Pretis, <http://www.felixpretis.org/>

### References

Hendry, D.F. and Krolzig, H.M. (2005): 'The properties of automatic Gets modelling'. *Economic Journal*, 115, C32-C61.

Pretis, F. (2015): 'Testing for time-varying predictive accuracy using bias-corrected indicator saturation'. Oxford Department of Economics Discussion Paper.

Pretis, Felix, Reade, James and Sucarrat, Genaro (2018): 'Automated General-to-Specific (GETS) Regression Modeling and Indicator Saturation for Outliers and Structural Breaks'. *Journal of Statistical Software* 86, Number 3, pp. 1-44

### See Also

[isat](#), [coef.gets](#), [plot.gets](#), [isatvar](#), [isattest](#)

### Examples

```
###Bias-correction of the coefficient path of the Nile data
#nile <- as.zoo(Nile)
#isat.nile <- isat(nile, sis=TRUE, iis=FALSE, plot=TRUE, t.pval=0.005)
#var <- isatvar(isat.nile)
#biascorr(b=var$const.path, b.se=var$const.se, p.alpha=0.005, T=length(var$const.path))

##Bias-correction of the coefficient path on artificial data
#set.seed(123)
#d <- matrix(0,100,1)
#d[35:55] <- 1
#e <- rnorm(100, 0, 1)
#y <- d*1 +e

#ys <- isat(y, sis=TRUE, iis=FALSE, t.pval=0.01)
```



```
#var <- isatvar(ys)
#biascorr(b=var$const.path, b.se=var$const.se, p.alpha=0.01, T=length(var$const.path))
```

coef.arx

*Extraction functions for 'arx' objects***Description**

Extraction functions for objects of class 'arx'

**Usage**

```
## S3 method for class 'arx'
coef(object, spec=NULL, ...)
## S3 method for class 'arx'
fitted(object, spec=NULL, ...)
## S3 method for class 'arx'
logLik(object, ...)
## S3 method for class 'arx'
plot(x, spec=NULL, col=c("red","blue"),
     lty=c("solid","solid"), lwd=c(1,1), ...)
## S3 method for class 'arx'
print(x, signif.stars=FALSE, ...)
## S3 method for class 'arx'
residuals(object, std=FALSE, ...)
## S3 method for class 'arx'
sigma(object, ...)
## S3 method for class 'arx'
summary(object, ...)
## S3 method for class 'arx'
vcov(object, spec=NULL, ...)
```

**Arguments**

object	an object of class 'arx'
x	an object of class 'arx'
spec	NULL, "mean", "variance" or, in some instances, "both". When NULL is a valid value, then it is automatically determined whether information pertaining to the mean or variance specification should be returned
signif.stars	logical. If TRUE, then P-values are additionally encoded visually, see printCoefmat
std	logical. If FALSE (default), then the mean residuals are returned. If TRUE, then the standardised residuals are returned
col	colours of actual (default=blue) and fitted (default=red) lines
lty	types of actual (default=solid) and fitted (default=solid) lines
lwd	widths of actual (default=1) and fitted (default=1) lines
...	additional arguments

**Value**

coef: a numeric vector containing parameter estimates  
 fitted: a `zoo` object with fitted values  
 logLik: log-likelihood (normal density)  
 plot: a plot of the fitted values and the residuals  
 print: a print of the estimation results  
 residuals: a `zoo` object with the residuals  
 sigma: the regression standard error ('SE of regression')  
 summary: a print of the items in the `arx` object  
 vcov: variance-covariance matrix

**Author(s)**

Felix Pretis, <http://www.felixpretis.org/>  
 James Reade, <https://sites.google.com/site/jjamesreade/>  
 Genaro Sucarrat, <http://www.sucarrat.net/>

**See Also**

`arx`

**Examples**

```
##simulate from an AR(1):
set.seed(123)
y <- arima.sim(list(ar=0.4), 40)

##simulate four independent Gaussian regressors:
xregs <- matrix(rnorm(4*40), 40, 4)

##estimate an 'arx' model: An AR(2) with intercept and four conditioning
##regressors in the mean, and log-ARCH(3) in the variance:
mymod <- arx(y, mc=TRUE, ar=1:2, mxreg=xregs, arch=1:3)

##print results:
print(mymod)

##plot the fitted vs. actual values, and the residuals:
plot(mymod)

##print the entries of object 'mymod':
summary(mymod)

##extract coefficient estimates (automatically determined):
coef(mymod)

##extract mean coefficients only:
coef(mymod, spec="mean")
```

```
##extract log-variance coefficients only:
coef(mymod, spec="variance")

##extract all coefficient estimates:
coef(mymod, spec="both")

##extract regression standard error:
sigma(mymod)

##extract log-likelihood:
logLik(mymod)

##extract variance-covariance matrix of mean equation:
vcov(mymod)

##extract variance-covariance matrix of log-variance equation:
vcov(mymod, spec="variance")

##extract and plot the fitted mean values (automatically determined):
mfit <- fitted(mymod)
plot(mfit)

##extract and plot the fitted variance values:
vfit <- fitted(mymod, spec="variance")
plot(vfit)

##extract and plot both the fitted mean and variance values:
vfit <- fitted(mymod, spec="both")
plot(vfit)

##extract and plot the fitted mean values:
vfit <- fitted(mymod, spec="mean")
plot(vfit)

##extract and plot residuals:
epshat <- residuals(mymod)
plot(epshat)

##extract and plot standardised residuals:
zhat <- residuals(mymod, std=TRUE)
plot(zhat)
```

---

coef.gets

*Extraction functions for 'gets' objects*

---

### **Description**

Extraction functions for objects of class 'gets'

**Usage**

```

## S3 method for class 'gets'
coef(object, spec=NULL, ...)
## S3 method for class 'gets'
fitted(object, spec=NULL, ...)
## S3 method for class 'gets'
logLik(object, ...)
## S3 method for class 'gets'
plot(x, spec=NULL, col=c("red","blue"),
     lty=c("solid","solid"), lwd=c(1,1), ...)
## S3 method for class 'gets'
predict(object, spec=NULL, n.ahead=12, newmxreg=NULL,
        newvxreg=NULL, newindex=NULL, n.sim=5000, innov=NULL, probs=NULL,
        ci.levels=NULL, quantile.type=7, return=TRUE, verbose=FALSE, plot=NULL,
        plot.options=list(), ...)
## S3 method for class 'gets'
print(x, ...)
## S3 method for class 'gets'
residuals(object, std=NULL, ...)
## S3 method for class 'gets'
sigma(object, ...)
## S3 method for class 'gets'
summary(object, ...)
## S3 method for class 'gets'
vcov(object, spec=NULL, ...)

```

**Arguments**

object	an object of class 'gets'
x	an object of class 'gets'
spec	NULL, "mean", "variance" or, in some instances, "both". When NULL is a valid value, then it is automatically determined whether information pertaining to the mean or variance specification should be returned
std	logical. If FALSE (default), then the mean residuals are returned. If TRUE, then the standardised residuals are returned
n.ahead	integer that determines how many steps ahead predictions should be generated (the default is 12)
newmxreg	a matrix of n.ahead rows and NCOL(mxreg) columns with the out-of-sample values of the mxreg regressors
newvxreg	a matrix of n.ahead rows and NCOL(vxreg) columns with the out-of-sample values of the vxreg regressors
newindex	NULL (default) or the date-index for the zoo object returned by predict.arx. If NULL, then the function uses the in-sample index to generate the out-of-sample index
n.sim	integer, the number of replications used for the generation of the forecasts

innov	NULL (default) or a vector of length <code>n.ahead * n.sim</code> containing the standardised errors (that is, zero mean and unit variance) used for the forecast simulations. If NULL, then a classical bootstrap procedure is used to draw from the standardised in-sample residuals
probs	NULL (default) or a vector with the quantile-levels (values strictly between 0 and 1) of the forecast distribution. If NULL, then no quantiles are returned unless <code>ci.levels</code> is non-NULL
ci.levels	NULL (default) or a vector with the confidence levels (expressed as values strictly between 0 and 1) of the forecast distribution. The upper and lower values of the confidence interval(s) are returned as quantiles
quantile.type	an integer between 1 and 9 that selects which algorithm to be used in computing the quantiles, see the argument <code>type</code> in <a href="#">quantile</a>
return	logical. If TRUE (default), then the out-of-sample predictions are returned. The value FALSE, which does not return the predictions, may be of interest if only a prediction plot is of interest
verbose	logical with default FALSE. If TRUE, then additional information (typically the quantiles and/or the simulated series) used in the generation of forecasts is returned. If FALSE, then only the forecasts are returned
plot	NULL (default) or logical. If NULL, then the value set by <code>options\$plot</code> (see <a href="#">options</a> ) determines whether a plot is produced or not. If TRUE, then the out-of-sample forecasts are plotted.
plot.options	a list of options related to the plotting of forecasts, see 'Details'
col	colours of fitted (default=red) and actual (default=blue) lines
lty	types of fitted (default=solid) and actual (default=solid) lines
lwd	widths of fitted (default=1) and actual (default=1) lines
...	additional arguments

### Details

The `plot.options` argument is a list that controls the prediction plot, see 'Details' in [predict.arx](#)

### Value

coef:	a numeric vector containing parameter estimates
fitted:	a <a href="#">zoo</a> object with fitted values
logLik:	a numeric, the log-likelihood (normal density)
plot:	a plot of the fitted values and the residuals
predict:	a vector of class <a href="#">zoo</a> containing the out-of-sample forecasts, or a matrix of class <a href="#">zoo</a> containing the out-of-sample forecasts together with prediction-quantiles, or - if <code>return=FALSE</code> - NULL
print:	a print of the estimation results
residuals:	a <a href="#">zoo</a> object with the residuals
sigma:	the regression standard error ('SE of regression')
summary:	a print of the items in the <a href="#">gets</a> object
vcov:	a variance-covariance matrix

**Author(s)**

Felix Pretis, <http://www.felixpretis.org/>  
James Reade, <https://sites.google.com/site/jjamesreade/>  
Genaro Sucarrat, <http://www.sucarrat.net/>

**See Also**

[getsm](#), [getsv](#), [isat](#)

**Examples**

```
##Simulate from an AR(1):
set.seed(123)
y <- arima.sim(list(ar=0.4), 100)

##Simulate four independent Gaussian regressors:
xregs <- matrix(rnorm(4*100), 100, 4)

##estimate an AR(2) with intercept and four conditioning
##regressors in the mean, and a log-ARCH(3) in the variance:
mymod <- arx(y, mc=TRUE, ar=1:2, mxreg=xregs, arch=1:3)

##General-to-Specific (GETS) model selection of the mean:
meanmod <- getsm(mymod)

##General-to-Specific (GETS) model selection of the variance:
varmod <- getsv(mymod)

##print results:
print(meanmod)
print(varmod)

##plot the fitted vs. actual values, and the residuals:
plot(meanmod)
plot(varmod)

##generate and plot predictions of the mean:
predict(meanmod, plot=TRUE)

##print the entries of object 'gets':
summary(meanmod)
summary(varmod)

##extract coefficients of the simplified (specific) model:
coef(meanmod) #mean spec
coef(varmod) #variance spec

##extract log-likelihood:
logLik(mymod)

##extract coefficient-covariance matrix of simplified
##(specific) model:
```

```

vcov(meanmod) #mean spec
vcov(varmod) #variance spec

##extract and plot the fitted values:
mfit <- fitted(meanmod) #mean fit
plot(mfit)
vfit <- fitted(varmod) #variance fit
plot(vfit)

##extract and plot residuals:
epshat <- residuals(meanmod)
plot(epshat)

##extract and plot standardised residuals:
zhat <- residuals(varmod)
plot(zhat)

```

---

coef.isat

*Extraction functions for 'isat' objects*


---

## Description

Extraction functions for objects of class 'isat'

## Usage

```

## S3 method for class 'isat'
coef(object, ...)
## S3 method for class 'isat'
fitted(object, ...)
## S3 method for class 'isat'
logLik(object, ...)
## S3 method for class 'isat'
plot(x, col=c("red","blue"), lty=c("solid","solid"),
     lwd=c(1,1), coef.path=TRUE, ...)
## S3 method for class 'isat'
predict(object, n.ahead=12, newmxreg=NULL, newindex=NULL,
         n.sim=2000, probs=NULL, ci.levels=NULL, quantile.type=7,
         return=TRUE, verbose=FALSE, plot=NULL, plot.options=list(), ...)
## S3 method for class 'isat'
print(x, ...)
## S3 method for class 'isat'
residuals(object, std=FALSE, ...)
## S3 method for class 'isat'
sigma(object, ...)
## S3 method for class 'isat'
summary(object, ...)
## S3 method for class 'isat'
vcov(object, ...)

```

**Arguments**

object	an object of class 'isat'
x	an object of class 'isat'
std	logical. If FALSE (default), then the mean residuals are returned. If TRUE, then the standardised residuals are returned
n.ahead	integer that determines how many steps ahead predictions should be generated (the default is 12)
newmxreg	a matrix of n.ahead rows and NCOL(mxreg) columns with the out-of-sample values of the mxreg regressors
newindex	NULL (default) or the date-index for the zoo object returned by predict.arx. If NULL, then the function uses the in-sample index to generate the out-of-sample index
n.sim	integer, the number of replications used for the generation of the forecasts
probs	NULL (default) or a vector with the quantile-levels (values strictly between 0 and 1) of the forecast distribution. If NULL, then no quantiles are returned unless ci.levels is non-NULL
ci.levels	NULL (default) or a vector with the confidence levels (expressed as values strictly between 0 and 1) of the forecast distribution. The upper and lower values of the confidence interval(s) are returned as quantiles
quantile.type	an integer between 1 and 9 that selects which algorithm to be used in computing the quantiles, see the argument type in <a href="#">quantile</a>
return	logical. If TRUE (default), then the out-of-sample predictions are returned. The value FALSE, which does not return the predictions, may be of interest if only a prediction plot is of interest
verbose	logical with default FALSE. If TRUE, then additional information (typically the quantiles and/or the simulated series) used in the generation of forecasts is returned. If FALSE, then only the forecasts are returned
plot	NULL (default) or logical. If NULL, then the value set by options\$plot (see <a href="#">options</a> ) determines whether a plot is produced or not. If TRUE, then the out-of-sample forecasts are plotted.
plot.options	a list of options related to the plotting of forecasts, see 'Details'
col	colours of fitted (default=red) and actual (default=blue) lines
lty	types of fitted (default=solid) and actual (default=solid) lines
lwd	widths of fitted (default=1) and actual (default=1) lines
coef.path	logical. Only applicable if there are retained indicators after the application of isat
...	additional arguments

**Details**

The plot.options argument is a list that controls the prediction plot, see 'Details' in [predict.arx](#)



**Value**

coef:	numeric vector containing parameter estimates
fitted:	a <code>zoo</code> object with fitted values
logLik:	a numeric, the log-likelihood (normal density)
plot:	plot of the fitted values and the residuals
predict:	a vector of class <code>zoo</code> containing the out-of-sample forecasts, or a matrix of class <code>zoo</code> containing the out-of-sample forecasts together with prediction-quantiles, or - if <code>return=FALSE</code> - NULL
print:	a print of the estimation results
residuals:	a <code>zoo</code> object with the residuals
sigma:	the regression standard error ('SE of regression')
summary:	a print of the items in the <code>isat</code> object
vcov:	variance-covariance matrix

**Author(s)**

Felix Pretis, <http://www.felixpretis.org/>  
 James Reade, <https://sites.google.com/site/jjamesreade/>  
 Genaro Sucarrat, <http://www.sucarrat.net/>

**See Also**

[paths](#), [terminals](#), [coef.gets](#), [getsm](#), [arx](#)

**Examples**

```
##step indicator saturation:
set.seed(123)
y <- rnorm(30)
isatmod <- isat(y)

##print results:
print(isatmod)

##plot the fitted vs. actual values, and the residuals:
plot(isatmod)

##print the entries of object 'isatmod':
summary(isatmod)

##extract coefficients of the simplified (specific) model:
coef(isatmod)

##extract log-likelihood:
logLik(isatmod)

##extract the coefficient-covariance matrix of simplified
##(specific) model:
```

```
vcov(isatmod)

##extract and plot the fitted values:
mfit <- fitted(isatmod)
plot(mfit)

##extract and plot (mean) residuals:
epshat <- residuals(isatmod)
plot(epshat)

##extract and plot standardised residuals:
zhat <- residuals(isatmod, std=TRUE)
plot(zhat)

##generate forecasts of the simplified (specific) model:
predict(isatmod, newmxreg=matrix(1,12,1), plot=TRUE)
```

---

diagnostics

*Diagnostics tests*


---

## Description

Auxiliary function (i.e. not intended for the average user) called by the [arx](#), [getsm](#), [getsv](#), [isat](#) and [getsFun](#) functions. The `diagnostics` function undertakes tests for autocorrelation, ARCH and non-normality in a residual series, and user-defined diagnostics provided via the `user.fun` argument (see details). The autocorrelation and ARCH tests are conducted as Ljung and Box (1979) tests for autocorrelation in the residuals and squared residuals, respectively, whereas the test for non-normality is that of Jarque and Bera (1980).

## Usage

```
diagnostics(x, ar.LjungB=c(1, 0.025), arch.LjungB=c(1, 0.025),
  normality.JarqueB=NULL, verbose=TRUE, user.fun=NULL, ...)
```

## Arguments

<code>x</code>	a <a href="#">list</a> , for example the estimation result of <a href="#">ols</a> . The tests for serial correlation, ARCH and normality look for an entry in the list named <code>std.residuals</code> or <code>residuals</code>
<code>ar.LjungB</code>	a two element vector or <code>NULL</code> . In the former case, the first element contains the AR-order, the second element the significance level. If <code>NULL</code> , then a test for autocorrelation is not conducted
<code>arch.LjungB</code>	a two element vector or <code>NULL</code> . In the former case, the first element contains the ARCH-order, the second element the significance level. If <code>NULL</code> , then a test for ARCH is not conducted

<code>normality.JarqueB</code>	NULL (the default) or a value between 0 and 1. In the latter case, a test for non-normality is conducted using a significance level equal to <code>normality.JarqueB</code> . If NULL, then no test for non-normality is conducted
<code>verbose</code>	logical. If TRUE, then a <code>data.frame</code> with the results of the diagnostics is returned. If FALSE, then the return-value is a logical that indicates whether the model passes the diagnostics (TRUE if it does, otherwise FALSE)
<code>user.fun</code>	NULL or a <code>list</code> with at least one entry, <code>name</code> (must be of class character), which should contain the name of the user-defined function. See details
<code>...</code>	further arguments (ignored) to accommodate deleted arguments from past versions of the functions

## Details

The argument `user.fun` enables the user to specify additional diagnostics. To do this, the argument should be a `list` with at least one entry, `name` (of class character), that contains the name of the user-defined function. By default, the user-defined function is looked for in the global environment (`.GlobalEnv`). However, this can be changed by adding an entry named `envir` that indicates where the user-defined function resides. If the `verbose` argument is set to FALSE, then an entry named `pval` must be provided. This entry should contain the chosen significance level or levels, i.e. either a scalar or a vector of length equal to the number of p-values returned by the user-defined diagnostics function (see examples). Additional entries in the `list` are passed on as arguments to the user-defined function. The user-defined function should refer to the named items of the estimation result `x` (see examples), and the value returned by the user-defined function should be a matrix of dimension  $m \times 3$ . Here,  $m$  is the number of diagnostic tests performed by the user-defined function. For example, if only a single test is performed, then  $m=1$  and so the returned value should be a  $1 \times 3$  matrix (or a vector of length 3). The three columns of the  $m \times 3$  matrix should contain, in the following order, 1) the value(s) of the test-statistic(s), 2) the degrees of freedom(s) (or NA if there are none) of the tests, and 3) the p-value(s) of the test(s). When checking whether the model passes the diagnostics or not, the p-value(s) is(are) checked against the value(s) in the entry named `pval` in the `list` provided to `user.fun`.

## Value

If `verbose=TRUE`:

a `data.frame` that contains the diagnostics results

If `verbose=FALSE`:

a logical indicating whether the residuals and/or model passes ALL the diagnostics (TRUE if it does, FALSE otherwise)

## Author(s)

Genaro Sucarrat, <http://www.sucarrat.net/>

## References

C. Jarque and A. Bera (1980): 'Efficient Tests for Normality, Homoscedasticity and Serial Independence'. *Economics Letters* 6, pp. 255-259

G. Ljung and G. Box (1979): 'On a Measure of Lack of Fit in Time Series Models'. *Biometrika* 66, pp. 265-270

### See Also

[arx](#), [getsm](#), [getsv](#), [isat](#), [getsFun](#)

### Examples

```
##generate some data:
set.seed(123)
vY <- rnorm(20) #the regressand
mX <- matrix(rnorm(3*20), 20, 3) #the regressors
est <- ols(vY,mX)

##return a data-frame with autocorrelation and ARCH diagnostics (default),
##and check whether they pass (the default p-value is 0.025):
diagnostics(est)
diagnostics(est, verbose=FALSE)

##add the Jarque-Bera normality test to the diagnostics (w/p-value=0.05):
diagnostics(est, normality.JarqueB=0.05)
diagnostics(est, normality.JarqueB=0.05, verbose=FALSE)

##user-defined Shapiro-Wilks test for non-normality of the residuals:
SWtest <- function(x, ...){
  tmp <- shapiro.test(x$residuals) #do test on est$residuals
  return( c(tmp$statistic, NA, tmp$p.value) )
}
diagnostics(est, user.fun=list(name="SWtest", pval=0.05))
diagnostics(est, user.fun=list(name="SWtest", pval=0.05), verbose=FALSE)
```

---

dropvar

*Drop variable*

---

### Description

Drops columns in a matrix to avoid perfect multicollinearity.

### Usage

```
dropvar(x, tol=1e-07, LAPACK=FALSE, silent=FALSE)
```

### Arguments

**x** a matrix, possibly less than full column rank.

**tol** numeric value. The tolerance for detecting linear dependencies among regressors, see [qr](#) function. Only used if LAPACK is FALSE

LAPACK	logical, TRUE or FALSE (default). If true use LAPACK otherwise use LINPACK, see <a href="#">qr</a> function
silent	logical, TRUE (default) or FALSE. Whether to print a notification whenever a regressor is removed

### Details

Original function `drop.coef` developed by Rune Haubo B. Christensen in package `ordinal`, <https://cran.r-project.org/package=ordinal>.

### Value

a matrix whose regressors linearly independent

### Author(s)

Rune Haubo B. Christensen, with modifications by Genaro Sucarrat, <http://www.sucarrat.net/>

### References

Rune H.B. Christensen (2014): 'ordinal: Regression Models for Ordinal Data'. <https://cran.r-project.org/package=ordinal>

### See Also

[isat](#)

### Examples

```
set.seed(1)
x <- matrix(rnorm(20), 5)
dropvar(x) #full rank, none are dropped

x[,4] <- x[,1]*2
dropvar(x) #less than full rank, last column dropped
```

---

eqwma	<i>Equally Weighted Moving Average (EqWMA) of the pth. exponentiated values</i>
-------	---

---

### Description

The function `eqwma` returns an Equally Weighted Moving Average (EqWMA) of the `pth.` exponentiated values lagged `k` times (the default of `k` is 1). Optionally, the absolute values are computed before averaging if `abs=TRUE`, and the natural log of the values is returned if `log=TRUE`. The function `leqwma` is a wrapper to `eqwma` with `abs=TRUE` and `log=TRUE`.

If `x` is financial return (possibly mean-corrected) and `p=2`, then this gives the so-called 'historical' model, also known as an integrated ARCH model where the ARCH coefficients all have the same

value with sum equal to one. In the log-variance specification the lag of  $\log(\text{EqWMA})$  is thus a financial volatility proxy. It may be an imperfect proxy compared with high-frequency data (which can also be included as regressors), but - in contrast to high-frequency data - is always available and easy to compute.

### Usage

```
eqwma(x, length=5, k=1, p=1, abs=FALSE, log=FALSE, as.vector=FALSE,
      lag=NULL, start=NULL)
leqwma(x, length=5, k=1, p=2, as.vector=FALSE, lag=NULL, start=NULL)
```

### Arguments

x	numeric vector, time-series or zoo object
length	integer or vector of integers each equal to or greater than 1. The length or lengths of the moving window or windows of averages
k	integer that determines how many periods the term(s) should be lagged. If 0 (or smaller), then the moving averages are not lagged
p	numeric value. The exponent $p$ in $x^p$ when <code>abs=FALSE</code> , and in $\text{abs}(x)^p$ when <code>abs=TRUE</code>
log	logical with default FALSE. If TRUE, then the logarithm of the moving average is returned
abs	logical with default FALSE. If TRUE, then $x$ is transformed to absolute values before $x$ is exponentiated
as.vector	logical with default FALSE. If TRUE, and if <code>length(length)==1</code> , then the result is returned as a vector. Otherwise the returned value is always a matrix
lag	deprecated
start	deprecated

### Details

The intended primary use of `eqwma` is to construct mixed frequency regressors for the mean specification of an `arx` model.

The intended primary use of `leqwma` is to construct volatility proxies for the log-variance specification in an `arx` model. In the latter case, the default is the lagged log of an equally weighted moving average of the squared residuals, where each average is made up of  $m$  observations. This is equivalent to an integrated ARCH( $p$ ) model where the  $p$  coefficients are all equal. For further details on the use of  $\log(\text{EqWMA})$  as a volatility proxy, see Sucarrat and Escribano (2012).

### Value

numeric matrix, vector or `zoo` object

### Author(s)

Genaro Sucarrat, <http://www.sucarrat.net/>

## References

Genaro Sucarrat and Alvaro Escribano (2012): 'Automated Financial Model Selection: General-to-Specific Modelling of the Mean and Volatility Specifications', *Oxford Bulletin of Economics and Statistics* 74, Issue no. 5 (October), pp. 716-735

Pretis, Felix, Reade, James and Sucarrat, Genaro (2018): 'Automated General-to-Specific (GETS) Regression Modeling and Indicator Saturation for Outliers and Structural Breaks'. *Journal of Statistical Software* 86, Number 3, pp. 1-44

## See Also

[zoo](#), [arx](#), [getsm](#), [getsv](#)

## Examples

```
##generate an iid normal series:
set.seed(123)
x <- rnorm(100)

##compute lag of EqWMA(20) for x^2:
eqwma(x, p=2)

##compute lag of EqWMA(5) and lag of EqWMA(10) for x:
eqwma(x, length=c(5,10))

##compute lag of log(EqWMA(20)) for x^2:
leqwma(x)

#compute lag of log(EqWMA(5)) and lag of log(EqWMA(8))
#for abs(x)^2:
leqwma(x, length=c(4,8))
```

---

ES

*Conditional Value-at-Risk (VaR) and Expected Shortfall (ES)*

---

## Description

Extract the in-sample conditional Value-at-Risk, or the in-sample conditional Expected Shortfall for the chosen risk level(s).

## Usage

```
ES(object, level=0.99, type=7, ...)
VaR(object, level=0.99, type=7, ...)
```

**Arguments**

object	an <code>arx</code> or <code>gets</code> object
level	the risk level(s), must be between 0 and 1
type	the method used to compute the empirical quantiles of the standardised residuals
...	arguments passed on (currently not used)

**Value**

A vector or matrix containing either the conditional Value-at-Risk (VaR) or the conditional Expected Shortfall (ES) for the chosen risk level(s).

**Author(s)**

Genaro Sucarrat, <http://www.sucarrat.net/>

**See Also**

[arx](#), [getsm](#), [getsv](#)

**Examples**

```
##generate random variates, estimate model:
y <- rnorm(50)
mymodel <- arx(y, arch=1)

##extract 99% expected shortfall:
ES(mymodel)

##extract 99%, 95% and 90% expected shortfalls:
ES(mymodel, level=c(0.99, 0.95, 0.9))

##extract 99% value-at-risk:
VaR(mymodel)

##extract 99%, 95% and 90% values-at-risk:
VaR(mymodel, level=c(0.99, 0.95, 0.9))
```

---

eviews

*Exporting results to EViews and STATA*

---

**Description**

Functions that facilitate the export of results to the commercial econometric softwares EViews and STATA, respectively.

**Usage**

```
eviews(object, file=NULL, print=TRUE, return=FALSE)
stata(object, file=NULL, print=TRUE, return=FALSE)
```



**Arguments**

object	an arx, gets or isat object
file	filename, i.e. the destination of the exported data
print	logical. If TRUE, then the estimation code in EViews (or STATA) is printed
return	logical. If TRUE, then a list is returned

**Value**

Either printed text or a list (if return=TRUE)

**Author(s)**

Genaro Sucarrat, <http://www.sucarrat.net/>

**See Also**

[arx](#), [getsm](#), [getsv](#), [isat](#)

**Examples**

```
##simulate random variates, estimate model:
y <- rnorm(30)
mX <- matrix(rnorm(30*2), 30, 2)
mymod <- arx(y, mc=TRUE, mxreg=mX)

##print EViews code:
eviews(mymod)

##print Stata code:
stata(mymod)
```

---

gets

*General-to-Specific (GETS) Modelling*

---

**Description**

Generic function that enables new GETS and ISAT methods for new classes

**Usage**

```
gets(x, ...)
```

**Arguments**

x	an object used to select a method
...	further arguments passed to or from other methods

**Author(s)**

Genaro Sucarrat, <http://www.sucarrat.net/>

---

getsFun

*General-to-Specific (GETS) modelling function*

---

**Description**

Auxiliary function (i.e. not intended for the average user) that enables fast and efficient GETS-modelling with user-specified estimators and models, and user-specified diagnostics and goodness-of-fit criteria. The function is called by and relied upon by `getsv` and `isat`, and in future versions of the package the same will be the case for `getsm`.

**Usage**

```
getsFun(y, x, untransformed.residuals=NULL,
        user.estimate=list(name="ols"), gum.result=NULL, t.pval=0.05,
        wald.pval=t.pval, do.pet=TRUE, ar.LjungB=NULL, arch.LjungB=NULL,
        normality.JarqueB=NULL, user.diagnostics=NULL,
        gof.function=list(name="infocrit", method="sc"),
        gof.method=c("min", "max"), keep=NULL, include.gum=FALSE,
        include.1cut=FALSE, include.empty=FALSE, max.paths=NULL, turbo=FALSE,
        tol=1e-07, LAPACK=FALSE, max.regs=NULL, print.searchinfo=TRUE,
        alarm=FALSE)
```

**Arguments**

<code>y</code>	a numeric vector (with no missing values, i.e. no non-numeric 'holes')
<code>x</code>	a matrix with <code>NROW(x)</code> equal to <code>NROW(y)</code> , or <code>NULL</code>
<code>untransformed.residuals</code>	<code>NULL</code> (default) or, when <code>ols</code> is used with <code>method=6</code> , a numeric vector containing the untransformed residuals
<code>user.estimate</code>	a list. The first item should be named <code>name</code> and contain the name (a character) of the estimation function. Additional items, if any, in the list <code>user.estimate</code> are passed on as arguments to the estimator. The value returned by the estimator should be a list, see details
<code>gum.result</code>	a list with the estimation results of the General Unrestricted Model (GUM), or <code>NULL</code> (default). If the estimation results of the GUM are already available, then re-estimation of the GUM is skipped if the estimation results are provided via this argument
<code>t.pval</code>	numeric value between 0 and 1. The significance level used for the two-sided regressor significance t-tests
<code>wald.pval</code>	numeric value between 0 and 1. The significance level used for the Parsimonious Encompassing Tests (PETs). By default, it is the same as <code>t.pval</code>

<code>do.pet</code>	logical. If TRUE (default), then a Parsimonious Encompassing Test (PET) against the GUM is undertaken at each regressor removal for the joint significance of all the deleted regressors along the current path. If FALSE, then a PET is not undertaken at each regressor removal
<code>ar.LjungB</code>	a two element vector or NULL. In the former case, the first element contains the AR-order, the second element the significance level. If NULL, then a test for autocorrelation is not conducted
<code>arch.LjungB</code>	a two element vector or NULL. In the former case, the first element contains the ARCH-order, the second element the significance level. If NULL, then a test for ARCH is not conducted
<code>normality.JarqueB</code>	NULL or a value between 0 and 1. In the latter case, a test for non-normality is conducted using a significance level equal to <code>normality.JarqueB</code> . If NULL, then no test for non-normality is conducted
<code>user.diagnostics</code>	NULL (default) or a list with two entries, <code>name</code> and <code>pval</code> . The first item ( <code>name</code> ) should contain the name of the user-defined function, and must be of class character. The the second item should contain the chosen significance level or levels, i.e. either a scalar or a vector of length equal to the number of p-values returned by the user-defined diagnostics function, see details
<code>gof.function</code>	a list. The first item should be named <code>name</code> and contain the name (a character) of the Goodness-of-Fit (GOF) function used. Additional items in the list <code>gof.function</code> are passed on as arguments to the GOF-function. The value returned by the GOF-function should be a numeric value (of length 1)
<code>gof.method</code>	a character. Determines whether the best Goodness-of-Fit is a minimum or maximum
<code>keep</code>	NULL or an integer vector that indicates which regressors to be excluded from removal in the search
<code>include.gum</code>	logical. If TRUE, then the GUM (i.e. the starting model) is included among the terminal models. If FALSE (default), then the GUM is not included
<code>include.1cut</code>	logical. If TRUE, then the 1-cut model is added to the list of terminal models. If FALSE (default), then the 1-cut is not added, unless it is a terminal model in one of the paths
<code>include.empty</code>	logical. If TRUE, then the empty model is added to the list of terminal models. If FALSE (default), then the empty model is not added, unless it is a terminal model in one of the paths
<code>max.paths</code>	NULL (default) or an integer greater than 0. If NULL, then there is no limit to the number of paths. If an integer (e.g. 1), then this integer constitutes the maximum number of paths searched (e.g. a single path)
<code>turbo</code>	logical. If TRUE, then (parts of) paths are not searched twice (or more) unnecessarily, thus yielding a significant potential for speed-gain. However, the checking of whether the search has arrived at a point it has already been comes with a slight computational overhead. Accordingly, if <code>turbo=TRUE</code> , then the total search time might in fact be higher than if <code>turbo=FALSE</code> . This happens if estimation is very fast, say, less than quarter of a second. Hence the default is FALSE

<code>tol</code>	numeric value (default = $1e-07$ ). The tolerance for detecting linear dependencies in the columns of the variance-covariance matrix when computing the Wald-statistic used in the Parsimonious Encompassing Tests (PETs), see the <a href="#"><code>qr.solve</code></a> function
LAPACK	currently not used
<code>max.regs</code>	integer. The maximum number of regressions along a deletion path. It is not recommended that this is altered
<code>print.searchinfo</code>	logical. If TRUE (default), then a print is returned whenever simplification along a new path is started
<code>alarm</code>	logical. If TRUE, then a sound or beep is emitted (in order to alert the user) when the model selection ends

### Details

The value returned by the estimator specified in `user.estimator` should be a list containing at least six items: "coefficients", "df", "vcov", "logl", "n" and "k". The item "coefficients" should be a vector of length  $NCOL(x)$  containing the estimated coefficients. The item named "df" is used to compute the p-values associated with the t-statistics, i.e. `coef/std.err`. The item named "vcov" contains the (symmetric) coefficient-covariance matrix of the estimated coefficients. The items "logl" (the log-likelihood), "n" (the number of observations) and "k" (the number of estimated parameters; not necessarily equal to the number of coefficients) are used to compute the information criterion. Finally, the estimator MUST be able to handle NULL regressor-matrices (i.e. `is.null(x)=TRUE` or  $NCOL(x)=0$ ). In this case, then the first three items (i.e. "coefficients", "df" and "vcov") can - and should - be NULL. The argument `user.diagnostics` enables the user to specify additional - or alternative - diagnostics, see [diagnostics](#).

### Value

The returned value, a [list](#), depends on the `user.estimator`. For the default, see [ols](#) with `method=3`.

### Author(s)

Genaro Sucarrat, <http://www.sucarrat.net/>

### References

- C. Jarque and A. Bera (1980): 'Efficient Tests for Normality, Homoscedasticity and Serial Independence'. *Economics Letters* 6, pp. 255-259
- G. Ljung and G. Box (1979): 'On a Measure of Lack of Fit in Time Series Models'. *Biometrika* 66, pp. 265-270

### See Also

[ols](#), [diagnostics](#), [infocrit](#), [getsv](#)

## Examples

```
##aim: do gets on the x-part (i.e. the covariates) of an arma-x model.
##create the user-defined estimator (essentially consists of adding,
##renaming and re-organising the items returned by the chosen
##estimator):
myEstimator <- function(y, x)
{
  tmp <- arima(y, order=c(1,0,1), xreg=x)

  #rename and re-organise:
  result <- list()
  result$coefficients <- tmp$coef[-c(1:3)]
  result$vcov <- tmp$var.coef
  result$vcov <- result$vcov[-c(1:3),-c(1:3)]
  result$loglik <- tmp$loglik
  result$n <- tmp$nobs
  result$k <- NCOL(x)
  result$df <- result$n - result$k

  return(result)
}

##generate some data:
##a series w/structural break and eleven step-dummies near the break
set.seed(123)
eps <- arima.sim(list(ar=0.4, ma=0.1), 60)
x <- coredata(sim(eps, which.ones=25:35)) #eleven step-dummies
y <- 4*x[, "sis30"] + eps #create shift upwards at observation 30
plot(y)

##estimate the gum and then do gets in a single step:
getsFun(y, x, user.estimator=list(name="myEstimator"))

##estimate the gum and then do gets in two steps:
mygum <- myEstimator(y,x)
#getsFun(y, x, user.estimator=list(name="myEstimator"), gum.result=mygum)
```

---

getsm

*General-to-Specific (GETS) Modelling of an AR-X model with log-ARCH-X errors*

---

## Description

The starting model, an object of the 'arx' class, is referred to as the General Unrestricted Model (GUM). The `getsm` function undertakes multi-path GETS modelling of the mean specification, whereas `getsv` does the same for the log-variance specification. The diagnostic tests are undertaken on the standardised residuals, and the `keep` option enables regressors to be excluded from possible removal.

**Usage**

```
##gets of mean specification:
getsm(object, t.pval=0.05, wald.pval=t.pval, vcov.type=NULL, do.pet=TRUE,
      ar.LjungB=list(lag=NULL, pval=0.025), arch.LjungB=list(lag=NULL, pval=0.025),
      normality.JarqueB=NULL, user.diagnostics=NULL,
      info.method=c("sc","aic","hq"), keep=NULL, include.gum=FALSE,
      include.empty=FALSE, max.paths=NULL, max.regs=NULL, zero.adj=NULL,
      vc.adj=NULL, verbose=TRUE, print.searchinfo=TRUE, estimate.specific=TRUE,
      plot=NULL, alarm=FALSE)

##gets of variance specification:
getsv(object, t.pval=0.05, wald.pval=t.pval, do.pet=TRUE,
      ar.LjungB=list(lag=NULL, pval=0.025), arch.LjungB=list(lag=NULL, pval=0.025),
      normality.JarqueB=NULL, user.diagnostics=NULL,
      info.method=c("sc","aic","hq"), keep=c(1), include.gum=FALSE,
      include.1cut=FALSE, include.empty=FALSE, max.paths=NULL, turbo=FALSE,
      max.regs=NULL, zero.adj=NULL, vc.adj=NULL, print.searchinfo=TRUE,
      estimate.specific=TRUE, plot=NULL, alarm=FALSE)
```

**Arguments**

object	an object of class 'arx'
t.pval	numeric value between 0 and 1. The significance level used for the two-sided regressor significance t-tests
wald.pval	numeric value between 0 and 1. The significance level used for the Parsimonious Encompassing Tests (PETs). By default, it is the same as t.pval
vcov.type	the type of variance-covariance matrix used. If NULL (default), then the type used in the estimation of the 'arx' object is used. This can be overridden by either "ordinary" (i.e. the ordinary variance-covariance matrix) or "white" (i.e. the White (1980) heteroscedasticity robust variance-covariance matrix)
do.pet	logical. If TRUE (default), then a Parsimonious Encompassing Test (PET) against the GUM is undertaken at each regressor removal for the joint significance of all the deleted regressors along the current path. If FALSE, then a PET is not undertaken at each regressor removal
ar.LjungB	a two-item list with names lag and pval, or NULL. In the former case lag contains the order of the Ljung and Box (1979) test for serial correlation in the standardised residuals, and pval contains the significance level. If lag=NULL (default), then the order used is that of the estimated 'arx' object. If ar.Ljungb=NULL, then the standardised residuals are not checked for serial correlation
arch.LjungB	a two-item list with names lag and pval, or NULL. In the former case, lag contains the order of the Ljung and Box (1979) test for serial correlation in the squared standardised residuals, and pval contains the significance level. If lag=NULL (default), then the order used is that of the estimated 'arx' object. If arch.Ljungb=NULL, then the standardised residuals are not checked for ARCH

<code>normality.JarqueB</code>	a value between 0 and 1, or NULL. In the former case, the Jarque and Bera (1980) test for non-normality is conducted using a significance level equal to the numeric value. If NULL, then no test for non-normality is undertaken
<code>user.diagnostics</code>	NULL or a <code>list</code> with two entries, <code>name</code> and <code>pval</code> , see the <code>user.fun</code> argument in <a href="#">diagnostics</a>
<code>info.method</code>	character string, "sc" (default), "aic" or "hq", which determines the information criterion to be used when selecting among terminal models. The abbreviations are short for the Schwarz or Bayesian information criterion (sc), the Akaike information criterion (aic) and the Hannan-Quinn (hq) information criterion
<code>keep</code>	the regressors to be excluded from removal in the specification search. Note that <code>keep=c(1)</code> is obligatory when using <code>getsv</code> . This excludes the log-variance intercept from removal. The regressor numbering is contained in the <code>reg.no</code> column of the GUM
<code>include.gum</code>	logical. If TRUE, then the GUM (i.e. the starting model) is included among the terminal models. If FALSE (default), then the GUM is not included
<code>include.1cut</code>	logical. If TRUE, then the 1-cut model is added to the list of terminal models. If FALSE (default), then the 1-cut is not added, unless it is a terminal model in one of the paths
<code>include.empty</code>	logical. If TRUE, then an empty model is included among the terminal models, if it passes the diagnostic tests, even if it is not equal to one of the terminals. If FALSE (default), then the empty model is not included (unless it is one of the terminals)
<code>max.paths</code>	NULL (default) or an integer greater than 0. If NULL, then there is no limit to the number of paths. If an integer (e.g. 1), then this integer constitutes the maximum number of paths searched (e.g. a single path)
<code>turbo</code>	logical. If TRUE, then (parts of) paths are not searched twice (or more) unnecessarily, thus yielding a significant potential for speed-gain. However, the checking of whether the search has arrived at a point it has already been comes with a slight computational overhead. Accordingly, if <code>turbo=TRUE</code> , then the total search time might in fact be higher than if <code>turbo=FALSE</code> . This happens if estimation is very fast, say, less than quarter of a second. Hence the default is FALSE
<code>max.regs</code>	integer. The maximum number of regressions along a deletion path. It is not recommended that this is altered
<code>zero.adj</code>	numeric value between 0 and 1. The quantile adjustment for zero values. The default 0.1 means the zero residuals are replaced by the 10 percent quantile of the absolute residuals before taking the logarithm
<code>vc.adj</code>	logical. If TRUE (default), then the log-variance intercept is adjusted by the estimate of $E[\ln(z^2)]$ . This adjustment is needed for the conditional scale of $e$ to be equal to the conditional standard deviation. If FALSE, then the log-variance intercept is not adjusted
<code>verbose</code>	logical. TRUE (default) returns (slightly) more output than FALSE

<code>print.searchinfo</code>	logical. If TRUE (default), then a print is returned whenever simplification along a new path is started
<code>estimate.specific</code>	logical. IF TRUE (default), then the specific model is estimated after model selection
<code>plot</code>	NULL or logical. If TRUE, then the fitted values and the residuals of the final model are plotted after model selection. If FALSE, then they are not. If NULL (default), then the value set by <code>options</code> determines whether a plot is produced or not
<code>alarm</code>	logical. If TRUE, then a sound or beep is emitted (in order to alert the user) when the model selection ends

### Details

See Pretis, Reade and Sucarrat (2018)

### Value

A list of class 'gets'

### Author(s)

Genaro Sucarrat, <http://www.sucarrat.net/>

### References

- C. Jarque and A. Bera (1980): 'Efficient Tests for Normality, Homoscedasticity and Serial Independence'. *Economics Letters* 6, pp. 255-259
- G. Ljung and G. Box (1979): 'On a Measure of Lack of Fit in Time Series Models'. *Biometrika* 66, pp. 265-270
- Pretis, Felix, Reade, James and Sucarrat, Genaro (2018): 'Automated General-to-Specific (GETS) Regression Modeling and Indicator Saturation for Outliers and Structural Breaks'. *Journal of Statistical Software* 86, Number 3, pp. 1-44

### See Also

Extraction functions: [coef.gets](#), [fitted.gets](#), [paths](#), [plot.gets](#), [print.gets](#), [residuals.gets](#), [summary.gets](#), [terminals](#), [vcov.gets](#)

Related functions: [arx](#), [isat](#), [eqwma](#), [leqwma](#), [zoo](#), [getsFun](#)

### Examples

```
##Simulate from an AR(1):
set.seed(123)
y <- arima.sim(list(ar=0.4), 80)

##Simulate four independent Gaussian regressors:
```



```

xregs <- matrix(rnorm(2*80), 80, 2)

##estimate an AR(2) with intercept and four conditioning
##regressors in the mean, and a log-ARCH(3) with log(xregs^2) as
##regressors in the log-variance:
gum01 <- arx(y, mc=TRUE, ar=1:2, mxreg=xregs, arch=1:3,
  vxreg=log(xregs^2))

##GETS model selection of the mean:
meanmod01 <- getsm(gum01)

##GETS model selection of the log-variance:
varmod01 <- getsv(gum01)

##GETS model selection of the mean with the mean intercept
##excluded from removal:
meanmod02 <- getsm(gum01, keep=1)

##GETS model selection of the mean with non-default
#serial-correlation diagnostics settings:
meanmod03 <- getsm(gum01, ar.LjungB=list(pval=0.05))

##GETS model selection of the mean with very liberal
##(20 percent) significance levels:
meanmod04 <- getsm(gum01, t.pval=0.2)

##GETS model selection of log-variance with all the
##log-ARCH terms excluded from removal:
varmod03 <- getsv(gum01, keep=2:4)

```

---

hpdata

*Hoover and Perez (1999) data*


---

## Description

Data used by Hoover and Perez (1999) in their evaluation of General-to-Specific (GETS) modelling. A detailed description of the data is found in their Table 1 (page 172). The data are quarterly, comprise 20 variables (the first variable is the quarterly index) and runs from 1959:1 to 1995:1. This corresponds to 145 observations. The original source of the data is Citibank.

## Usage

```
data(hpdata)
```

## Format

Date a factor that contains the (quarterly) dates of the observations  
DCOINC index of four coincident indicators

GD GNP price deflator  
GGEQ government purchases of goods and services  
GGFEQ federal purchases of goods and services  
GGFR federal government receipts  
GNPQ GNP  
GYDQ disposable personal income  
GPIQ gross private domestic investment  
FMRRR total member bank reserves  
FMBASE monetary base (federal reserve bank of St. Louis)  
FM1DQ M1  
FM2DQ M2  
FSDJ Dow Jones stock price  
FYAAAC Moody's AAA corporate bond yield  
LHC labour force (16 years+, civilian)  
LHUR unemployment rate  
MU unfilled orders (manufacturing, all industries)  
MO new orders (manufacturing, all industries)  
GCQ personal consumption expenditure

### Details

The data have been used for comparison and illustration of GETS model selection in several studies of the GETS methodology, including Hendry and Krolzig (1999, 2005), Doornik (2009) and Sucarrat and Escribano (2012).

### Source

Retrieved 14 October 2014 from: <http://www.csus.edu/indiv/p/perezs/Data/data.htm>

### References

- David F. Hendry and Hans-Martin Krolzig (1999): 'Improving on 'Data mining reconsidered' by K.D. Hoover and S.J Perez', *Econometrics Journal*, Vol. 2, pp. 202-219.
- David F. Hendry and Hans-Martin Krolzig (2005): 'The properties of automatic Gets modelling', *Economic Journal* 115, C32-C61.
- Jurgen Doornik (2009): 'Autometrics', in Jennifer L. Castle and Neil Shephard (eds), 'The Methodology and Practice of Econometrics: A Festschrift in Honour of David F. Hendry', Oxford University Press, Oxford, pp. 88-121.
- Pretis, Felix, Reade, James and Sucarrat, Genaro (2018): 'Automated General-to-Specific (GETS) Regression Modeling and Indicator Saturation for Outliers and Structural Breaks'. *Journal of Statistical Software* 86, Number 3, pp. 1-44.

**Examples**

```
##load Hoover and Perez (1999) data:
data(hpdata)

##make quarterly data-matrix of zoo type:
newhpdata <- zooreg(hpdata[,-1], start=c(1959,1), frequency=4)

##plot data:
plot(newhpdata)

##transform data to log-differences in percent:
dloghpdata <- diff(log(newhpdata))*100

##plot log-differenced data:
plot(dloghpdata)
```

iim

*Make Indicator Matrices (Impulses, Steps, Trends)***Description**

Auxiliary functions to make, respectively, matrices of impulse indicators (`iim`), step indicators (`sim`) and trend indicators (`tim`)

**Usage**

```
##make matrix of impulse indicators:
iim(x, which.ones = NULL)

##make matrix of step indicators:
sim(x, which.ones = NULL)

##make matrix of trend indicators:
tim(x, which.ones = NULL, log.trend = FALSE)
```

**Arguments**

<code>x</code>	either an integer (the length of the series in question) or a series (a vector or matrix) from which to use the time-series index to make indicators of
<code>which.ones</code>	the locations of the impulses. If <code>NULL</code> (the default), then all impulses are returned
<code>log.trend</code>	logical. If <code>TRUE</code> , then the natural log is applied on the trends

**Details**

If `x` is a series or vector of observations, then the index of `x` will be used for the labelling of the impulses, and in the returned `zoo` object.

Note: For `sim` and `tim` the first indicator is removed, since it is exactly colinear with the others.

**Value**

A `zoo` matrix containing the impulses

**Author(s)**

Genaro Sucarrat, <http://www.sucarrat.net/>

**See Also**

[isat](#), [zoo](#)

**Examples**

```
##generate series:
y <- rnorm(40)

##make matrix of impulse indicators:
mIIM <- iim(40)

##make matrix of step-indicators, but only every third:
mSIM <- sim(y, which.ones=seq(1,40,3))

##give quarterly time-series attributes to y-series:
y <- zooreg(y, frequency=4, end=c(2015,4))

##make matrix of trend-indicators with quarterly labels:
mTIM <- tim(y)
```

---

infldata

*Quarterly Norwegian year-on-year CPI inflation*

---

**Description**

Quarterly Norwegian year-on-year CPI inflation from 1989(1) to 2015(4).

**Usage**

```
data("infldata")
```

**Format**

A data frame with 108 observations on the following 5 variables:

`date` a factor containing the dates

`infl` year-on-year inflation

`q2dum` a dummy variable equal to 1 in quarter 2 and 0 otherwise

`q3dum` a dummy variable equal to 1 in quarter 3 and 0 otherwise

`q4dum` a dummy variable equal to 1 in quarter 4 and 0 otherwise

**Source**

Statistics Norway (SSB): <http://www.ssb.no/>. The raw data comprise monthly CPI data obtained via <http://www.ssb.no/tabell/08183/>.

**References**

Pretis, Felix, Reade, James and Sucarrat, Genaro (2018): 'Automated General-to-Specific (GETS) Regression Modeling and Indicator Saturation for Outliers and Structural Breaks'. *Journal of Statistical Software* 86, Number 3, pp. 1-44

**Examples**

```
data(infldata)
infldata <- zooreg(infldata[,-1], frequency=4, start=c(1989,1))
plot(infldata[, "infl"])
```

---

infocrit

*Computes the Average Value of an Information Criterion*


---

**Description**

Given a log-likelihood, the number of observations and the number of estimated parameters, the average value of a chosen information criterion is computed. This facilitates comparison of models that are estimated with a different number of observations, e.g. due to different lags.

**Usage**

```
infocrit(x, method=c("sc", "aic", "aicc", "hq"))

info.criterion(logl, n=NULL, k=NULL, method=c("sc", "aic", "aicc", "hq"))
```

**Arguments**

x	a list that contains, at least, three items: logl (a numeric, the log-likelihood), k (a numeric, usually the number of estimated parameters) and n (a numeric, the number of observations)
method	character, either "sc" (default), "aic", "aicc" or "hq"
logl	numeric, the value of the log-likelihood
n	integer, number of observations
k	integer, number of parameters

**Details**

Contrary to **AIC** and **BIC**, `info.criterion` computes the average criterion value (i.e. division by the number of observations). This facilitates comparison of models that are estimated with a different number of observations, e.g. due to different lags.

**Value**

infocrit: a numeric (i.e. the value of the chosen information criterion)  
 info.criterion: a list with elements

method	type of information criterion
n	number of observations
k	number of parameters
value	the value on the information criterion

**Author(s)**

Genaro Sucarrat, <http://www.sucarrat.net/>

**References**

- H. Akaike (1974): 'A new look at the statistical model identification'. IEEE Transactions on Automatic Control 19, pp. 716-723
- E. Hannan and B. Quinn (1979): 'The determination of the order of an autoregression'. Journal of the Royal Statistical Society B 41, pp. 190-195
- C.M. Hurvich and C.-L. Tsai (1989): 'Regression and Time Series Model Selection in Small Samples'. Biometrika 76, pp. 297-307
- Pretis, Felix, Reade, James and Sucarrat, Genaro (2018): 'Automated General-to-Specific (GETS) Regression Modeling and Indicator Saturation for Outliers and Structural Breaks'. Journal of Statistical Software 86, Number 3, pp. 1-44
- G. Schwarz (1978): 'Estimating the dimension of a model'. The Annals of Statistics 6, pp. 461-464

---

isat

*Indicator Saturation*

---

**Description**

The `isat` function undertakes multi-path indicator saturation to detect outliers and mean-shifts using impulses (IIS), step-shifts (SIS), or trend-indicators (TIS). Indicators are partitioned into blocks and selected over at a chosen level of significance (`t.pval`) using the `getsm` function.

**Usage**

```
isat(y, mc=TRUE, ar=NULL, ewma=NULL, mxreg=NULL, iis=FALSE, sis=TRUE,
     tis=FALSE, uis=FALSE, blocks=NULL, ratio.threshold=0.8, max.block.size=30,
     t.pval=0.001, wald.pval=t.pval,
     vcov.type= c("ordinary", "white", "newey-west"), do.pet=FALSE, ar.LjungB=NULL,
     arch.LjungB=NULL, normality.JarqueB=NULL, info.method=c("sc", "aic", "hq"),
     user.diagnostics=NULL, user.estimator=NULL, gof.function=NULL,
     gof.method = c("min", "max"), include.gum=NULL, include.1cut=FALSE,
     include.empty=FALSE, max.paths=NULL, parallel.options=NULL, turbo=FALSE,
     tol=1e-07, LAPACK=FALSE, max.regs=NULL, print.searchinfo=TRUE, plot=NULL,
     alarm=FALSE)
```

**Arguments**

<code>y</code>	numeric vector, time-series or <code>zoo</code> object. Missing values in the beginning and at the end of the series is allowed, as they are removed with the <code>na.trim</code> command
<code>mc</code>	logical. TRUE (default) includes an intercept in the mean specification, whereas FALSE does not
<code>ar</code>	integer vector, say, <code>c(2,4)</code> or <code>1:4</code> . The AR-lags to include in the mean specification
<code>ewma</code>	either NULL (default) or a list with arguments sent to the <code>eqwma</code> function. In the latter case a lagged moving average of <code>y</code> is included as a regressor
<code>mxreg</code>	numeric vector or matrix, say, a <code>zoo</code> object, of conditioning variables. Note that missing values in the beginning or at the end of the series is allowed, as they are removed with the <code>na.trim</code> command. Note also that, if both <code>y</code> and <code>mxreg</code> are <code>zoo</code> objects, then their samples are chosen to match
<code>iis</code>	logical. If TRUE, impulse indicator saturation is performed.
<code>sis</code>	logical. If TRUE, step indicator saturation is performed.
<code>tis</code>	logical. If TRUE, trend indicator saturation is performed.
<code>uis</code>	a matrix of regressors, or a list of matrices.
<code>blocks</code>	NULL (default), an integer (the number of blocks) or a user-specified list that indicates how blocks should be put together. If NULL, then the number of blocks is determined automatically
<code>ratio.threshold</code>	Minimum ratio of variables in each block to total observations to determine the block size, default=0.8. Only relevant if <code>blocks = NULL</code>
<code>max.block.size</code>	Maximum size of block of variables to be selected over, default=30. Block size used is the maximum of given by either the <code>ratio.threshold</code> and <code>max.block.size</code>
<code>t.pval</code>	numeric value between 0 and 1. The significance level used for the two-sided regressor significance t-tests
<code>wald.pval</code>	numeric value between 0 and 1. The significance level used for the Parsimonious Encompassing Tests (PETs)
<code>vcov.type</code>	the type of variance-covariance matrix used. If NULL (default), then the type used is that of the 'arx' object. This can be overridden by either "ordinary" (i.e. the ordinary variance-covariance matrix) or "white" (i.e. the White (1980) heteroscedasticity robust variance-covariance matrix)
<code>do.pet</code>	logical. If TRUE, then a Parsimonious Encompassing Test (PET) against the GUM is undertaken at each regressor removal for the joint significance of all the deleted regressors along the current path. If FALSE (default), then a PET is not undertaken at each regressor removal. By default, the numeric value is the same as that of <code>t.pval</code>
<code>ar.LjungB</code>	a two-item list with names <code>lag</code> and <code>pval</code> , or NULL (default). In the former case <code>lag</code> contains the order of the Ljung and Box (1979) test for serial correlation in the standardised residuals, and <code>pval</code> contains the significance level. If <code>lag=NULL</code> (default), then the order used is that of the estimated 'arx' object. If <code>ar.LjungB=NULL</code> , then the standardised residuals are not checked for serial correlation

<code>arch.LjungB</code>	a two-item list with names <code>lag</code> and <code>pval</code> , or <code>NULL</code> (default). In the former case, <code>lag</code> contains the order of the Ljung and Box (1979) test for serial correlation in the squared standardised residuals, and <code>pval</code> contains the significance level. If <code>lag=NULL</code> (default), then the order used is that of the estimated 'arx' object. If <code>arch.Ljungb=NULL</code> , then the standardised residuals are not checked for ARCH
<code>normality.JarqueB</code>	<code>NULL</code> (the default) or a value between 0 and 1. In the latter case, a test for non-normality is conducted using a significance level equal to <code>normality.JarqueB</code> . If <code>NULL</code> , then no test for non-normality is conducted
<code>info.method</code>	character string, "sc" (default), "aic" or "hq", which determines the information criterion to be used when selecting among terminal models. The abbreviations are short for the Schwarz or Bayesian information criterion (sc), the Akaike information criterion (aic) and the Hannan-Quinn (hq) information criterion
<code>user.diagnostics</code>	<code>NULL</code> or a <a href="#">list</a> with two entries, <code>name</code> and <code>pval</code> , see the <code>user.fun</code> argument in <a href="#">diagnostics</a>
<code>user.estimator</code>	<code>NULL</code> or a <a href="#">list</a> with at least one entry, <code>name</code> , see the <code>user.estimator</code> argument in <a href="#">getsFun</a>
<code>gof.function</code>	<code>NULL</code> or a <a href="#">list</a> with at least one entry, <code>name</code> , see the <code>user.estimator</code> argument in <a href="#">getsFun</a>
<code>gof.method</code>	<code>NULL</code> or a character that determines whether the best Goodness-of-Fit is a minimum or maximum
<code>include.gum</code>	ignored (temporarily deprecated)
<code>include.1cut</code>	logical. If <code>TRUE</code> , then the 1-cut model is included among the terminal models, if it passes the diagnostic tests, even if it is not equal to one of the terminals. If <code>FALSE</code> (default), then the 1-cut model is not included (unless it is one of the terminals)
<code>include.empty</code>	logical. If <code>TRUE</code> , then an empty model is included among the terminal models, if it passes the diagnostic tests, even if it is not equal to one of the terminals. If <code>FALSE</code> (default), then the empty model is not included (unless it is one of the terminals)
<code>max.paths</code>	<code>NULL</code> (default) or an integer indicating the maximum number of paths to search
<code>parallel.options</code>	<code>NULL</code> or an integer, i.e. the number of cores/threads to be used for parallel computing (implemented w/ <code>makeCluster</code> and <code>parLapply</code> )
<code>turbo</code>	logical. If <code>TRUE</code> , then (parts of) paths are not searched twice (or more) unnecessarily, thus yielding a significant potential for speed-gain. However, the checking of whether the search has arrived at a point it has already been comes with a slight computational overhead. Accordingly, if <code>turbo=TRUE</code> , then the total search time might in fact be higher than if <code>turbo=FALSE</code> . This happens if estimation is very fast, say, less than quarter of a second. Hence the default is <code>FALSE</code>
<code>tol</code>	numeric value (default = <code>1e-07</code> ). The tolerance for detecting linear dependencies in the columns of the regressors (see <a href="#">qr</a> function). Only used if LAPACK is <code>FALSE</code> (default)



LAPACK	logical. If TRUE, then use LAPACK. If FALSE (default), then use LINPACK (see <a href="#">qr</a> function)
max.regs	integer. The maximum number of regressions along a deletion path. It is not recommended that this is altered
print.searchinfo	logical. If TRUE (default), then a print is returned whenever simplification along a new path is started, and whenever regressors are dropped due to exact multicollinearity
plot	NULL or logical. If TRUE, then the fitted values and the residuals of the final model are plotted after model selection. If NULL (default), then the value set by <a href="#">options</a> determines whether a plot is produced or not.
alarm	logical. If TRUE, then a sound is emitted (in order to alert the user) when the model selection ends

### Details

Multi-path indicator saturation using impulses (IIS), step-shifts (SIS), or trend-indicators (TIS). Indicators are partitioned into sequential blocks (as of beta version 0.7) where the block intervals are defined by the ratio of variables to observations in each block and a specified maximum block size. Indicators are selected over using the [getsm](#) function. Retained indicators in each block are combined and re-selected over. Fixed covariates that are not selected over can be included in the regression model either in the mxreg matrix, or for auto-regressive terms through the ar specification. See Hendry, Johansen and Santos (2007) and Castle, Doornik, Hendry, and Pretis (2015)

### Value

A list of class 'gets'

### Author(s)

Felix Pretis, <http://www.felixpretis.org/>  
 James Reade, <https://sites.google.com/site/jjamesreade/>  
 Genaro Sucarrat, <http://www.sucarrat.net/>

### References

- Castle, Jennifer, L., Doornik, Jurgen, A., Hendry, David F., and Pretis, Felix (2015): 'Detecting Location Shifts during Model Selection by Step-Indicator Saturation', *Econometrics*, vol 3:2, 240-264.
- Hendry, David, F., Johansen, Soren, and Santos, Carlos (2007): 'Automatic selection of indicators in a fully saturated regression'. *Computational Statistics*, vol 23:1, pp.317-335.
- Pretis, Felix, Reade, James and Sucarrat, Genaro (2018): 'Automated General-to-Specific (GETS) Regression Modeling and Indicator Saturation for Outliers and Structural Breaks'. *Journal of Statistical Software* 86, Number 3, pp. 1-44

**See Also**

Extraction functions for 'gets' objects: [coef.gets](#), [fitted.gets](#), [paths](#), [plot.gets](#), [print.gets](#), [residuals.gets](#), [summary.gets](#), [terminals](#), [vcov.gets](#)

Related functions: [arx](#), [eqwma](#), [leqwma](#), [zoo](#), [getsFun](#)

**Examples**

```
##SIS using the Nile data
data(Nile)
isat(Nile, sis=TRUE, iis=FALSE, plot=TRUE, t.pval=0.005)

##SIS using the Nile data in an autoregressive model
#isat(Nile, ar=1:2, sis=TRUE, iis=FALSE, plot=TRUE, t.pval=0.005)

##HP Data
##load Hoover and Perez (1999) data:
#data(hpdata)

##make quarterly data-matrix of zoo type
##(GCQ = personal consumption expenditure):
#y <- zooreg(hpdata$GCQ, 1959, frequency=4)

##transform data to log-differences:
#dlogy <- diff(log(y))

##run isat with step impulse saturation on four
##lags and a constant 1 percent significance level:
#isat(dlogy, ar=1:4, sis=TRUE, t.pval =0.01)

##Example with additional covariates entering through mxreg:

##(GYDQ = disposable personal income):
#x <- zooreg(hpdata$GYDQ, 1959, frequency=4)

##transform data to log-differences:
#dlogx <- diff(log(x))

##run isat with step impulse saturation on four
##lags and a constant 1 percent significance level:
#isat(dlogy, mxreg=dlogx, ar=1:4, sis=TRUE, t.pval =0.01)
```

---

isatdates

---

*Extracting Indicator Saturation Breakdates*


---

**Description**

Takes an [isat](#) object and extracts the break dates together with their estimated coefficients.

**Usage**

```
isatdates(x)
```

**Arguments**

x                    an `isat` object

**Details**

The function extracts the breakdates determined by `isat` for `iis`, `sis`, and `tis`, together with their estimated coefficients and standard errors.

**Value**

Returns a list of three elements (one for `iis`, `sis`, and `tis`). Each element lists the name of the break variable, the time index of the break (labelled 'date'), the index of the break date, the estimated coefficient, the standard error of the estimated coefficient, as well as the corresponding t-statistic and p-value.

**Author(s)**

Felix Pretis, <http://www.felixpretis.org/>

**References**

Pretis, F., Reade, J., & Sucarrat, G. (2018). Automated General-to-Specific (GETS) regression modeling and indicator saturation methods for the detection of outliers and structural breaks. *Journal of Statistical Software*, 86(3).

**See Also**

`isat`

**Examples**

```
###Break date extraction of the Nile data
nile <- as.zoo(Nile)
isat.nile <- isat(nile, sis=TRUE, iis=FALSE, plot=TRUE, t.pval=0.005)
isatdates(isat.nile)
```

isatloop

*Repeated Impulse Indicator Saturation***Description**

Runs `isat` repeatedly at pre-specified significance levels to yield multiple iterations used in `outlierscaletest`.

**Usage**

```
isatloop(num=c(seq(from=20, to=1, by=-1)), t.pval.spec = FALSE,
print=FALSE, y, ar=NULL, iis=TRUE, sis=FALSE, ...)
```

**Arguments**

<code>num</code>	numeric, target expected number of outliers under the null hypothesis, or target proportion of outliers if <code>t.pval.spec==TRUE</code>
<code>t.pval.spec</code>	logical, if TRUE, then <code>num</code> specifies proportion rather than number of targeted outliers
<code>print</code>	logical, if TRUE, then iterations are printed
<code>y</code>	numeric vector, time-series or <code>zoo</code> object. Missing values in the beginning and at the end of the series is allowed, as they are removed with the <code>na.trim</code> command
<code>ar</code>	integer vector, say, <code>c(2,4)</code> or <code>1:4</code> . The AR-lags to include in the mean specification
<code>iis</code>	logical, whether to use <code>iis</code>
<code>sis</code>	logical, whether to use <code>sis</code> , default is FALSE
<code>...</code>	any argument from <code>isat</code> can also be used in <code>isatloop</code>

**Details**

The function repeatedly runs `isat` detecting outliers in a model of `y` at different chosen target levels of significance specified in `num`. The output of this function is used as the input for the `outlierscaletest` function. All additional arguments from `isat` can be passed to `isatloop`.

**Value**

Returns a list of two items. The first item is the number of observations. The second item is a dataframe containing the expected and observed proportion (and number of outliers) for each specified significance level of selection.

**Author(s)**

Felix Pretis, <http://www.felixpretis.org/>

## References

- Jiao, X. & Pretis, F. (2019). Testing the Presence of Outliers in Regression Models. Discussion Paper.
- Pretis, F., Reade, J., & Sucarrat, G. (2018). Automated General-to-Specific (GETS) regression modeling and indicator saturation methods for the detection of outliers and structural breaks. *Journal of Statistical Software*, 86(3).

## See Also

[isat](#), [outlierscaletest](#)

## Examples

```
###Repeated isat models using the Nile dataset
### where p-values are chosen such that the expected number of outliers under the null
### corresponds to 1, 2, 3, 4 and 5.
nile <- as.zoo(Nile)
isat.nile.loop <- isatloop(y=nile, iis=TRUE, num=c(1,2, 3, 4, 5))
```

---

isattest

*Indicator Saturation Test*

---

## Description

Takes an 'isat' object returned by the `isat` function as input and returns the results of a hypothesis test on the time-varying intercept or long-run equilibrium against a specified null-hypothesis for a chosen level of significance - see Pretis (2015).

## Usage

```
isattest(x, hnull=0, lr=FALSE, ci.pval=0.99, plot=NULL, plot.turn=FALSE,
  conscorr=FALSE, effcorr=FALSE, mcor = 1, biascorr=FALSE, mxfull = NULL,
  mxbreak=NULL)
```

## Arguments

- |                      |  |
|----------------------|--|
| <code>x</code>       | a 'gets' object obtained with the <a href="#">isat</a> function  |
| <code>hnull</code>   | numeric. the null-hypothesis value to be tested against.   |
| <code>lr</code>      | logical. If TRUE and 'x' contains autoregressive elements, then <code>isattest</code> tests on the long-run equilibrium path. See Pretis (2015). |
| <code>ci.pval</code> | numeric between 0 and 1. Default is 0.99, the level of significance for the confidence interval of the test against 'hnull'.                     |
| <code>plot</code>    | logical. If TRUE, then a plot showing the coefficient path and bias relative to 'hnull' is shown.  |

<code>plot.turn</code>	logical. If TRUE, then the plot output adds the time of the breaks to the plot showing the bias relative to 'hnull'.
<code>biascorr</code>	logical. If TRUE, then the coefficient path is bias-corrected using <code>biascorr</code> . This is only valid for the non-dynamic test without additional covariates.
<code>conscorr</code>	logical. If TRUE then the Johansen and Nielsen (2016) impulse-indicator consistency correction is applied to estimated residual variance.
<code>effcorr</code>	logical. If TRUE then the Johansen and Nielsen (2016) m-step efficiency correction is applied to estimated standard errors of 'fixed' regressors.
<code>mcor</code>	integer. The m-step efficiency correction factor, where $m=mcor$ .
<code>mxfull</code>	string. The name of the full-sample variable when constructing the coefficient path of user-specified break variables.
<code>mxbreak</code>	string. The name of the break variables used to construct the coefficient path of user-specified break variables.

### Details

The function tests the coefficient path (or long-run equilibrium path) against a specified null hypothesis at a chosen level of significance. If conducted on an `isat` model of a forecast error or relative forecast differential, then this corresponds to the test of time-varying predictive accuracy of Pretis (2015). The resulting output plot shows the coefficient path in the top panel (where 'hnull' is plotted as dotted lines), with the bias (significant difference relative to 'hnull') in the lower panel. If `mxfull` and `mxbreak` are specified, then the function tests on the coefficient path of the user-specified variable, where `mxfull` denotes the full-sample variable name, to which the `mxbreak` variables are added. To correct for the under-estimation of the residual variance, the argument `conscorr` implements the Johansen and Nielsen (2016) consistency correction, and `effcorr` adds the efficiency correction for standard errors on fixed regressors which are not selected over.

### Value

A  $T \times 4$  matrix (with  $T$  = number of observations) where the first two columns denote the confidence interval of the coefficient path (or the long-run equilibrium path if 'lr=TRUE'). The third and fourth column denote the bias of the coefficient path relative to the chosen null-hypothesis, where 'bias.high' denotes the bias when the series tested is above the hypothesized value, and 'bias.low' denotes the bias when the series tested is significantly below the hypothesized value.

### Author(s)

Felix Pretis, <http://www.felixpretis.org/>

### References

- Johansen, S., & Nielsen, B. (2016): 'Asymptotic theory of outlier detection algorithms for linear time series regression models.' *Scandinavian Journal of Statistics*, 43(2), 321-348.
- Pretis, F. (2015): 'Testing for time-varying predictive accuracy using bias-corrected indicator saturation'. Oxford Department of Economics Discussion Paper.
- Hendry, David, F., Johansen, Soren, and Santos, Carlos (2007): 'Automatic selection of indicators in a fully saturated regression'. *Computational Statistics*, vol 23:1, pp.317-335.

**See Also**

[isat](#), [coef.gets](#), [plot.gets](#), [biascorr](#), [isatvar](#)

**Examples**

```
##Using artificial data:
#set.seed(123)
#d <- matrix(0,100,1)
#d[35:55] <- 1
#e <- rnorm(100, 0, 1)
#y <- d*2 +e
#plot(y, type="l")

##Static Test against hnull=0 using bias-correction:

#ys <- isat(y, sis=TRUE, iis=FALSE, tis=FALSE, t.pval=0.01)
#isattest(ys, hnull=0, lr=FALSE, ci.pval = 0.99, plot.turn = FALSE, biascorr=TRUE)

##Dynamic Test of the long-run equilibrium against hnull=2 with breakpoints
##labelled in the plot:

#ys <- isat(y, sis=TRUE, iis=FALSE, tis=FALSE, t.pval=0.01, ar=1:2)
#isattest(ys, hnull=2, lr=TRUE, ci.pval = 0.99, plot.turn = TRUE, biascorr=FALSE)
```

---

isatvar

*Variance of the coefficient path*


---

**Description**

Takes an 'isat' object returned by the `isat` function as input and returns the coefficient path of the constant (and long-run equilibrium if 'lr' is specified) together with its approximate variance and standard errors. If `mxfull` and `mxbreak` are specified, then the function returns the coefficient path of the user-specified variable.

**Usage**

```
isatvar(x, lr=FALSE, conscorr=FALSE, effcorr=FALSE, mcor = 1,
        mxfull = NULL, mxbreak=NULL)
```

**Arguments**

<code>x</code>	a 'gets' object obtained with the <code>isat</code> function
<code>lr</code>	logical. If TRUE and 'x' contains autoregressive elements, then <code>isatvar</code> also returns the long-run equilibrium coefficient path with its variance and standard deviation. See Pretis (2015).
<code>conscorr</code>	logical. If TRUE then the Johansen and Nielsen (2016) impulse-indicator consistency correction is applied to estimated residual variance.

effcorr	logical. If TRUE then the Johansen and Nielsen (2016) m-step efficiency correction is applied to estimated standard errors of ‘fixed’ regressors.
mcor	integer. The m-step efficiency correction factor, where m=mcor.
mxfull	string. The name of the full-sample variable when constructing the coefficient path of user-specified break variables.
mxbreak	string. The name of the break variables used to construct the coefficient path of user-specified break variables.

## Details

The function computes the approximate variance and standard errors of the intercept term with structural breaks determined by `isat`. This permits hypothesis testing and plotting of approximate confidence intervals for the intercept in the presence of structural breaks. For dynamic autoregressive models in `isat` the `lr` argument returns the time-varying long-run equilibrium together with its approximate variance and standard errors. If `mxfull` and `mxbreak` are specified, then the function returns the coefficient path of the user-specified variable, where `mxfull` denotes the full-sample variable name, to which the `mxbreak` variables are added. To correct for the under-estimation of the residual variance, the argument `conscorr` implements the Johansen and Nielsen (2016) consistency correction, and `effcorr` adds the efficiency correction for standard errors on fixed regressors which are not selected over.

## Value

If `lr=FALSE`: A  $T \times 4$  matrix (with  $T$  = number of observations) where the first column denotes the coefficient path relative to the full sample coefficient, the second column the coefficient path of the intercept, the third the approximate variance of the coefficient path, and the fourth column the approximate standard errors of the coefficient path. If `lr=TRUE`: A  $T \times 7$  matrix where the first four columns are identical to the `lr=FALSE` case, and the additional columns denote the long-run equilibrium coefficient path, together with the approximate variance and standard errors of the long-run equilibrium coefficient path.

## Author(s)

Felix Pretis, <http://www.felixpretis.org/>  
James Reade, <https://sites.google.com/site/jjamesreade/>

## References

- Pretis, F. (2015): ‘Testing for time-varying predictive accuracy using bias-corrected indicator saturation’. Oxford Department of Economics Working Paper.
- Johansen, S., & Nielsen, B. (2016): ‘Asymptotic theory of outlier detection algorithms for linear time series regression models.’ *Scandinavian Journal of Statistics*, 43(2), 321-348.
- Pretis, Felix, Reade, James and Sucarrat, Genaro (2018): ‘Automated General-to-Specific (GETS) Regression Modeling and Indicator Saturation for Outliers and Structural Breaks’. *Journal of Statistical Software* 86, Number 3, pp. 1-44

## See Also

[isat](#), [coef.gets](#), [plot.gets](#), [biascorr](#), [isattest](#)



## Examples

```
##Variance in presence of a break
#nile <- as.zoo(Nile)
#isat.nile <- isat(nile, sis=TRUE, iis=FALSE, plot=FALSE, t.pval=0.005)
#var <- isatvar(isat.nile)

#plot(nile)
#lines(isat.nile$mean.fit, col="red")
#lines(isat.nile$mean.fit + 2*var$const.se, col="blue", lty=3)
#lines(isat.nile$mean.fit - 2*var$const.se, col="blue", lty=3)

##Variance when there is no break
#set.seed(1)
#x <- as.zoo(rnorm(100, 0, 1))
#isat.x <- isat(x, sis=TRUE, iis=FALSE, plot=TRUE, t.pval=0.005)
#var.x <- isatvar(isat.x)

#plot(x)
#lines(isat.x$mean.fit, col="red")
#lines(isat.x$mean.fit + 2*var.x[,2], col="blue", lty=3)
#lines(isat.x$mean.fit - 2*var.x[,2], col="blue", lty=3)

##Variance of the long-run equilibrium coefficient path

#nile <- as.zoo(Nile)
#isat.nile <- isat(nile, sis=TRUE, iis=FALSE, plot=TRUE, t.pval=0.005, ar=1:2)
#var <- isatvar(isat.nile, lr=TRUE)
```

---

isatvarcorrect	<i>Consistency and Efficiency Correction for Impulse Indicator Saturation</i>
----------------	---

---

## Description

Takes an `isat` object and corrects the estimates of the error variance and the estimated standard errors of 'forced' regressors.

## Usage

```
isatvarcorrect(x, mcor=1)
```

## Arguments

<code>x</code>	an <code>isat</code> object
<code>mcor</code>	integer, number of iterations in the correction. Default = 1.

## Details

Impulse indicator saturation results in an under-estimation of the error variance as well as the variance of regressors not selected over. The magnitude of the inconsistency increases with the p-value of selection (`t.pval`). The function takes an `isat` object and applies the impulse indicator consistency (`isvarcor`) and efficiency correction (`isvareffcor`) of the estimated error variance and the estimated variance of regressors not selected over. See Johansen and Nielsen (2016a) and (2016b).

## Value

Returns an `isat` object in which the estimated standard errors, t-statistics, p-values, standard error of the regression, and log-likelihood are consistency and efficiency corrected when using impulse indicator saturation (`iis=TRUE`).

## Author(s)

Felix Pretis, <http://www.felixpretis.org/>

## References

Johansen, S., & Nielsen, B. (2016a). Asymptotic theory of outlier detection algorithms for linear time series regression models. *Scandinavian Journal of Statistics*, 43(2), 321-348.

Johansen, S., & Nielsen, B. (2016b). Rejoinder: Asymptotic Theory of Outlier Detection Algorithms for Linear. *Scandinavian Journal of Statistics*, 43(2), 374-381.

Pretis, F., Reade, J., & Sucarrat, G. (2018). Automated General-to-Specific (GETS) regression modeling and indicator saturation methods for the detection of outliers and structural breaks. *Journal of Statistical Software*, 86(3).

## See Also

`isat`, `isvarcor`, `isvareffcor`

## Examples

```
###Consistency and Efficiency Correction of Impulse Indicator Estimates
nile <- as.zoo(Nile)
isat.nile <- isat(nile, sis=FALSE, iis=TRUE, plot=TRUE, t.pval=0.1)
isat.nile.corrected <- isatvarcorrect(isat.nile)

isat.nile$sigma2
isat.nile.corrected$sigma2
```

---

isvarcor                      *IIS Consistency Correction*

---

**Description**

Consistency correction for estimate of residual variance when using impulse indicator saturation.

**Usage**

```
isvarcor(t.pval, sigma)
```

**Arguments**

t.pval	numeric value. the p-value of selection in the impulse indicator saturation model.
sigma	numeric value. The estimated standard deviation of the residuals from the impulse indicator saturation model.

**Details**

The Johansen and Nielsen (2016) impulse-indicator consistency correction for the estimated residual standard deviation.

**Value**

a data frame containing the corrected standard deviation `$sigma.cor` and the correction factor used `$corxi`

**Author(s)**

Felix Pretis, <http://www.felixpretis.org/>

**References**

Johansen, S., & Nielsen, B. (2016): 'Asymptotic theory of outlier detection algorithms for linear time series regression models.' *Scandinavian Journal of Statistics*, 43(2), 321-348.

Pretis, Felix, Reade, James and Sucarrat, Genaro (2018): 'Automated General-to-Specific (GETS) Regression Modeling and Indicator Saturation for Outliers and Structural Breaks'. *Journal of Statistical Software* 86, Number 3, pp. 1-44

**See Also**

[isatvar](#)

**Examples**

```
isvarcor(t.pval=0.05, sigma=2)
```

---

`isvareffcor`*IIS Efficiency Correction*

---

**Description**

Efficiency correction for the estimates of coefficient standard errors on fixed regressors.

**Usage**

```
isvareffcor(t.pval, se, m=1)
```

**Arguments**

<code>t.pval</code>	numeric value. the p-value of selection in the impulse indicator saturation model.
<code>se</code>	numeric value or vector. The estimated standard errors of the coefficients on fixed regressors in impulse indicator saturation model.
<code>m</code>	integer. The m-step correction factor.

**Details**

The Johansen and Nielsen (2016) impulse-indicator efficiency correction for the estimated standard errors on fixed regressors in impulse indicator models.

**Value**

a data frame containing the corrected standard deviation `$se.cor` and the correction factor used `$eta.m`

**Author(s)**

Felix Pretis, <http://www.felixpretis.org/>

**References**

Johansen, S., & Nielsen, B. (2016): 'Asymptotic theory of outlier detection algorithms for linear time series regression models.' *Scandinavian Journal of Statistics*, 43(2), 321-348.

Pretis, Felix, Reade, James and Sucarrat, Genaro (2018): 'Automated General-to-Specific (GETS) Regression Modeling and Indicator Saturation for Outliers and Structural Breaks'. *Journal of Statistical Software* 86, Number 3, pp. 1-44

**See Also**

[isatvar](#)

**Examples**

```
isvareffcor(t.pval=0.05, se=2, m=1)
```

---

`mvrnormsim`*Simulate from a Multivariate Normal Distribution*

---

**Description**

Produces one or more samples from the specified multivariate normal distribution. Used in [outlierscaletest](#).

**Usage**

```
mvrnormsim(n = 1, mu, Sigma, tol = 1e-6, empirical = FALSE)
```

**Arguments**

<code>n</code>	the number of samples required.
<code>mu</code>	a vector giving the means of the variables.
<code>Sigma</code>	a positive-definite symmetric matrix specifying the covariance matrix of the variables.
<code>tol</code>	tolerance (relative to largest variance) for numerical lack of positive-definiteness in <code>Sigma</code> .
<code>empirical</code>	logical. If true, <code>mu</code> and <code>Sigma</code> specify the empirical not population mean and covariance matrix.

**Details**

Original function `mvrnorm` developed by Venables, W. N. & Ripley. in package MASS, <https://CRAN.R-project.org/package=MASS>.

**Value**

If `n = 1` a vector of the same length as `mu`, otherwise an `n` by `length(mu)` matrix with one sample in each row.

**Author(s)**

Venables, W. N. & Ripley, with modifications by Felix Pretis, <http://www.felixpretis.org/>

**References**

Venables, W. N. & Ripley, B. D. (2019): 'MASS: Support Functions and Datasets for Venables and Ripley's MASS'. <https://CRAN.R-project.org/package=MASS>

Venables, W. N. & Ripley, B. D. (2002) Modern Applied Statistics with S. Fourth Edition. Springer, New York. ISBN 0-387-95457-0

**See Also**

[outlierscaletest](#)

**Examples**

```
Sigma <- matrix(c(3,2,1,7),2,2)
mvrnormsim(n=2, mu=c(1,2), Sigma)
```

---

ols *OLS estimation*

---

**Description**

OLS estimation with the QR decomposition and, for some options, computation of variance-covariance matrices

**Usage**

```
ols(y, x, untransformed.residuals=NULL, tol=1e-07, LAPACK=FALSE, method=3, ...)
```

**Arguments**

y	numeric vector, the regressand
x	numeric matrix, the regressors
untransformed.residuals	NULL (default) or, when <code>ols</code> is used with <code>method=6</code> , a numeric vector containing the untransformed residuals
tol	numeric value. The tolerance for detecting linear dependencies in the columns of the regressors, see the <code>qr</code> function. Only used if LAPACK is FALSE
LAPACK	logical, TRUE or FALSE (default). If true use LAPACK otherwise use LINPACK, see the <code>qr</code> function
method	an integer, 1 to 6, that determines the estimation method
...	further arguments (currently ignored)

**Details**

`method = 1` or `2` only returns the OLS coefficient estimates together with the QR-information. `method = 1` is slightly faster than `method=2`. `method=3` returns, in addition, the ordinary variance-covariance matrix of the OLS estimator. `method=4` returns the White (1980) heteroscedasticity robust variance-covariance matrix in addition to the information returned by `method=3`, whereas `method=5` does the same except that the variance-covariance matrix now is that of Newey and West (1987). `method=6` undertakes OLS estimation of a log-variance model, see Pretis, Reade and Sucarrat (2018, Section 4).

**Value**

A list with items depending on `method`

**Author(s)**

Genaro Sucarrat, <http://www.sucarrat.net/>

## References

W. Newey and K. West (1987): 'A Simple Positive Semi-Definite, Heteroskedasticity and Autocorrelation Consistent Covariance Matrix', *Econometrica* 55, pp. 703-708.

F. Pretis, J. Reade and G. Sucarrat (2018): 'Automated General-to-Specific (GETS) Regression Modeling and Indicator Saturation for Outliers and Structural Breaks', *Journal of Statistical Software* 86, Issue 3, pp. 1-44, DOI: <https://doi.org/10.18637/jss.v086.i03>

H. White (1980): 'A Heteroskedasticity-Consistent Covariance Matrix and a Direct Test for Heteroskedasticity', *Econometrica* 48, pp. 817-838.

## See Also

[qr](#), [solve.qr](#)

---

outlierscaletest

*Sum and Sup Scaling Outlier Tests*

---

## Description

Computes the Sum and Supremum Scaling Tests for the overall presence of outliers based on Jiao and Pretis (2019).

## Usage

```
outlierscaletest(x, nsim = 10000)
```

## Arguments

x	list, output of the <a href="#">isatloop</a> function
nsim	integer, number of replications to simulate critical values for the Sup test

## Details

The function takes the output of the [isatloop](#) function and computes the Scaling Sum and Supremum Tests for the presence of outliers from Jiao and Pretis (2019). The test compares the expected and observed proportion of outliers over the range of different significance levels of selection specified in [isatloop](#). The Sum test compares the sum of deviations against the standard normal distribution, the Sup test compares the supremum of deviations against critical values simulated with `nsim` replications. The null hypothesis is that the observed proportion of outliers scales with the proportion of outliers under the null of no outliers.

## Value

Returns a list of two `htest` objects. The first providing the results of the Sum test on the sum of the deviation of outliers against a standard normal distribution. The second providing the results on the supremum of the deviation of outliers against simulated critical values.

**Author(s)**

Xiyu Jiao, & Felix Pretis, <http://www.felixpretis.org/>

**References**

Jiao, X. & Pretis, F. (2019). Testing the Presence of Outliers in Regression Models. Discussion Paper.

Pretis, F., Reade, J., & Sucarrat, G. (2018). Automated General-to-Specific (GETS) regression modeling and indicator saturation methods for the detection of outliers and structural breaks. *Journal of Statistical Software*, 86(3).

**See Also**

[isat](#), [isatloop](#)

**Examples**

```
###Repeated isat models using the Nile dataset
### where p-values are chosen such that the expected number of outliers under the null
### corresponds to 1, 2, ..., 20. Then computing the Outlier Scaling Tests:

#nile <- as.zoo(Nile)
#isat.nile.loop <- isatloop(y=nile)
#outlierscaletest(isat.nile.loop)
```

---

outliertest

*Jiao and Pretis Outlier Proportion and Count Tests*

---

**Description**

Tests whether the proportion (or number) of outliers detected using impulse indicator saturation is different from the proportion (or number) of outliers expected under the null hypothesis of no outliers using the Jiao and Pretis (2019) proportion and count outlier tests.

**Usage**

```
outliertest(x, noutl=NULL, t.pval=NULL, T=NULL,
m=1, infly=FALSE, alternative="two.sided")
```

**Arguments**

x	an <a href="#">isat</a> object
noutl	integer, number of detected outliers if no <a href="#">isat</a> object is provided i.e. x=NULL
t.pval	numeric, between 0 and 1. Selection p-value used in indicator saturation if no <a href="#">isat</a> object is provided i.e. x=NULL



T	integer, sample sized used in indicator saturation if no <code>isat</code> object is provided i.e. <code>x=NULL</code>
m	integer, number of iterations in variance computation, default=1
infty	logical, argument used for variance computation
alternative	"two-sided", "less", "greater", alternative hypothesis of outlier test.

## Details

The function computes the estimated proportion of outliers (gauge) based on impulse indicator saturation and constructs the proportion and count outlier test statistics from Jiao and Pretis (2019). The null hypothesis is that the proportion (or count) of outliers is not different than the proportion (or count) of outliers detected under the null hypothesis of no outliers. The first test compares the estimated proportion of outliers scaled by its estimated variance against a standard normal distribution. The second test compares the number of outliers against a Poisson distribution. If an `isat` object is provided in `x`, then the function automatically extracts the detected impulses and computes the estimated outlier proportion. If no `isat` object is provided and `x=NULL`, then the tests can be conducted manually by providing the number of detected outliers (`nout1`), the sample size (`T`), and the chosen level of significance used to detect outliers (`t.pval`).

## Value

Returns a list of two `htest` objects. The first providing the results of the test on the proportion of outliers against a standard normal distribution. The second providing the results on the number of outliers against the Poisson distribution.

## Author(s)

Xiyu Jiao, & Felix Pretis, <http://www.felixpretis.org/>

## References

- Jiao, X. & Pretis, F. (2019). Testing the Presence of Outliers in Regression Models. Discussion Paper.
- Pretis, F., Reade, J., & Sucarrat, G. (2018). Automated General-to-Specific (GETS) regression modeling and indicator saturation methods for the detection of outliers and structural breaks. *Journal of Statistical Software*, 86(3).

## See Also

[isat](#)

## Examples

```
###Testing the Presence of Outliers in the Nile Data
nile <- as.zoo(Nile)
isat.nile <- isat(nile, sis=FALSE, iis=TRUE, plot=TRUE, t.pval=0.1)
outliertest(isat.nile)

###Testing the number of outliers when the sample is T=200,
```

```
### with 7 detected outliers at t.pval=0.05 if no isat object is provided:  
outliertest(x=NULL, noutl=7, t.pval=0.05, T=200)
```

---

paths

*Extraction functions for 'arx', 'gets' and 'isat' objects*

---

### Description

Extraction functions for objects of class 'arx', 'gets' and 'isat'

### Usage

```
paths(object, ...)  
terminals(object, ...)  
rsquared(object, adjusted=FALSE, ...)
```

### Arguments

object	an object of class 'arx', 'gets' or 'isat'
adjusted	logical. If TRUE the adjusted R-squared is returned
...	additional arguments

### Details

paths and terminals can only be applied on objects of class 'gets' and 'isat'

### Value

paths:	a <b>list</b> with the paths searched (each number refers to a regressor in the GUM)
terminals:	a <b>list</b> with the terminal models (each number refers to a regressor in the GUM)
rsquared:	a <b>numeric</b> , the R-squared of the regression, or adjusted R-squared if adjusted is set to TRUE

### Author(s)

Genaro Sucarrat, <http://www.sucarrat.net/>

### See Also

[getsm](#), [getsm](#), [getsv](#), [isat](#)

**Examples**

```
##Simulate from an AR(1):
set.seed(123)
y <- arima.sim(list(ar=0.4), 50)

##Simulate four independent Gaussian regressors:
xregs <- matrix(rnorm(4*50), 50, 4)

##estimate an AR(2) with intercept and four conditioning
##regressors in the mean:
mymod <- arx(y, mc=TRUE, ar=1:2, mxreg=xregs)
rsquared(mymod)
rsquared(mymod, adjusted=TRUE)

##General-to-Specific (GETS) modelling of the mean:
meanmod <- getsm(mymod)
rsquared(meanmod)
rsquared(meanmod, adjusted=TRUE)

##extract the paths searched:
paths(meanmod)

##extract the terminal models:
terminals(meanmod)
```

---

periodicdummies

*Make matrix of periodicity (e.g. seasonal) dummies*


---

**Description**

Auxiliary function that generates periodicity dummies (e.g. seasonal dummies) for regular time series. The function is similar to, but more general than, the `seasonaldummy` function in the `forecast` package.

**Usage**

```
periodicdummies(x, values=1)
```

**Arguments**

x	a regular time series (vector or matrix)
values	numeric of length 1 (default) or numeric vector of length equal to frequency(x)

**Value**

A matrix of class `zoo` with the periodicity dummies

**Author(s)**

Genaro Sucarrat, <http://www.sucarrat.net/>

**See Also**

[is.regular](#), [zooreg](#), [zoo](#), [ts](#)

**Examples**

```
##quarterly dummies:
x <- zooreg(rnorm(30), start=2000, frequency=4)
periodicdummies(x)

##monthly dummies:
y <- zooreg(rnorm(30), start=c(2000,1), frequency=12)
periodicdummies(y)
```

---

predict.arx

*Forecasting with 'arx' models*

---

**Description**

Generate out-of-sample forecasts up to `n` steps ahead for objects of class `arx`. Optionally, quantiles of the forecasts are also returned if any of the arguments `ci.levels` or `probs` are specified. The forecasts, confidence intervals and quantiles are obtained via simulation. By default, 5000 simulations is used, but this can be changed via the `n.sim` argument. Also by default, the simulations uses a classical bootstrap to sample from the standardised residuals. To use an alternative set of standardised innovations, for example the standard normal, use the `innov` argument. If `plot=TRUE`, then a plot of the forecasts is created.

**Usage**

```
## S3 method for class 'arx'
predict(object, spec=NULL, n.ahead=12, newmxreg=NULL,
        newvxreg=NULL, newindex=NULL, n.sim=5000, innov=NULL, probs=NULL,
        ci.levels=NULL, quantile.type=7, return=TRUE, verbose=FALSE,
        plot=NULL, plot.options=list(), ...)
```

**Arguments**

<code>object</code>	an object of class 'arx'
<code>spec</code>	NULL (default), "mean", "variance" or "both". If NULL, then it is automatically determined whether information pertaining to the mean or variance specification should be returned
<code>n.ahead</code>	integer that determines how many steps ahead predictions should be generated (the default is 12)

newmxreg	a matrix of n.ahead rows and NCOL(mxreg) columns with the out-of-sample values of the mxreg regressors
newvxreg	a matrix of n.ahead rows and NCOL(vxreg) columns with the out-of-sample values of the vxreg regressors
newindex	NULL (default) or the date-index for the zoo object returned by predict.arx. If NULL, then the function uses the in-sample index to generate the out-of-sample index
n.sim	integer, the number of replications used for the generation of the forecasts
innov	NULL (default) or a vector of length n.ahead * n.sim containing the standardised errors (that is, zero mean and unit variance) used for the forecast simulations. If NULL, then a classical bootstrap procedure is used to draw from the standardised in-sample residuals
probs	NULL (default) or a vector with the quantile-levels (values strictly between 0 and 1) of the forecast distribution. If NULL, then no quantiles are returned unless ci.levels is non-NULL
ci.levels	NULL (default) or a vector with the confidence levels (expressed as values strictly between 0 and 1) of the forecast distribution. The upper and lower values of the confidence interval(s) are returned as quantiles
quantile.type	an integer between 1 and 9 that selects which algorithm to be used in computing the quantiles, see the argument type in <a href="#">quantile</a>
return	logical. If TRUE (default), then the out-of-sample predictions are returned. The value FALSE, which does not return the predictions, may be of interest if only a prediction plot is of interest
verbose	logical with default FALSE. If TRUE, then additional information (typically the quantiles and/or the simulated series) used in the generation of forecasts is returned. If FALSE, then only the forecasts are returned
plot	NULL (default) or logical. If NULL, then the value set by options\$plot (see <a href="#">options</a> ) determines whether a plot is produced or not. If TRUE, then the out-of-sample forecasts are plotted.
plot.options	a list of options related to the plotting of forecasts, see 'Details'
...	additional arguments

### Details

The plot.options argument is a list that, optionally, can contain any of the following arguments:

- keep: integer greater than zero (the default is 12) that controls the number of in-sample actual values to plot
- line.at.origin: logical. If TRUE, then a vertical line is drawn at the forecast origin, that is, at the last in-sample observation
- start.at.origin: logical. If TRUE, then the drawing of the forecast line starts at the actual value of the forecast origin

- `dot.at.origin`: logical. If TRUE, then a dot is drawn at the forecast origin
- `hlines`: numeric vector that indicates where to draw grey horizontal grid lines
- `col`: numeric vector of length two that controls the colour of the plotted lines. The first value controls the colour of the forecasts and the fitted values, whereas the second controls the colour of the actual values
- `lty`: numeric vector of length two that controls the line type. The first value controls the line type of the forecast, whereas the second controls the line type of the actual and fitted values
- `lwd`: an integer that controls the width of the plotted lines (the default is 1)
- `ylim`: numeric vector of length two that contains the limits of the y-axis of the prediction plot
- `ylab`: a character that controls the text on the y-axis
- `main`: a character that controls the text in the overall title
- `legend.text`: a character vector of length two that controls how the forecast and actual lines should be named or referred to in the legend of the plot
- `fitted`: If TRUE, then the fitted values as well as actual values are plotted in-sample
- `newmactual`: numeric vector or NULL (default). Enables the plotting of actual values out-of-sample in the mean in addition to the forecasts
- `newvactual`: numeric vector or NULL (default). Enables the plotting of squared residuals ('actual values') out-of-sample in addition to the forecasts
- `shades.of.grey`: numeric vector of length `length(ci.levels)` that contains the shades of grey associated with the confidence intervals in the prediction plot. The shades can range from 100 (white) to 0 (black)

### Value

a vector of class `zoo` containing the out-of-sample forecasts, or a matrix of class `zoo` containing the out-of-sample forecasts together with prediction-quantiles, or NULL if `return=FALSE`

### Author(s)

Felix Pretis, <http://www.felixpretis.org/>  
James Reade, <https://sites.google.com/site/jjamesreade/>  
Genaro Sucarrat, <http://www.sucarrat.net/>

**See Also**[arx](#)**Examples**

```

##simulate from an AR(1):
set.seed(123)
y <- arima.sim(list(ar=0.4), 40)

##estimate AR(2) model with intercept:
mymod <- arx(y, mc=TRUE, ar=c(1,2))

##generate out-of-sample forecasts:
predict(mymod)

##same, but plot the predictions in addition:
#predict(mymod, plot=TRUE)

##same, but return also the quantiles of the confidence intervals:
#predict(mymod, ci.levels=c(0.50,0.90), plot=TRUE)

##same, but with non-default levels on the confidence intervals:
#predict(mymod, ci.levels=c(0.20,0.80, 0.99), plot=TRUE)

##same, but with more confidence intervals:
#predict(mymod, ci.levels=seq(0.20, 0.95, by=0.05), plot=TRUE)

##same, but with less rugged ci's (achieved by increasing n.sim):
##predict(mymod, ci.levels=seq(0.20, 0.95, by=0.05), n.sim=50000, plot=TRUE)

##same, but using standard normals (instead of bootstrap) in the simulations:
#n.sim <- 2000
#n.ahead <- 12 #the default on n.ahead
#predict(mymod, ci.levels=seq(0.20, 0.95, by=0.05), n.sim=n.sim,
# innov=rnorm(n.ahead*n.sim), plot=TRUE)

##make x-regressors:
x <- matrix(rnorm(40*3), 40, 3)

##estimate AR(1) model with intercept and covariates:
mymod <- arx(y, mc=TRUE, ar=1, mxreg=x)

##predict up to 5 steps ahead, setting x's to 0 out-of-sample:
predict(mymod, n.ahead=5, newmxreg=matrix(0,5,NCOL(x)))

##same, but do also plot:
#predict(mymod, n.ahead=5, newmxreg=matrix(0,5,NCOL(x)),
# plot=TRUE)

##estimate an AR(2) model w/intercept and a log-ARCH(1) specification
##on the variance:
#mymodel <- arx(y, mc=TRUE, ar=1:2, arch=1)

```

```

##generate forecasts of the conditional variances:
#predict(mymodel, spec="variance")

##same, but do also plot:
#predict(mymodel, spec="variance", plot=TRUE)

##illustrations of the usage of plot.options:
#mymodel <- arx(y, mc=TRUE, ar=1)
#predict(mymodel, plot=TRUE, plot.options=list(keep=1))
#predict(mymodel, plot=TRUE, plot.options=list(line.at.origin=TRUE))
#predict(mymodel, plot=TRUE, plot.options=list(start.at.origin=FALSE))
#predict(mymodel, plot=TRUE,
# plot.options=list(start.at.origin=FALSE, fitted=TRUE))
#predict(mymodel, plot=TRUE, plot.options=list(dot.at.origin=FALSE))
#predict(mymodel, plot=TRUE, plot.options=list(hlines=c(-2,-1,0,1,2)))
#predict(mymodel, plot=TRUE, plot.options=list(col=c("darkred","green")))
#predict(mymodel, plot=TRUE, plot.options=list(lty=c(3,2)))
#predict(mymodel, plot=TRUE, plot.options=list(lwd=3))
#predict(mymodel, plot=TRUE, plot.options=list(ylim=c(-8,8)))
#predict(mymodel, plot=TRUE, plot.options=list(ylab="User-specified y-axis"))
#predict(mymodel, plot=TRUE,
# plot.options=list(main="User-specified overall title"))
#predict(mymodel, plot=TRUE,
# plot.options=list(legend.text=c("User-specified 1","User-specified 2")))
#predict(mymodel, plot=TRUE, plot.options=list(fitted=TRUE))
#predict(mymodel, plot=TRUE, plot.options=list(newmactual=rep(0,6)))
#predict(mymodel, plot=TRUE, plot.options=list(shades.of.grey=c(95,50)))
#predict(mymodel, plot=TRUE, plot.options=list(shades.of.grey=c(50,95))) #invert shades

```

---

printtex

---

*Generate LaTeX code of an estimation result*


---

## Description

Convenience functions that generates LaTeX-code of an estimation result in equation-form. `printtex` can, in principle, be applied to any object for which `coef`, `vcov` and `logLik` methods exist. Note: The generated LaTeX-code contains an `eqnarray` environment, which requires that the `amsmath` package is loaded in the preamble of the LaTeX document.

## Usage

```

printtex(x, fitted.name=NULL, xreg.names=NULL, digits=4,
         intercept=TRUE, gof=TRUE, diagnostics=TRUE, nonumber=FALSE,
         nobs="T")
## S3 method for class 'arx'
toLatex(object, ...)
## S3 method for class 'gets'
toLatex(object, ...)

```



**Arguments**

<code>x</code>	an estimation result, e.g. <code>arx</code> , <code>gets</code> or <code>isat</code> object
<code>object</code>	an estimation result of class <code>arx</code> or <code>gets</code>
<code>fitted.name</code>	NULL or a user-specified name of left-hand side variable
<code>xreg.names</code>	NULL or a user-specified character vector with the names of regressors
<code>digits</code>	integer, the number of digits to be printed
<code>intercept</code>	logical or numeric. The argument determines whether one of the regressors is an intercept or not, or its location. If TRUE, then the intercept is assumed to be located at <code>coef(x)[1]</code> , and hence the regressor-name of location 1 is excluded from the print. If FALSE, then it is assumed that there is no intercept among the regressors. If numeric, then it is assumed that the regressors contain an intercept at the location equal to the numeric value
<code>gof</code>	logical, whether to include goodness-of-fit in the print
<code>diagnostics</code>	logical, whether to include diagnostics in the print
<code>nonumber</code>	logical, whether to remove or not (default) the equation-numbering
<code>nobs</code>	character, the notation to use to denote the number of observations
<code>...</code>	arguments passed on to <code>printtex</code>

**Details**

Currently, `toLatex.arx` and `toLatex.gets` are simply wrappers to `printtex`

**Value**

LaTeX code of an estimation result

**Author(s)**

Genaro Sucarrat, <http://www.sucarrat.net/>

**See Also**

`arx`, `getsm`, `getsv`, `isat`

**Examples**

```
##simulate random variates, estimate model:
y <- rnorm(30)
mX <- matrix(rnorm(30*2), 30, 2)
mymod <- arx(y, mc=TRUE, ar=1:3, mxreg=mX)

##print latex code of estimation result:
printtex(mymod)

##add intercept, at the end, to regressor matrix:
mX <- cbind(mX,1)
colnames(mX) <- c("xreg1", "xreg2", "intercept")
```

```

mymod <- arx(y, mxreg=mX)

##set intercept location to 3:
printtex(mymod, intercept=3)

```

---

recursive	<i>Recursive estimation</i>
-----------	-----------------------------

---

### Description

Recursive estimation of coefficients and standard errors

### Usage

```

recursive(object, spec="mean", std.errors=TRUE, from=40, tol=1e-07,
          LAPACK=FALSE, plot=TRUE, return=TRUE)

```

### Arguments

object	an <a href="#">arx</a> , gets or isat object
spec	'mean' or 'variance'. If 'mean' (default), the the recursive estimates of the mean-equation are estimated
std.errors	logical. If TRUE (default), then the coefficient standard errors are also computed
from	integer. The starting point of the recursion
tol	numeric. The tolerance for linear dependency among regressors
LAPACK	logical, TRUE or FALSE (default). If true use LAPACK otherwise use LINPACK, see <a href="#">qr</a> function
plot	NULL or logical. If TRUE, then the recursive coefficient estimates are plotted. If NULL (default), then the value set by <a href="#">options</a> determines whether a plot is produced or not.
return	logical. If TRUE (default), then the recursive estimates are returned in a list

### Value

If return=TRUE, then a [list](#) is returned with the following components:

estimates	a <a href="#">zoo</a> matrix with the recursive estimates
standard.errors	a <a href="#">zoo</a> matrix with the standard errors

### Author(s)

Genaro Sucarrat, <http://www.sucarrat.net/>

**See Also**

[ols](#), [qr](#), [solve.qr](#)

**Examples**

```
##generate random variates, estimate model:
y <- rnorm(100)
mX <- matrix(rnorm(4*100), 100, 4)
mymodel <- arx(y, mc=TRUE, mxreg=mX)

##compute recursive estimates and plot them:
recursive(mymodel)
```

---

regressorsMean

*Create the regressors of the mean equation*


---

**Description**

The function generates the regressors of the mean equation in an [arx](#) model. The returned value is a matrix with the regressors and, by default, the regressand in column one. By default, observations (rows) with missing values are removed in the beginning and the end with [na.trim](#), and the returned matrix is a [zoo](#) object.

**Usage**

```
regressorsMean(y, mc = FALSE, ar = NULL, ewma = NULL, mxreg = NULL,
  return.regressand = TRUE, return.as.zoo = TRUE, na.trim=TRUE)
```

**Arguments**

<code>y</code>	numeric vector, time-series or <a href="#">zoo</a> object.
<code>mc</code>	logical. TRUE includes an intercept, whereas FALSE (default) does not.
<code>ar</code>	either NULL (default) or an integer vector, say, <code>c(2, 4)</code> or <code>1:4</code> with the AR-terms. The AR-lags to include in the mean specification. If NULL, then no lags are included.
<code>ewma</code>	either NULL (default) or a <a href="#">list</a> with arguments sent to the <a href="#">eqwma</a> function. In the latter case a lagged moving average of <code>y</code> is included as a regressor.
<code>mxreg</code>	either NULL (default) or a numeric vector or matrix, say, a <a href="#">zoo</a> object, of conditioning variables. Note that, if both <code>y</code> and <code>mxreg</code> are zoo objects, then their samples are matched.
<code>return.regressand</code>	logical. TRUE, the default, includes the regressand as column one in the returned matrix.
<code>return.as.zoo</code>	TRUE, the default, returns the matrix as a <a href="#">zoo</a> object.
<code>na.trim</code>	TRUE, the default, removes observations with NA-values in the beginning and the end with <a href="#">na.trim</a> .

**Value**

A matrix, by default of class `zoo`, with the regressand as column one (the default).

**Author(s)**

Genaro Sucarrat, <http://www.sucarrat.net/>

**See Also**

`arx`, `isat`, `zoo` and `na.trim`.

**Examples**

```
##generate some data:
y <- rnorm(10) #regressand
x <- matrix(rnorm(10*5), 10, 5) #regressors

##create regressors (examples):
regressorsMean(y, mxreg=x)
regressorsMean(y, mxreg=x, return.regressand=FALSE)
regressorsMean(y, mc=TRUE, ar=1:3, mxreg=x)
regressorsMean(log(y^2), mc=TRUE, ar=c(2,4))

##let y and x be time-series:
y <- ts(y, frequency=4, end=c(2018,4))
x <- ts(x, frequency=4, end=c(2018,4))
regressorsMean(y, mxreg=x)
regressorsMean(y, mc=TRUE, ar=1:3, mxreg=x)
regressorsMean(log(y^2), mc=TRUE, ar=c(2,4))

##missing values (NA):
y[1] <- NA
x[10,3] <- NA
regressorsMean(y, mxreg=x)
regressorsMean(y, mxreg=x, na.trim=FALSE)
```

---

so2data

*UK SO2 Data*

---

**Description**

UK Annual Total Anthropogenic Sulphur Dioxide (SO<sub>2</sub>) Emissions 1946-2005.

**Usage**

```
data("so2data")
```

**Format**

A data frame with 60 observations on the following 4 variables.

year Year of observation

uk\_tot\_so2 UK annual total anthropogenic SO2 emissions in gigagrams

Luk\_tot\_so2 Log of UK annual total anthropogenic SO2 emissions

DLuk\_tot\_so2 First difference of Log UK annual total anthropogenic SO2 emissions

**Details**

Data reports the total estimated anthropogenic SO2 emissions aggregated over coal, petroleum, biomass combustion, smelting, fuel processing, and other processes.

**Source**

Smith, SJ, J van Aardenne, Z Klimont, RJ Andres, A Volke, and S Delgado Arias. (2011). Anthropogenic Sulfur Dioxide Emissions, 1850-2005: National and Regional Data Set by Source Category, Version 2.86. Data distributed by the NASA Socioeconomic Data and Applications Center (SEDAC), CIESIN, Columbia University, Palisades, New York. Available at

<http://sedac.ciesin.columbia.edu/data/set/haso2-anthro-sulfur-dioxide-emissions-1850-2005-v2-86>

**References**

Pretis, Felix, Reade, James and Sucarrat, Genaro (2018): 'Automated General-to-Specific (GETS) Regression Modeling and Indicator Saturation for Outliers and Structural Breaks'. *Journal of Statistical Software* 86, Number 3, pp. 1-44

Smith, SJ, J van Aardenne, Z Klimont, RJ Andres, A Volke, and S Delgado Arias. (2011). Anthropogenic Sulfur Dioxide Emissions: 1850-2005, *Atmospheric Chemistry and Physics*, 11:1101-1116.

**Examples**

```
data(so2data)

##create annual zoo object:
newso2data<- zooreg(so2data[,-1], start=1946, frequency=1)

##plot UK annual total anthropogenic SO2 emissions:
plot(newso2data$uk_tot_so2)
```

---

`sp500data`*Daily Standard and Poor's 500 index data*

---

**Description**

Daily Standard and Poor's 500 (SP500) index data from 3 January 1950 to 8 March 2016.

**Usage**

```
data("sp500data")
```

**Format**

A data frame with 16652 observations on the following 7 variables:

Date the dates

Open the opening values of the index

High the daily maximum value of the index

Low the daily minimum value of the index

Close the closing values of the index

Volume the traded volume

Adj.Close the adjusted closing values of the index

**Source**

Yahoo Finance, retrieved 9 March 2016

**References**

Pretis, Felix, Reade, James and Sucarrat, Genaro (2018): 'Automated General-to-Specific (GETS) Regression Modeling and Indicator Saturation for Outliers and Structural Breaks'. *Journal of Statistical Software* 86, Number 3, pp. 1-44

**Examples**

```
data(sp500data)
sp500data <- zoo(sp500data[, -1], order.by = as.Date(sp500data[, "Date"]))
plot(window(sp500data, start = as.Date("2000-01-03")))
```

---

`vargaugeiis`*Variance of the Impulse Indicator Saturation Gauge*

---

**Description**

Computes the variance of the gauge (false-positive rate of outliers under the null of no outliers) in impulse indicator saturation based on Jiao and Pretis (2019).

**Usage**

```
vargaugeiis(t.pval, T, infty=FALSE, m=1)
```

**Arguments**

<code>t.pval</code>	numeric, between 0 and 1. Selection p-value used in indicator saturation.
<code>T</code>	integer, sample sized used in indicator saturation.
<code>m</code>	integer, number of iterations in variance computation, default=1
<code>infty</code>	logical, argument used for variance computation

**Details**

The function computes the variance of the Gauge (false-positive rate of outliers in impulse indicator saturation) for a given level of significance of selection (`t.pval`) and sample size (`T`) based on Jiao and Pretis (2019). This is an auxilliary function used within the `outliertest` function.

**Value**

Returns a dataframe of the variance and standard deviation of the gauge, as well the asymptotic variance and standard deviation.

**Author(s)**

Felix Pretis, <http://www.felixpretis.org/>

**References**

Jiao, X. & Pretis, F. (2019). Testing the Presence of Outliers in Regression Models. Discussion Paper.

Pretis, F., Reade, J., & Sucarrat, G. (2018). Automated General-to-Specific (GETS) regression modeling and indicator saturation methods for the detection of outliers and structural breaks. *Journal of Statistical Software*, 86(3).

**See Also**

[isat](#), [outliertest](#)

**Examples**

```
###Computing the variance of the gauge under the null for a sample of T=200 observations:  
vargaugeiis(t.pval=0.05, T=200, infity=FALSE, m=1)
```



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