

Package ‘astsa’

August 21, 2025

Type Package

Title Applied Statistical Time Series Analysis

Version 2.3

Date 2026-09-01

Depends R (>= 3.5)

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BugReports <https://github.com/nickpoison/astsa/issues>

Description Contains data sets and scripts for analyzing time series in both the frequency and time domains including state space modeling as well as supporting the texts Time Series Analysis and Its Applications: With R Examples (5th ed), by R.H. Shumway and D.S. Stoffer. Springer Texts in Statistics, 2025, <[DOI:10.1007/978-3-031-70584-7](https://doi.org/10.1007/978-3-031-70584-7)>, and Time Series: A Data Analysis Approach Using R (2nd ed). Chapman-Hall, 2026, <<https://www.routledge.com/Time-Series-A-Data-Analysis-Approach-Using-R/Shumway-Stoffer/p/book/9781041031642>>. Most scripts are designed to require minimal input to produce aesthetically pleasing output for ease of use in live demonstrations and course work.

URL <https://dsstoffer.github.io/>, <https://nickpoison.github.io/>

License GPL (>= 2)

LazyLoad yes

LazyData yes

NeedsCompilation no

Repository CRAN

Date/Publication 2025-08-21 08:30:18 UTC

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astsa-package	<i>Applied Statistical Time Series Analysis (more than just data)</i>
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Description

Contains data sets and scripts for analyzing time series in both the frequency and time domains including state space modeling as well as supporting the texts [Time Series Analysis and Its Applications: With R Examples \(5th ed, 2025\)](#) and [Time Series: A Data Analysis Approach Using R, \(2nd ed, 2026\)](#). Most scripts are designed to need minimal input to produce aesthetically pleasing output for ease of use in live demonstrations and in course work.

Details

Package:	astsa
Type:	Package
Version:	2.3
Date:	2026-01-01
License:	GPL (>= 2)
LazyLoad:	yes
LazyData:	yes

Warning

If loaded, the dplyr package corrupts the base scripts filter and lag among other things. In this case, whenever you analyze time series data, we suggest you either:

- (1) Detach it if it's loaded but not being used:

```
detach(package:dplyr)
```

- (2) If you want to use it, fix it:

```
library(dplyr, exclude = c("filter", "lag")) # load it without the culprits
dlag    = dplyr::lag      # and fix ...
dfilter = dplyr::filter   # ... the blunders
```

then use dlag and dfilter for dplyr scripts
and lag and filter can be use as originally intended.

- (3) Or just take back the commands:

```
filter = stats::filter
lag     = stats::lag
```

In this case you can still use

```
dlag    = dplyr::lag
dfilter = dplyr::filter
```

for dplyr.

Author(s)

David Stoffer <stoffer@pitt.edu>

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

acf1

*Plot and print ACF or PACF of a time series***Description**

Produces a plot (and a printout) of the sample ACF or PACF. The zero lag value of the ACF is removed.

Usage

```
acf1(series, max.lag = NULL, plot = TRUE, main = NULL, ylim = NULL, pacf = FALSE,
      ylab = NULL, xlab = NULL, na.action = na.pass, ...)
```

Arguments

<code>series</code>	The data (must be more than 2 observations). Does not have to be a time series object.
<code>max.lag</code>	Maximum lag. Can be omitted. Unless $n < 60$, defaults to $\sqrt{n} + 10$ or at least 4 seasons if the series is seasonal.
<code>plot</code>	If TRUE (default), a graph is produced and the values are rounded and listed. If FALSE, no graph is produced and the values are listed but not rounded by the script.
<code>main</code>	Title of graphic; defaults to name of series.
<code>ylim</code>	Specify limits for the y-axis.
<code>pacf</code>	If TRUE, the sample PACF is returned instead of ACF.
<code>ylab</code>	Change y-axis label from default.
<code>xlab</code>	Change x-axis label from default.
<code>na.action</code>	How to handle missing data; default is <code>na.pass</code>
<code>...</code>	Additional arguments passed to tsplot

Details

Will print and/or plot the sample ACF or PACF (if `pacf=TRUE`). The zero lag of the ACF (which is always 1) has been removed. If `plot=TRUE`, a graph is produced and the values are rounded and listed. If `plot=FALSE`, no graph is produced and the values are listed but not rounded by the script. The error bounds are approximate white noise bounds, $-1/n \pm 2/\sqrt{n}$; no other option is given.

Value

ACF	The sample ACF or PACF
-----	------------------------

Author(s)

D.S. Stoffer

References

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In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[acf2](#), [acfm](#), [ccf2](#)

Examples

```
acf1(rnorm(100))

acf1(sarima.sim(ar=.9), pacf=TRUE)

# show it to your mom:
acf1(soi, col=2:7, lwd=4, gg=TRUE)
```

acf2

Plot and print ACF and PACF of a time series

Description

Produces a simultaneous plot (and a printout) of the sample ACF and PACF on the same scale. The zero lag value of the ACF is removed.

Usage

```
acf2(series, max.lag = NULL, plot = TRUE, main = NULL, ylim = NULL,
      na.action = na.pass, ...)
```

Arguments

<code>series</code>	The data (must be more than 2 observations). Does not have to be a time series object.
<code>max.lag</code>	Maximum lag. Can be omitted. Defaults to $\sqrt{n} + 10$ unless $n < 60$. If the series is seasonal, this will be at least 4 seasons by default.
<code>plot</code>	If TRUE (default), a graph is produced and the values are rounded and listed. If FALSE, no graph is produced and the values are listed but not rounded by the script.
<code>main</code>	Title of graphic; defaults to name of series.
<code>ylim</code>	Specify limits for the y-axis.
<code>na.action</code>	How to handle missing data; default is <code>na.pass</code>
<code>...</code>	Additional arguments passed to tsplot

Details

Will print and/or plot the sample ACF and PACF on the same scale. The zero lag of the ACF (which is always 1) has been removed. If `plot=TRUE`, a graph is produced and the values are rounded and listed. If `plot=FALSE`, no graph is produced and the values are listed but not rounded by the script. The error bounds are approximate white noise bounds, $-1/n \pm 2/\sqrt{n}$; no other option is given.

Value

ACF	The sample ACF
PACF	The sample PACF

Author(s)

D.S. Stoffer

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[acf1](#), [acfm](#), [ccf2](#)

Examples

```
acf2(rnorm(100))

acf2(rnorm(100), 25, main='') # no title

acf2(rnorm(100), plot=FALSE)[,'ACF'] # print only ACF

acf2(soi, col=2:7, lwd=4, gg=TRUE) # mother's day present
```

acfm

ACF and CCF for Multiple Time Series

Description

Produces a grid of plots of the sample ACF (diagonal) and CCF (off-diagonal). The values are returned invisibly.

Usage

```
acfm(series, max.lag = NULL, na.action = na.pass, ylim = NULL,
      acf.highlight = TRUE, plot = TRUE, ...)
```

Arguments

<code>series</code>	Multiple time series (at least 2 columns of time series with more than 3 observations).
<code>max.lag</code>	Maximum lag. Can be omitted. Unless $n < 60$, defaults to $\sqrt{n} + 10$ or at least 4 seasons if the series is seasonal.
<code>na.action</code>	How to handle missing data; default is <code>na.pass</code>
<code>ylim</code>	Specify limits for the all correlation axes. If <code>NULL</code> (default) the values are a little wider than the min and max of all values.
<code>acf.highlight</code>	If <code>TRUE</code> (default), the diagonals (ACFs) are highlighted.
<code>plot</code>	If <code>TRUE</code> (default), you get a wonderful graphic.
<code>...</code>	Additional arguments passed to tsplot

Details

Produces a grid of plots of the sample ACF (diagonal) and CCF (off-diagonal). The plots in the grid are estimates of $\text{corr}\{x(t+\text{LAG}), y(t)\}$. Thus x leads y if LAG is positive and x lags y if LAG is negative. If `plot` is `FALSE`, then there is no graphic.

Value

The correlations are returned invisibly.

Author(s)

D.S. Stoffer

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[acf1](#), [acf2](#), [ccf2](#)

Examples

```
acfm(diff(log(econ5)), gg=TRUE)

acfm(diff(log(econ5)), 2, plot=FALSE)
```

ar.boot

*Bootstrap Distribution of AR Model Parameters***Description**

Performs a nonparametric bootstrap to obtain the distribution of the AR model parameters.

Usage

```
ar.boot(series, order.ar, nboot = 500, seed = NULL, plot = TRUE, ...)
```

Arguments

series	time series data (univariate only)
order.ar	autoregression order - must be specified
nboot	number of bootstrap iterations (default is 500)
seed	seed for the bootstrap sampling (default is NULL)
plot	if TRUE (default) and order.ar > 1, returns a scatterplot matrix of the bootstrapped parameters, - the diagonals show a histogram.
...	if plot=TRUE and order.ar > 1, graphical parameters sent to tspairs

Details

For a specified series, finds the bootstrap distribution of the Yule-Walker estimates of $\phi_1, \dots, \phi_p$ in the AR model specified by order.ar,

$$x_t = \mu + \phi_1(x_{t-1} - \mu) + \dots + \phi_p(x_{t-p} - \mu) + w_t,$$

where w_t is white noise. The data are centered by the estimate of μ prior to the bootstrap simulations.

The script displays a number of quantiles of the bootstrapped estimates, the means, the biases, and the root mean squared errors.

Value

Returned invisibly as a list:

phi.star	[[1]] bootstrapped AR parameters
x.sim	[[2]] bootstrapped data
mean.star	[[3]] bootstrapped mean
var.star	[[4]] bootstrapped noise variance
yw.fit	[[5]] results of Yule-Walker fit to the data

Author(s)

D.S. Stoffer

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

Examples

```
## Not run:

u = ar.boot(rec, 2)
dev.new()
tspairs(u[[1]], hist=FALSE) # another view of results
head(u[[1]])                # some bootied AR parameters
head(u[[2]][,1:5])          # some bootied data

## End(Not run)
```

ar.mcmc

*Fit Bayesian AR Model***Description**

Uses Gibbs sampling to fit an AR model to time series data.

Usage

```
ar.mcmc(xdata, porder, n.iter = 1000, n.warmup = 100, plot = TRUE,
        prior_var_phi = 50, prior_sig_a = 1, prior_sig_b = 2, ...)
```

Arguments

xdata	time series data (univariate only)
porder	autoregression order
n.iter	number of iterations for the sampler
n.warmup	number of startup iterations for the sampler (these are removed)
plot	if TRUE (default) returns the draws after warmup (diagonal) and a scatterplot matrix of the draws (off-diagonal)
prior_var_phi	prior variance of the vector of AR coefficients; see details

prior_sig_a	first prior for the variance component; see details
prior_sig_b	second prior for the variance component; see details
...	additional graphic parameters passed to tspairs

Details

Assumes a normal-inverse gamma model,

$$x_t = \phi_0 + \phi_1 x_{t-1} + \cdots + \phi_p x_{t-p} + \sigma z_t,$$

where z_t is standard Gaussian noise. With Φ being the $(p+1)$ -dimensional vector of the ϕ s, the priors are $\Phi \mid \sigma \sim N(0, \sigma^2 V_0)$ and $\sigma^2 \sim IG(a, b)$, where $V_0 = \gamma^2 I$. Defaults are given for the hyperparameters, but the user may choose (a, b) as `(prior_sig_a, prior_sig_b)` and γ^2 as `prior_var_phi`.

The algorithm is efficient and converges quickly. Further details can be found in Chapter 6 of the 5th edition of the Springer text.

Value

In addition to the graphics (if `plot` is `TRUE`), the draws of each parameter (`phi0`, `phi1`, ..., `sigma`) are returned invisibly and means, standard deviations, and various quantiles are displayed.

Author(s)

D.S. Stoffer

Source

Based on the script `arp.mcmc` used in Douc, Moulines, & Stoffer, D. (2014). *Nonlinear Time Series: Theory, Methods and Applications with R Examples*. CRC press. ISBN 9781466502253.

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

Examples

```
## Not run:

u = ar.mcmc(rec, 2)

dev.new()
tspairs(u, hist=FALSE, col.diag=6) # another view
```

```

dev.new()
tsplot(u, ncol=2, col=4, xlab='Index') # traces only

apply(u, 2, ESS)    # effective sample sizes

## End(Not run)

```

ar1miss

AR with Missing Values

Description

Data used in Chapter 6 Problems

Format

The format is: Time-Series [1:100] with NA for missing values.

Details

Simulated AR(1) with $\phi = .9$ and $\sigma_w = 1$ and ~10% of the values missing at random (in this case, there happen to be 15 missing values). A similar data set can be generated as follows:

```

x  = sarima.sim(ar=.9, n=100)
u  = sample(c(NA,0), replace=TRUE, size=100, prob=c(.1,.9))
arm = x + u

```

Another approach if you want exactly 10 missing values is this:

```

arms = sarima.sim(ar=.9, n=100)
arms[sample(1:100, size=10)] = NA

```

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

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arf	<i>Simulated ARFIMA</i>
-----	-------------------------

Description

1000 simulated observations from an ARFIMA(1, 1, 0) model with $\phi = .75$ and $d = .4$.

Format

The format is: Time-Series [1:1000] from 1 to 1000: -0.0294 0.7487 -0.3386 -1.0332 -0.2627 ...

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

arma.spec	<i>Spectral Density of an ARMA Model</i>
-----------	--

Description

Gives the ARMA spectrum, tests for causality, invertibility, and common zeros.

Usage

```
arma.spec(ar = 0, ma = 0, var.noise = 1, n.freq = 500, main = NULL,
          frequency = 1, ylim = NULL, plot = TRUE, ...)
```

Arguments

ar	vector of AR parameters
ma	vector of MA parameters
var.noise	variance of the noise
n.freq	number of frequencies
main	title of graphic; default is "ARMA" with orders "(p, q)"
frequency	for seasonal models, adjusts the frequency scale
ylim	optional; specify limits for the y-axis
plot	if TRUE (default), produces a graphic
...	additional arguments

Details

The basic call is `arma.spec(ar, ma)` where `ar` and `ma` are vectors containing the model parameters. Use `log='y'` if you want the plot on a log scale. If the model is not causal or invertible an error message is given. If there are approximate common zeros, a spectrum will be displayed and a warning will be given; e.g., `arma.spec(ar=.9, ma=-.9)` will yield a warning and the plot will be the spectrum of white noise. Note for example that `arma.spec(ar=1, ma=1)` will bring up an error with warnings that the model is not causal and not invertible.

Value

<code>freq</code>	frequencies - returned invisibly
<code>spec</code>	spectral ordinates - returned invisibly

Author(s)

D.S. Stoffer

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

Examples

```
arma.spec(ar = c(1, -.9), ma = .8)

arma.spec(ar = c(1, -.9), log='y')

arma.spec(ar = -.9, ma=.89, gg=TRUE) # almost white noise

arma.spec(ar=c(rep(0,11),.4), ma=.5, col=5, lwd=3, frequency=12)
```

 ARMAtoAR

Convert ARMA Process to Infinite AR Process

Description

Gives the π -weights in the invertible representation of an ARMA model.

Usage

```
ARMAtoAR(ar = 0, ma = 0, lag.max=20)
```

Arguments

ar	vector of AR coefficients
ma	vector of MA coefficients
lag.max	number of pi-weights desired

Value

A vector of coefficients. For an ARMA model $\phi(B)x_t = \theta(B)w_t$, returns the coefficients $\{\pi_j; j = 1, 2, \dots\}$ in the invertible representation $w_t = \pi(B)x_t$.

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

Examples

```
ARMAtoAR(ar=.9, ma=.5, 10)
```

astsa.col	<i>astsa color palette with transparency or a color wheel</i>
-----------	---

Description

The script does one of two things. [1] Easily modify the opacity level of the astsa colors. [2] Create a color wheel of a chosen number of colors from a base color.

Usage

```
astsa.col(col=1, alpha=1, wheel=FALSE, pie=FALSE, num, sat=NULL, val=NULL, ...)
```

Arguments

col	Either 1. a numerical vector representing the colors to change transparency level - - OR- 2. a color used as a base for a wheel (if a vector, the first color is chosen) - see Examples
alpha	factor in [0, 1] setting the opacity of all colors. Smaller values are more transparent.

wheel	if TRUE, produces a color wheel of num colors from a base color.
pie	if TRUE, produces a pie chart of the chosen colors.
num	an integer specifying the number of desired colors in the wheel. If missing, the user is prompted to enter a number.
sat	factor in [0, 1] setting the 'saturation' (intensity) of the colors. If NULL, the saturation from the base color is used.
val	factor in [0, 1] setting the 'value' (brightness) of the colors. If NULL, the value from the base color is used.
...	other graphical parameters for the pie chart; see pie .

Details

The astsa color palette is attached when the package is attached. The colors follow the R pattern of shades of: (1) black, (2) red, (3) green, (4) blue, (5) cyan, (6) magenta, (7) gold, (8) gray. The opacity of these colors can be changed easily using this script. Values are recycled, e.g., col=9 is the same as col=1.

Additionally, a color wheel can be generated by specifying a base color and inputting the number of desired colors. Using [hsv](#), the script moves the 'hue' around the circle in equal steps holding 'saturation' and 'value' constant. This may be overridden by entering an alternate 'saturation' or 'value'.

In either application, a pie chart can be displayed to help in choosing the desired color scheme.

Value

- [1] a *color vector* using the astsa color palette at the chosen transparency level
- OR -
- [2] a *color wheel* of a chosen number num of colors from a base color

Author(s)

D.S.Stoffer

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

Examples

```
# View the astsa palette
astsa.col(1:8, pie=TRUE)
legend('topright', legend=astsa.col(1:8), fill=1:8, title='Hex Color Code')
```

```

# Plotting 2 series that touch (but in a nice way)
tsplot(cbind(gtemp_land, gtemp_ocean), col=astsa.col(c(4,2), .5), lwd=2, spaghetti=TRUE,
       type='o', pch=20, ylab="Temperature Deviations", addLegend=TRUE, location='topleft',
       legend=c("Land Only", "Ocean Only"), gg=TRUE)

# The hsv values for Dodgerblue3 (or astsa color 4)?
rgb2hsv(col2rgb(4))

# Wheels of fortune
vanna = par(no.readonly = TRUE)
par(mar=rep(0, 4))
layout( matrix(c(1,3,2,3), 2) )
astsa.col(4, wheel=TRUE, num=8, pie=TRUE)
astsa.col(4, wheel=TRUE, num=8, pie=TRUE, sat=.6, val=.9)
astsa.col(2, wheel=TRUE, num=100, pie=TRUE, border=FALSE, labels=NA)

# I'd like to solve the puzzle
par(vanna) # reset graphic device
x = replicate(6, sarima.sim(ar=c(1.5,-.75), n=120))
tsplot(x, spag=TRUE, col=astsa.col(4, alpha=.7, wheel=TRUE, num=6), lwd=12)

```

autoParm

autoParm - Structural Break Estimation Using AR Models

Description

Uses minimum description length (MDL) to fit piecewise AR processes with the goal of detecting changepoints in time series. Optimization is accomplished via a genetic algorithm (GA).

Usage

```

autoParm(xdata, Pi.B = NULL, Pi.C = NULL, PopSize = 70, generation = 70, P0 = 20,
        Pi.P = 0.3, Pi.N = 0.3, NI = 7)

```

Arguments

xdata	time series (of length n at least 100) to be analyzed; the ts attributes are stripped prior to the analysis
Pi.B	probability of being a breakpoint in initial stage; default is 10/n. Does not need to be specified.
Pi.C	probability of conducting crossover; default is (n-10)/n. Does not need to be specified.
PopSize	population size (default is 70); the number of chromosomes in each generation. Does not need to be specified.
generation	number of iterations; default is 70. Does not need to be specified.

P0	maximum AR order; default is 20. If larger than 20, it is reset to 20. Does not need to be specified.
Pi.P	probability of taking parent's gene in mutation; default is 0.3. Does not need to be specified.
Pi.N	probability of taking -1 in mutation; default is 0.3 Does not need to be specified.
NI	number if islands; default is 7. Does not need to be specified.

Details

Details may be found in Davis, Lee, & Rodriguez-Yam (2006). Structural break estimation for nonstationary time series models. JASA, 101, 223-239. [doi:10.1198/016214505000000745](https://doi.org/10.1198/016214505000000745)

Value

Returns three values, (1) the breakpoints including the endpoints, (2) the number of segments, and (3) the segment AR orders. See the examples.

Note

The GA is a stochastic optimization procedure and consequently will give different results at each run. It is a good idea to run the algorithm a few times before coming to a final decision.

Author(s)

D.S. Stoffer

Source

The code is adapted from R code provided to us by Rex Cheung (<https://www.linkedin.com/in/rexcheung>).

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[autoSpec](#)

Examples

```
## Not run:

##-- simulation
x1 = sarima.sim(ar=c(1.69, -.81), n=500)
x2 = sarima.sim(ar=c(1.32, -.81), n=500)
x = c(x1, x2)

##-- look at the data
tsplot(x)

##-- run procedure
autoParm(x)

##-- output (yours will be slightly different -
##--         the nature of GA)
# returned breakpoints include the endpoints
# $breakpoints
# [1] 1 514 1000
#
# $number_of_segments
# [1] 2
#
# $segment_AR_orders
# [1] 2 2

## End(Not run)
```

autoSpec

autoSpec - Changepoint Detection of Narrowband Frequency Changes

Description

Uses changepoint detection to discover if there have been slight changes in frequency in a time series. The autoSpec procedure uses minimum description length (MDL) to do nonparametric spectral estimation with the goal of detecting changepoints. Optimization is accomplished via a genetic algorithm (GA).

Usage

```
autoSpec(xdata, Pi.B = NULL, Pi.C = NULL, PopSize = 70, generation = 70, m0 = 10,
         Pi.P = 0.3, Pi.N = 0.3, NI = 7, taper = .5, min.freq = 0, max.freq = .5)
```

Arguments

xdata	time series (of length n at least 100) to be analyzed; the ts attributes are stripped prior to the analysis
Pi.B	probability of being a breakpoint in initial stage; default is 10/n. Does not need to be specified.

Pi.C	probability of conducting crossover; default is $(n-10)/n$. Does not need to be specified.
PopSize	population size (default is 70); the number of chromosomes in each generation. Does not need to be specified.
generation	number of iterations; default is 70. Does not need to be specified.
m0	maximum width of the Bartlett kernel is $2*m0 + 1$; default is 10. If larger than 20, m0 is reset to 20. Does not need to be specified.
Pi.P	probability of taking parent's gene in mutation; default is 0.3. Does not need to be specified.
Pi.N	probability of taking -1 in mutation; default is 0.3 Does not need to be specified.
NI	number if islands; default is 7. Does not need to be specified.
taper	half width of taper used in spectral estimate; .5 (default) is full taper Does not need to be specified.
min.freq, max.freq	the frequency range (min.freq, max.freq) over which to calculate the Whittle likelihood; the default is (0, .5). Does not need to be specified. If min > max, the roles are reversed, and reset to the default if either is out of range.

Details

Details may be found in Stoffer, D. S. (2023). AutoSpec: Detection of narrowband frequency changes in time series. *Statistics and Its Interface*, 16(1), 97-108. doi:10.4310/21SIH703

Value

Returns three values, (1) the breakpoints including the endpoints, (2) the number of segments, and (3) the segment kernel orders. See the examples.

Note

The GA is a stochastic optimization procedure and consequently will give different results at each run. It is a good idea to run the algorithm a few times before coming to a final decision.

Author(s)

D.S. Stoffer

Source

The genetic algorithm code is adapted from R code provided to us by Rex Cheung (<https://www.linkedin.com/in/rexcheung/>). The code originally supported Aue, Cheung, Lee, & Zhong (2014). Segmented model selection in quantile regression using the minimum description length principle. *JASA*, 109, 1241-1256. A similar version also supported Davis, Lee, & Rodriguez-Yam (2006). Structural break estimation for nonstationary time series models. *JASA*, 101, 223-239.

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[autoParm](#)

Examples

```
## Not run:

##-- simulation
set.seed(1)
num = 500
t   = 1:num
w   = 2*pi/25
d   = 2*pi/150
x1  = 2*cos(w*t)*cos(d*t) + rnorm(num)
x2  = cos(w*t) + rnorm(num)
x   = c(x1,x2)

##-- plot and periodogram (all action below 0.1)
tsplot(x, main='not easy to see the change')
mvspec(x)

##-- run procedure
autoSpec(x, max.freq=.1)

##-- output (yours will be slightly different -
##--          the nature of GA)
# returned breakpoints include the endpoints
# $breakpoints
# [1] 1 503 1000
#
# $number_of_segments
# [1] 2
#
# $segment_kernel_orders_m
# [1] 2 4

##-- plot everything
par(mfrow=c(3,1))
tsplot(x, col=4)
abline(v=503, col=6, lty=2, lwd=2)
```

```

mvspec(x[1:502], kernel=bart(2), taper=.5, main='segment 1', col=4, xlim=c(0,.25))
mvspec(x[503:1000], kernel=bart(4), taper=.5, main='segment 2', col=4, xlim=c(0,.25))

## End(Not run)

```

bart

Bartlett Kernel

Description

Smoothing (triangular) kernel that decreases one unit from the center.

Usage

```
bart(m)
```

Arguments

m non-negative integer specifying the kernel width, which is $2m + 1$. If **m** has length larger than one, the convolution of the kernel is returned.

Details

Uses kernel from the stats package to construct a Bartlett (triangular) kernel of width $2m + 1$; see `help(kernel)` for further details.

Value

Returns an object of class `tskernel` with the coefficients, the kernel dimension, and attribute "Bartlett".

Author(s)

D.S. Stoffer

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

Examples

```

bart(4) # for a list
plot(bart(4), ylim=c(.01,.21)) # for a graph

```

BCJ

*Daily Returns of Three Banks***Description**

Daily returns of three banks, 1. Bank of America [boa], 2. Citibank [citi], and 3. JP Morgan Chase [jpm], from 2005 to 2017.

Format

The format is: Time-Series [1:3243, 1:3] from 2005 to 2017: -0.01378 -0.01157 -0.00155 -0.01084 0.01252 ... with column names "boa" "citi" "jpm" .

Source

Gong & Stoffer (2021). A Note on Efficient Fitting of Stochastic Volatility Models. *Journal of Time Series Analysis*, 42(2), 186-200.

<https://github.com/nickpoison/Stochastic-Volatility-Models>

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

Examples

```
tsplot(BCJ, col=2:4)
```

beamd

*Infrasonic Signal from a Nuclear Explosion***Description**

Infrasonic signal from a nuclear explosion.

Usage

```
data(beamd)
```


Format

A data frame with 2048 observations (rows) on 3 numeric variables (columns): sensor1, sensor2, sensor3.

Details

This is a data frame consisting of three columns (that are not time series objects). The data are an infrasonic signal from a nuclear explosion observed at sensors on a triangular array.

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

birth	<i>U.S. Monthly Live Births</i>
-------	---------------------------------

Description

Monthly live births (adjusted) in thousands for the United States, 1948-1979.

Format

The format is: Time-Series [1:373] from 1948 to 1979: 295 286 300 278 272 268 308 321 313 308 ...

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

blood*Daily Blood Work with Missing Values*

Description

Multiple time series of measurements made for 91 days on the three variables, log(white blood count) [WBC], log(platelet) [PLT] and hematocrit [HCT]. Missing data code is NA.

Format

Time-Series [1:91, 1:3] from 1 to 91: 2.33 1.89 2.08 1.82 1.82 ...

..\$: NULL ..\$: chr [1:3] "WBC" "PLT" "HCT"

Details

This data set is used in Chapter 6 for a missing data example.

Source

Jones, R.H. (1984). Fitting multivariate models to unequally spaced data. In *Time Series Analysis of Irregularly Observed Data*, pp. 158-188. E. Parzen, ed. Lecture Notes in Statistics, 25, New York: Springer-Verlag.

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[HCT](#), [PLT](#), [WBC](#)

Examples

```
tsplot(blood, type='o', pch=19, cex=1.1, col=2:4, gg=TRUE, xlab='day')
```

bnrf1ebv*Nucleotide sequence - BNRF1 Epstein-Barr*

Description

Nucleotide sequence of the BNRF1 gene of the Epstein-Barr virus (EBV): 1=A, 2=C, 3=G, 4=T.
The data are used in Chapter 7.

Format

The format is: Time-Series [1:3954] from 1 to 3954: 1 4 3 3 1 1 3 1 3 1 ...

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

bnrf1hvs*Nucleotide sequence - BNRF1 of Herpesvirus saimiri*

Description

Nucleotide sequence of the BNRF1 gene of the herpesvirus saimiri (HVS): 1=A, 2=C, 3=G, 4=T.
The data are used in Chapter 7.

Format

The format is: Time-Series [1:3741] from 1 to 3741: 1 4 3 2 4 4 3 4 4 4 ...

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

cardox*Monthly Carbon Dioxide Levels at Mauna Loa*

Description

Monthly mean carbon dioxide (in ppm) measured at Mauna Loa Observatory, Hawaii. This is an update to co2 in the datasets package.

Format

The format is: Time-Series [1:781] from 1958 to 2023: 316 317 318 317 316 ...

Details

The carbon dioxide data measured as the mole fraction in dry air, on Mauna Loa constitute the longest record of direct measurements of CO₂ in the atmosphere. They were started by C. David Keeling of the Scripps Institution of Oceanography in March of 1958 at a facility of the National Oceanic and Atmospheric Administration. NOAA started its own CO₂ measurements in May of 1974, and they have run in parallel with those made by Scripps since then. Data are reported as a dry mole fraction defined as the number of molecules of carbon dioxide divided by the number of molecules of dry air multiplied by one million (ppm).

Due to the eruption of the Mauna Loa Volcano, measurements from Mauna Loa Observatory were suspended as of Nov. 29, 2022. Observations starting in December 2022 are from a site at the Maunakea Observatories, approximately 21 miles north of the Mauna Loa Observatory.

Source

<https://gml.noaa.gov/ccgg/trends/>

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

ccf2

*Cross Correlation***Description**

Calculates and plots the sample CCF of two time series.

Usage

```
ccf2(x, y, max.lag = NULL, main = NULL, ylab = "CCF", plot = TRUE,
      na.action = na.pass, type = c("correlation", "covariance"), ...)
```

Arguments

<code>x, y</code>	univariate time series
<code>max.lag</code>	maximum lag for which to calculate the CCF. Can be omitted. Unless $n < 50$, defaults to $\sqrt{n} + 10$ or at least 3 seasons if the series is seasonal.
<code>main</code>	plot title - if NULL, uses x and y names
<code>ylab</code>	vertical axis label; default is 'CCF'
<code>plot</code>	if TRUE (default) a graphic is produced and the values are returned invisibly. Otherwise, the values are returned.
<code>na.action</code>	how to handle missing values; default is <code>na.pass</code>
<code>type</code>	default is cross-correlation; an option is cross-covariance
<code>...</code>	additional arguments passed to tsplot

Details

This will produce a graphic of the sample $\text{corr}[x(t+\text{lag}), y(t)]$ from $-\text{max.lag}$ to max.lag . Also, the (rounded) values of the CCF are returned invisibly unless `plot=FALSE`. Similar details apply to the cross-covariance.

Author(s)

D.S. Stoffer

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[pre.white](#), [acf1](#), [acf2](#), [acfm](#)

Examples

```
ccf2(soi, rec, plot=FALSE, max.lag=6) # now you see it
ccf2(soi, rec)                        # now you don't

# happy birthday mom
ccf2(soi, rec, col=astsa.col(4, wheel=TRUE, num=6), lwd=4, gg=TRUE)
```

chicken

Monthly price of a pound of chicken

Description

Poultry (chicken), Whole bird spot price, Georgia docks, US cents per pound

Format

The format is: Time-Series [1:180] from August 2001 to July 2016: 65.6 66.5 65.7 64.3 63.2 ...

Source

<https://www.indexmundi.com/commodities/>

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

climhyd*Lake Shasta inflow data*

Description

Lake Shasta inflow data. This is a data frame.

Format

A data frame with 454 observations (rows) on the following 6 numeric variables (columns): Temp, DewPt, CldCvr, WndSpd, Precip, Inflow.

Details

The data are 454 months of measured values for the climatic variables: air temperature, dew point, cloud cover, wind speed, precipitation, and inflow, at Lake Shasta, California.

Lake Shasta is a man-made reservoir that is situated on ancestral lands of the Shasta Indian Nation. The area around the lake was once a thriving homeland for the Shasta. The construction of the reservoir submerged many of these significant cultural sites, leading to displacement and hardship for the Shasta people. The California Gold Rush drew in outsiders in the late 1840s. For the Shasta, this was a devastating process as thousands of miners who did not respect the Shasta or their homeland, operated along various waterways. Introduction of diseases and fighting against white people, rapidly reduced the number of Shasta. For more information, see <https://www.legendsofamerica.com/shasta-indians>.

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

cmort*Cardiovascular Mortality from the LA Pollution study*

Description

Average weekly cardiovascular mortality in Los Angeles County; 508 six-day smoothed averages obtained by filtering daily values over the 10 year period 1970-1979.

Format

The format is: Time-Series [1:508] from 1970 to 1980: 97.8 104.6 94.4 98 95.8 ...

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[lap](#), [lap.xts](#)

cpg

Hard Drive Cost per GB

Description

Median annual cost per gigabyte (GB) of storage.

Format

The format is: Time-Series [1:29] from 1980 to 2008: 213000.00 295000.00 260000.00 175000.00 160000.00 ...

Source

The data are retail prices per GB taken from a sample of manufacturers listed at <https://mkomo.com/cost-per-gigabyte>. From there, the median cost per GB was computed by year.

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

detrend	<i>Detrend a Time Series</i>
---------	------------------------------

Description

Returns a time series with the trend removed. The trend can be estimated using polynomial regression or using a lowess fit.

Usage

```
detrend(series, order = 1, lowess = FALSE, lowspan = 2/3)
```

Arguments

series	The time series to be detrended.
order	Order of the polynomial used to estimate the trend with a linear default (order=1) unless lowess is TRUE.
lowess	If TRUE, lowess is used to find the trend. The default is FALSE.
lowspan	The smoother span used for lowess.

Value

The detrended series is returned.

Author(s)

D.S. Stoffer

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[trend](#)

Examples

```
tsplot( cbind(salmon, detrend(salmon)), gg=TRUE, main='Norwegian Salmon USD/KG' )
```

djia	<i>Dow Jones Industrial Average</i>
------	-------------------------------------

Description

Daily DJIA values from April 2006 - April 2016

Format

The format is: xts [1:2518, 1:5] 11279 11343 11347 11337 11283 ...
 - attr(*, "class")= chr [1:2] "xts" "zoo"
 ..\$: chr [1:5] "Open" "High" "Low" "Close" "Volume"

Source

The data were obtained via the TTR package and Yahoo financial data. Unfortunately, this does not work now. It seems like the R package quantmod is a good bet and Yahoo still has financial data.

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

dna2vector	<i>Convert DNA Sequence to Indicator Vectors</i>
------------	--

Description

Takes a string (e.g., a DNA sequence) of general form (e.g., FASTA) and converts it to a sequence of indicator vectors for use with the Spectral Envelope (specenv).

Usage

```
dna2vector(data, alphabet = NULL)
```

Arguments

data	A single string.
alphabet	The particular alphabet being used. The default is alphabet=c("A", "C", "G", "T").

Details

Takes a string of categories and converts it to a matrix of indicators. The data can then be used by the script [specenv](#), which calculates the Spectral Envelope of the sequence (or subsequence). Many different type of sequences can be used, including FASTA and GenBank, as long as the data is a string of categories.

The indicator vectors (as a matrix) are returned invisibly in case the user forgets to put the results in an object wherein the screen would scroll displaying the entire sequence. In other words, the user should do something like `xdata = dna2vector(data)` where `data` is the original sequence.

If the DNA sequence is in a FASTA file, say `sequence.fasta`, the following code can be used to read the data into the session, create the indicator sequence and save it as a compressed R data file:

```
fileName <- 'sequence.fasta'      # name of FASTA file
data      <- readChar(fileName, file.info(fileName)$size) # input the sequence
myseq     <- dna2vector(data)      # convert it to indicators

##== to compress and save the data ==##
save(myseq, file='myseq.rda')
##== and then load it when needed ==##
load('myseq.rda')
```

Value

matrix of indicator vectors; returned invisibly

Author(s)

D.S. Stoffer

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[specenv](#)

Examples

```
# Epstein-Barr virus (entire sequence included in astsa)
xdata = dna2vector(EBV)
head(xdata)

# part of EBV with 1, 2, 3, 4 for "A", "C", "G", "T"
```

```
xdata = dna2vector(bnrf1ebv)
head(xdata)

# raw GenBank sequence
data <-
c("1 agaattcgctc ttgctctatt cacccttact tttcttcttg cccgttctct ttcttagtat
  61 gaatccagta tgcctgcctg taattgttgc gccctacctc ttttgctgg cggctattgc")
xdata = dna2vector(data, alphabet=c('a', 'c', 'g', 't'))
head(xdata)

# raw FASTA sequence
data <-
c("AGAATTCGCTCTTGCTCTATTACCCCTTACTTTTCTTCTTGCCGTTCTCTTCTTAGTATGAATCCAGTA
  TGCCTGCCTGTAATTGTTGCGCCCTACCTCTTTGGCTGGCGGCTATTGCCGCTCGTGTTTCACGGCCT")
xdata = dna2vector(data)
head(xdata)
```

EBV

*Entire Epstein-Barr Virus (EBV) Nucleotide Sequence***Description**

EBV nucleotide sequence - 172281 bp as a single string

Format

The format is: chr "AGAATTCGCTCTT ..."

Note

EBV is not useful on its own, but using 'dna2vector', different regions can be explored. For example, `ebv = dna2vector(EBV)`

Source

<https://www.ncbi.nlm.nih.gov/nuccore/V01555.2>

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[dna2vector](#)

econ5

*Five Quarterly Economic Series***Description**

Multiple time series of quarterly U.S. unemployment, GNP, consumption, and government and private investment, from 1948-III to 1988-II.

Format

Multiple time series with 161 observations (rows) on the following 5 numeric variables (columns): unemp, gnp, consum, govinv, prinv.

Source

Young, P.C. and Pedregal, D.J. (1999). Macro-economic relativity: government spending, private investment and unemployment in the USA 1948-1998. *Structural Change and Economic Dynamics*, 10, 359-380.

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

EM

*EM Algorithm for State Space Models***Description**

Estimation of the parameters in general linear state space models via the EM algorithm. Missing data may be entered as NA or as zero (0), however, use NAs if zero (0) can be an observation. Inputs in both the state and observation equations are allowed. This script replaces EM0 and EM1.

Usage

```
EM(y, A, mu0, Sigma0, Phi, Q, R, Ups = NULL, Gam = NULL, input = NULL,
  max.iter = 100, tol = 1e-04)
```

Arguments

y	data matrix (n × q), vector or time series, n = number of observations, q = number of series. Use NA or zero (0) for missing data, however, use NAs if zero (0) can be an observation.
A	measurement matrices; can be constant or an array with dimension dim=c(q,p,n) if time varying. Use NA or zero (0) for missing data.
mu0	initial state mean vector (p × 1)
Sigma0	initial state covariance matrix (p × p)
Phi	state transition matrix (p × p)
Q	state error matrix (p × p)
R	observation error matrix (q × q - diagonal only)
Ups	state input matrix (p × r); leave as NULL (default) if not needed
Gam	observation input matrix (q × r); leave as NULL (default) if not needed
input	NULL (default) if not needed or a matrix (n × r) of inputs having the same row dimension (n) as y
max.iter	maximum number of iterations
tol	relative tolerance for determining convergence

Details

This script replaces EM0 and EM1 by combining all cases and allowing inputs in the state and observation equations. It uses version 1 of the new [Ksmooth](#) script (hence correlated errors is not allowed).

The states x_t are p-dimensional, the data y_t are q-dimensional, and the inputs u_t are r-dimensional for $t = 1, \dots, n$. The initial state is $x_0 \sim N(\mu_0, \Sigma_0)$.

The general model is

$$\begin{aligned} x_t &= \Phi x_{t-1} + \Upsilon u_t + w_t & w_t &\sim iid N(0, Q) \\ y_t &= A_t x_{t-1} + \Gamma u_t + v_t & v_t &\sim iid N(0, R) \end{aligned}$$

where $w_t \perp v_t$. The observation noise covariance matrix is assumed to be diagonal and it is forced to diagonal otherwise.

The measurement matrices A_t can be constant or time varying. If time varying, they should be entered as an array of dimension dim = c(q,p,n). Otherwise, just enter the constant value making sure it has the appropriate $q \times p$ dimension.

Value

Phi	Estimate of Phi
Q	Estimate of Q
R	Estimate of R
Ups	Estimate of Upsilon (NULL if not used)
Gam	Estimate of Gamma (NULL if not used)

<code>mu0</code>	Estimate of initial state mean
<code>Sigma0</code>	Estimate of initial state covariance matrix
<code>like</code>	-log likelihood at each iteration
<code>niter</code>	number of iterations to convergence
<code>cvg</code>	relative tolerance at convergence

Note

The script does not allow for constrained estimation directly, however, constrained estimation is possible with some extra manipulations. There is an example of constrained estimation using EM at [FUN WITH ASTSA](#), where the fun never stops.

Author(s)

D.S. Stoffer

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[Kfilter](#), [Ksmooth](#)

Examples

```
# example used for ssm()
# x[t] = Ups + Phi x[t-1] + w[t]
# y[t] = x[t] + v[t]
y = gtemp_land
A = 1; Phi = 1.03; Ups = 0.01
Q = 0.0001; R = 0.1
mu0 = -0.5; Sigma0 = 0.001
input = rep(1, length(y))
( em = EM(y, A, mu0, Sigma0, Phi, Q, R, Ups, Gam=NULL, input) )
```

ENSO*El Nino - Southern Oscillation Index*

Description

Southern Oscillation Index (SOI), 1/1951 to 10/2022; anomalies are departures from the 1981-2010 base period.

Format

The format is: Time-Series [1:862] from 1951 to 2022: 2.0 1.1 -0.3 -0.8 -1.1 -0.7 -1.5 -0.3 -0.7 -0.7 ...

Details

The El Niño - Southern Oscillation (ENSO) is a recurring climate pattern involving changes in the temperature of waters in the central and eastern tropical Pacific Ocean. This data set is an update to [soi](#).

Source

<https://www.ncei.noaa.gov/access/monitoring/enso/soi>

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[soi](#)

EQ5

*Seismic Trace of Earthquake number 5***Description**

Seismic trace of an earthquake [two phases or arrivals along the surface, the primary wave ($t = 1, \dots, 1024$) and the shear wave ($t = 1025, \dots, 2048$)] recorded at a seismic station.

Format

The format is: Time-Series [1:2048] from 1 to 2048: 0.01749 0.01139 0.01512 0.01477 0.00651 ...

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[eqexp](#)

EQcount

*Earthquake Counts***Description**

Series of annual counts of major earthquakes (magnitude 7 and above) in the world between 1900 and 2006.

Format

The format is: Time-Series [1:107] from 1900 to 2006: 13 14 8 10 16 26 ...

Source

Zucchini and MacDonald (2009). Hidden Markov Models for Time Series: An Introduction using R. CRC Press.

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

eqexp

Earthquake and Explosion Seismic Series

Description

This is a data frame of the earthquake and explosion seismic series used throughout the text.

Format

A data frame with 2048 observations (rows) on 17 variables (columns). Each column is a numeric vector.

Details

The matrix has 17 columns, the first eight are earthquakes, the second eight are explosions, and the last column is the Novaya Zemlya event of unknown origin.

The column names are: EQ1, EQ2, ..., EQ8; EX1, EX2, ..., EX8; NZ. The first 1024 observations correspond to the P wave, the second 1024 observations correspond to the S wave.

All events in the data set were on or near land and were distributed uniformly over Scandinavia so as to minimize the possibility that discriminators might be keying on location or land-sea differences. The events are earthquakes ranging in magnitude from 2.74 to 4.40 and explosions in the range 2.13 to 2.19. Also added is an event of uncertain origin that was located in the Novaya Zemlya region of Russia. All events except the Russian event occurred in the Scandinavian peninsula and were recorded by seismic arrays located in Norway by Norwegian and Arctic experimental seismic stations (NORESS, ARCESS) and in Finland by Finnish experimental seismic stations (FINESS).

No.	Type	Date	Array	Magnitude	Latitude	Longitude
1	EQ	6/16/92	FINESS	3.22	65.5	22.9
2	EQ	8/24/91	ARCESS	3.18	65.7	32.1
3	EQ	9/23/91	NORESS	3.15	64.5	21.3
4	EQ	7/4/92	FINESS	3.60	67.8	15.1
5	EQ	2/19/92	ARCESS	3.26	59.2	10.9
6	EQ	4/13/92	NORESS	4.40	51.4	6.1
7	EQ	4/14/92	NORESS	3.38	59.5	5.9
8	EQ	5/18/92	NORESS	2.74	66.9	13.7
9	EX	3/23/91	ARCESS	2.85	69.2	34.3
10	EX	4/13/91	FINESS	2.60	61.8	30.7

11	EX	4/26/91	ARCESS	2.95	67.6	33.9
12	EX	8/3/91	ARCESS	2.13	67.6	30.6
13	EX	9/5/91	ARCESS	2.32	67.1	21.0
14	EX	12/10/91	FINCESS	2.59	59.5	24.1
15	EX	12/29/91	ARCESS	2.96	69.4	30.8
16	EX	3/25/92	NORESS	2.94	64.7	30.8
17	NZ	12/31/92	NORESS	2.50	73.6	55.2

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

Examples

```
## Not run:

# view all series
# first 2 rows EQs - second 2 rows EXs
# 5th row NZ event

tsplot(eqexp, ncol=4, col=1:8)

## End(Not run)
```

ESS	<i>Effective Sample Size (ESS)</i>
-----	------------------------------------

Description

Estimates the ESS of a given vector of samples.

Usage

```
ESS(trace, tol = 1e-08, BIC = TRUE, digits=2)
```

Arguments

<code>trace</code>	vector of sampled values from an MCMC run (univariate only)
<code>tol</code>	ESS is returned as zero if the estimated spectrum at frequency zero is less than this value
<code>BIC</code>	if TRUE (default), <code>spec0</code> is obtained using BIC; otherwise, AIC is used. See the details.
<code>digits</code>	integer indicating the approximate number of decimal places to be used

Details

Uses `spec.ic` to estimate the spectrum of the input at frequency zero (`spec0`). Then, ESS is estimated as `ESS = length(trace)*var(trace)/spec0`. See Examples for multivariate case.

ESS is discussed in detail in Example 6.31 of *Time Series Analysis and Its Applications: With R Examples* (5th ed, 2025).

Value

Returns the estimated ESS of the input.

Author(s)

D.S. Stoffer

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

Examples

```
# How autocorrelation affects ESS
x = matrix(NA, 500, 3) # sample size is 500
phi = c(0,.3,.9)       # no, low, high
for (i in 1:3) x[,i] = sarima.sim(ar=phi[i])
apply(x, 2, ESS, digits=-1)
```

EXP6

Seismic Trace of Explosion number 6

Description

Seismic trace of an explosion [two phases or arrivals along the surface, the primary wave ($t = 1, \dots, 1024$) and the shear wave ($t = 1025, \dots, 2048$)] recorded at a seismic station.

Format

The format is: Time-Series [1:2048] from 1 to 2048: -0.001837 -0.000554 -0.002284 -0.000303 -0.000721 ...

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[eqexp](#)

FDR

Basic False Discovery Rate

Description

Computes the basic false discovery rate given a vector of p-values and returns the index of the maximal p-value satisfying the FDR condition.

Usage

```
FDR(pvals, qlevel = 0.05)
```

Arguments

pvals	a vector of pvals on which to conduct the multiple testing
qlevel	the proportion of false positives desired

Value

fdr.id	NULL if no significant tests, or the index of the maximal p-value satisfying the FDR condition.
--------	---

Note

This is used primarily in Chapter 7.

Source

Built off of <https://www.stat.berkeley.edu/~paciorek/code/fdr/fdr.R>.

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

ffbs

Forward Filtering Backward Sampling

Description

FFBS algorithm for state space models

Usage

```
ffbs(y, A, mu0, Sigma0, Phi, sQ, sR, Ups = NULL, Gam = NULL, input = NULL)
```

Arguments

y	Data matrix, vector or time series.
A	Observation matrix. Can be constant or an array with dim=c(q,p,n) if time varying.
mu0	Initial state mean.
Sigma0	Initial state covariance matrix.
Phi	State transition matrix.
sQ	State error covariance matrix is $Q = sQ \%*\% t(sQ)$ – see details below. In the univariate case, it is the standard deviation.
sR	Observation error covariance matrix is $R = sR \%*\% t(sR)$ – see details below. In the univariate case, it is the standard deviation.
Ups	State input matrix.
Gam	Observation input matrix.
input	matrix or vector of inputs having the same row dimension as y.

Details

For a linear state space model, the FFBS algorithm provides a way to sample a state sequence $x_{0:n}$ from the posterior $\pi(x_{0:n} \mid \Theta, y_{1:n})$ with parameters Θ and data $y_{1:n}$.

The general model is

$$x_t = \Phi x_{t-1} + \Upsilon u_t + sQ w_t \quad w_t \sim iid N(0, I)$$

$$y_t = A_t x_{t-1} + \Gamma u_t + sR v_t \quad v_t \sim iid N(0, I)$$

where $w_t \perp v_t$. Consequently the state noise covariance matrix is $Q = sQ sQ'$ and the observation noise covariance matrix is $R = sR sR'$ and sQ, sR do not have to be square as long as everything is conformable.

x_t is p-dimensional, y_t is q-dimensional, and u_t is r-dimensional. Note that $sQ w_t$ has to be p-dimensional, but w_t does not, and $sR v_t$ has to be q-dimensional, but v_t does not.

Value

Xs	An array of sampled states
X0n	The sampled initial state (because R is 1-based)

Note

The script uses [Kfilter](#). If A_t is constant wrt time, it is not necessary to input an array; see the example. The example below is just one pass of the algorithm; see the example at [FUN WITH ASTSA](#) for the real fun.

Author(s)

D.S. Stoffer

Source

Chapter 6 of the Shumway and Stoffer Springer text.

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

Examples

```
## Not run:

## -- this is just one pass --##
# generate some data
set.seed(1)
sQ = 1; sR = 3; n = 100
mu0 = 0; Sigma0 = 10; x0 = rnorm(1,mu0,Sigma0)
w = rnorm(n); v = rnorm(n)
x = c(x0 + sQ*w[1]); y = c(x[1] + sR*v[1]) # initialize
for (t in 2:n){
  x[t] = x[t-1] + sQ*w[t]
  y[t] = x[t] + sR*v[t]
```

```

    }

## run one pass of FFBS, plot data, states and sampled states
run = ffbs(y, A=1, mu0=0, Sigma0=10, Phi=1, sQ=1, sR=3)
tsplot(cbind(y,run$Xs), spaghetti=TRUE, type='o', col=c(8,4), pch=c(1,NA))
legend('topleft', legend=c("y(t)", "xs(t)"), lty=1, col=c(8,4), bty="n", pch=c(1,NA))

## End(Not run)

```

flu

*Monthly pneumonia and influenza deaths in the U.S., 1968 to 1978.***Description**

Monthly pneumonia and influenza deaths per 10,000 people in the United States for 11 years, 1968 to 1978.

Usage

```
data(flu)
```

Format

The format is: Time-Series [1:132] from 1968 to 1979: 0.811 0.446 0.342 0.277 0.248 ...

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

fmri

*fMRI - complete data set***Description**

Data (as a vector list) from an fMRI experiment in pain, listed by location and stimulus. The data are BOLD signals when a stimulus was applied for 32 seconds and then stopped for 32 seconds. The signal period is 64 seconds and the sampling rate was one observation every 2 seconds for 256 seconds ($n = 128$). The number of subjects under each condition varies.

Details

The LOCATIONS of the brain where the signal was measured were [1] Cortex 1: Primary Somatosensory, Contralateral, [2] Cortex 2: Primary Somatosensory, Ipsilateral, [3] Cortex 3: Secondary Somatosensory, Contralateral, [4] Cortex 4: Secondary Somatosensory, Ipsilateral, [5] Caudate, [6] Thalamus 1: Contralateral, [7] Thalamus 2: Ipsilateral, [8] Cerebellum 1: Contralateral and [9] Cerebellum 2: Ipsilateral.

The TREATMENTS or stimuli (and number of subjects in each condition) are [1] Awake-Brush (5 subjects), [2] Awake-Heat (4 subjects), [3] Awake-Shock (5 subjects), [4] Low-Brush (3 subjects), [5] Low-Heat (5 subjects), and [6] Low-Shock (4 subjects). Issue the command `summary(fmri)` for further details. In particular, awake (Awake) or mildly anesthetized (Low) subjects were subjected levels of periodic brushing (Brush), application of heat (Heat), and mild shock (Shock) effects.

As an example, `fmri$LT6` (Location 1, Treatment 6) will show the data for the four subjects receiving the Low-Shock treatment at the Cortex 1 location; note that `fmri[[6]]` will display the same data.

Source

Joseph F. Antognini, Michael H. Buonocore, Elizabeth A. Disbrow, Earl Carstens, Isoflurane anesthesia blunts cerebral responses to noxious and innocuous stimuli: a fMRI study, *Life Sciences*, Volume 61, Issue 24, 1997, Pages PL349-PL354, ISSN 0024-3205, [https://doi.org/10.1016/S0024-3205\(97\)00960-0](https://doi.org/10.1016/S0024-3205(97)00960-0).

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

fmri1

fMRI Data Used in Chapter 1

Description

A data frame that consists of average fMRI BOLD signals at eight locations.

Usage

```
data(fmri1)
```

Format

The format is: mts [1:128, 1:9]

Details

Multiple time series consisting of fMRI BOLD signals at eight locations (in columns 2-9, column 1 is time period), when a stimulus was applied for 32 seconds and then stopped for 32 seconds. The signal period is 64 seconds and the sampling rate was one observation every 2 seconds for 256 seconds ($n = 128$). The columns are labeled: "time" "cort1" "cort2" "cort3" "cort4" "thal1" "thal2" "cere1" "cere2".

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).
The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.
In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.
The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[fmri](#)

gas	<i>Gas Prices</i>
-----	-------------------

Description

New York Harbor conventional regular gasoline weekly spot price FOB (in cents per gallon) from 2000 to mid-2010.

Format

The format is: Time-Series [1:545] from 2000 to 2010: 70.6 71 68.5 65.1 67.9 ...

Details

Pairs with series oil

Source

Data were obtained from: https://www.eia.gov/dnav/pet/pet_pri_spt_s1_w.htm

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).
The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.
In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.
The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also[oil](#)

gdp

*Quarterly U.S. GDP***Description**

Seasonally adjusted quarterly U.S. GDP from 1947(1) to 2018(3).

Format

The format is: Time-Series [1:287] from 1947 to 2018: 2033 2028 2023 2055 2086 ...

Source

<https://tradingeconomics.com/united-states/gdp>

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also[GDP, GNP, gnp](#)

GDP23

*Quarterly U.S. GDP - updated to 2023***Description**

Seasonally adjusted quarterly U.S. GDP from 1947(1) to 2023(1).

Format

The format is: Time-Series [1:305] from 1947 to 2023: 243.164 245.968 249.585 259.745 ...

Source

<https://fred.stlouisfed.org/series/GDP>

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[gdp](#), [GNP](#), [gnp](#)

gnp	<i>Quarterly U.S. GNP</i>
-----	---------------------------

Description

Seasonally adjusted quarterly U.S. GNP from 1947(1) to 2002(3).

Format

The format is: Time-Series [1:223] from 1947 to 2002: 1489 1497 1500 1524 1547 ...

Source

<https://fred.stlouisfed.org/series/GNP>

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[GNP](#), [GDP](#), [gdp](#)

GNP23

*Quarterly U.S. GNP - updated to 2023***Description**

Seasonally adjusted quarterly U.S. GNP from 1947(1) to 2003(1).

Format

The format is: Time-Series [1:305] from 1947 to 2023: 244.142 247.063 250.716 260.981 ...

Source

<https://fred.stlouisfed.org/series/GNP>

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[gnp](#), [GDP](#), [gdp](#)

Grid

*A Better Add Grid to a Plot***Description**

Adds a grid to an existing plot with major and minor ticks. Works like R graphics `grid()` but the grid lines are solid and gray and minor ticks are produced by default.

Usage

```
Grid(nx = NULL, ny = nx, col = gray(0.9), lty = 1, lwd = par("lwd"), equilogs = TRUE,
     minor = TRUE, nxm = 2, nym = 2, tick.ratio = 0.5, xm.grid = TRUE, ym.grid = TRUE, ...)
```

Arguments

<code>nx, ny</code>	number of cells of the grid in x and y direction. When NULL, as per default, the grid aligns with the tick marks on the corresponding default axis (i.e., tickmarks as computed by <code>axTicks</code>). When NA, no grid lines are drawn in the corresponding direction.
<code>col</code>	color of the grid lines.
<code>lty</code>	line type of the grid lines.
<code>lwd</code>	line width of the grid lines.
<code>equilogs</code>	logical, only used when log coordinates and alignment with the axis tick marks are active. Setting <code>equilogs = FALSE</code> in that case gives non equidistant tick aligned grid lines.
<code>minor</code>	logical with TRUE (default) adding minor ticks.
<code>nxm, nym</code>	number of intervals in which to divide the area between major tick marks on the x-axis (y-axis). If <code>minor=TRUE</code> , should be > 1 or no minor ticks will be drawn.
<code>tick.ratio</code>	ratio of lengths of minor tick marks to major tick marks. The length of major tick marks is retrieved from <code>par("tck")</code> .
<code>xm.grid, ym.grid</code>	if TRUE (default), adds grid lines at minor x-axis, y-axis ticks.
<code>...</code>	other graphical parameters;

Author(s)

D.S. Stoffer

Source

The code for `grid()` in R graphics and `minor.tick()` from the Hmisc package were combined.

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[grid](#)

gtemp.month

*Monthly global average surface temperatures by year***Description**

Monthly global average surface temperatures by year. The temperature of the air measured 2 meters above the ground, encompassing land, sea, and in-land water surfaces.

Format

A data frame with 12 monthly observations (as rows) for the years 1975-2023 (as columns in reverse order).

Details

Temperature of air at 2m above the surface of land, sea or in-land waters. 2m temperature is calculated by interpolating between the lowest model level and the Earth's surface, taking account of the atmospheric conditions. Technical details at <https://cds.climate.copernicus.eu/datasets/reanalysis-era5-pressure>

Source

<https://ourworldindata.org/grapher/monthly-average-surface-temperatures-by-year>

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

Examples

```
## Not run:

# functional data plot showing warming trend
tsplot(gtemp.month, spaghetti=TRUE, col=rainbow(49, start=.2, v=.8, rev=TRUE), ylab='\u00b0C',
       xlab='Month', xaxt='n', main='Mean Monthly Global Temperature', lwd=c(3,rep(1,47),3))
axis(1, labels=Months, at=1:12)
text(10, 13, '1975')
text(10.3, 15.5, '2023')

## End(Not run)
```

gtemp_both	<i>Global mean land and open ocean temperature deviations, 1850-2023</i>
------------	--

Description

Annual temperature anomalies (in degrees centigrade) averaged over the Earth's land and ocean area from 1850 to 2023. Anomalies are with respect to the 1991-2020 average.

Format

The format is: Time-Series [1:174] from 1850 to 2023: -0.24 -0.25 -0.27 -0.15 -0.05 -0.16 -0.29 -0.32 -0.19 -0.04 ...

Source

<https://www.ncei.noaa.gov/access/monitoring/climate-at-a-glance/global/time-series/>

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[gtemp_land](#), [gtemp_ocean](#)

gtemp_land	<i>Global mean land temperature deviations, 1850-2023</i>
------------	---

Description

Annual temperature anomalies (in degrees centigrade) averaged over the Earth's land area from 1850 to 2023. Anomalies are with respect to the 1991-2020 average.

Format

The format is: Time-Series [1:174] from 1850 to 2023: -0.50 -0.60 -0.50 -0.50 -0.20 -0.50 -0.80 -0.40 -0.40 -0.10 ...

Source

<https://www.ncei.noaa.gov/access/monitoring/climate-at-a-glance/global/time-series/>

References

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The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[gtemp_ocean](#), [gtemp_both](#)

gtemp_ocean

Global mean ocean temperature deviations, 1850-2023

Description

Annual sea surface temperature anomalies averaged over the part of the ocean that is free of ice at all times (open ocean) from 1850 to 2023. Anomalies are with respect to the 1991-2020 average.

Format

The format is: Time-Series [1:174] from 1850 to 2023: -0.12 -0.08 -0.14 0.04 0.04 0.00 -0.05 -0.27 -0.09 0.01 ...

Source

<https://www.ncei.noaa.gov/access/monitoring/climate-at-a-glance/global/time-series/>

References

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The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[gtemp_land](#), [gtemp_both](#)

Hare	<i>Snowshoe Hare</i>
------	----------------------

Description

This is one of the classic studies of predator-prey interactions, the 90-year data set is the number, in thousands, of snowshoe hare pelts purchased by the Hudson's Bay Company of Canada. While this is an indirect measure of predation, the assumption is that there is a direct relationship between the number of pelts collected and the number of hare and lynx in the wild.

Format

The format is: Time-Series [1:91] from 1845 to 1935: 19.6 19.6 19.6 12 28 ...

Note

This data set pairs with [Lynx](#). The data are in units of one thousand.

Source

From Odum's "Fundamentals of Ecology", p. 191. Data listed at:
<http://people.whitman.edu/~hundleedr/courses/M250F03/M250.html>
 scroll down to: Chapter 6, Difference Equations.

NB: For some reason, there is not a secure link to the site above.

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[Lynx](#)

HCT

*Hematocrit Levels***Description**

HCT: Measurements made for 91 days on the three variables, log(white blood count) [WBC], log(platelet) [PLT] and hematocrit [HCT]. Missing data code is 0 (zero).

Format

The format is: Time-Series [1:91] from 1 to 91: 30 30 28.5 34.5 34 32 30.5 31 33 34 ...

Details

See Examples 6.1 and 6.9 for more details.

Source

Jones, R.H. (1984). Fitting multivariate models to unequally spaced data. In *Time Series Analysis of Irregularly Observed Data*, pp. 158-188. E. Parzen, ed. Lecture Notes in Statistics, 25, New York: Springer-Verlag.

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[blood](#), [PLT](#), [WBC](#)

hor

*Hawaiian occupancy rates***Description**

Quarterly Hawaiian hotel occupancy rate (percent of rooms occupied) from 1982-I to 2015-IV

Format

The format is: Time-Series [1:136] from 1982 to 2015: 79 65.9 70.9 66.7 ...

Source

<https://dbedt.hawaii.gov/economic/qser/tourism/>

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

Examples

```
tsplot(hor, type='c')           # plot data and
text(hor, labels=1:4, col=c(1,4,2,6), cex=.9) # add quarter labels
```

 jj

Johnson and Johnson Quarterly Earnings Per Share

Description

Johnson and Johnson quarterly earnings per share, 84 quarters (21 years) measured from the first quarter of 1960 to the last quarter of 1980.

Format

The format is: Time-Series [1:84] from 1960 to 1981: 0.71 0.63 0.85 0.44 0.61 0.69 0.92 0.55 0.72 0.77 ...

Details

The data were provided (personal communication) by Professor Paul Griffin, <https://gsm.ucdavis.edu/profile/paul-griffin> of the Graduate School of Management, University of California, Davis. This data set is also included with the R distribution as JohnsonJohnson.

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

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The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

Kfilter

*Quick Kalman Filter***Description**

Returns both the predicted and filtered values for various linear state space models; it also evaluates the likelihood at the given parameter values.

Usage

```
Kfilter(y, A, mu0, Sigma0, Phi, sQ, sR, Ups = NULL, Gam = NULL,
        input = NULL, S = NULL, version = 1)
```

Arguments

y	data matrix (n × q), vector or time series, n = number of observations. Use NA or zero (0) for missing data.
A	can be constant or an array with dimension dim=c(q,p,n) if time varying (see details). Use NA or zero (0) for missing data.
mu0	initial state mean vector (p × 1)
Sigma0	initial state covariance matrix (p × p)
Phi	state transition matrix (p × p)
sQ	state error pre-matrix (see details)
sR	observation error pre-matrix (see details)
Ups	state input matrix (p × r); leave as NULL (default) if not needed
Gam	observation input matrix (q × r); leave as NULL (default) if not needed
input	NULL (default) if not needed or a matrix (n × r) of inputs having the same row dimension (n) as y
S	covariance matrix between the (not premultiplied) state and observation errors; not necessary to specify if not needed and only used if version=2. See details for more information.
version	either 1 (default) or 2; version 2 allows for correlated errors

Details

This script replaces Kfilter0, Kfilter1, and Kfilter2 by combining all cases. The major difference is how to specify the covariance matrices; in particular, sQ = t(cQ) and sR = t(cR) where cQ and cR were used in Kfilter0-1-2 scripts.

The states x_t are p-dimensional, the data y_t are q-dimensional, and the inputs u_t are r-dimensional for $t = 1, \dots, n$. The initial state is $x_0 \sim N(\mu_0, \Sigma_0)$.

The measurement matrices A_t can be constant or time varying. If time varying, they should be entered as an array of dimension dim = c(q,p,n). Otherwise, just enter the constant value making sure it has the appropriate $q \times p$ dimension.

Version 1 (default): The general model is

$$x_t = \Phi x_{t-1} + \Upsilon u_t + sQ w_t \quad w_t \sim iid N(0, I)$$

$$y_t = A_t x_{t-1} + \Gamma u_t + sR v_t \quad v_t \sim iid N(0, I)$$

where $w_t \perp v_t$. Consequently the state noise covariance matrix is $Q = sQ sQ'$ and the observation noise covariance matrix is $R = sR sR'$ and sQ, sR do not have to be square as long as everything is conformable. Notice the specification of the state and observation covariances has changed from the original scripts.

NOTE: If it is easier to model in terms of Q and R , simply input the square root matrices $sQ = Q^{.5}$ and $sR = R^{.5}$.

Version 2 (correlated errors): The general model is

$$x_{t+1} = \Phi x_t + \Upsilon u_{t+1} + sQ w_t \quad w_t \sim iid N(0, I)$$

$$y_t = A_t x_{t-1} + \Gamma u_t + sR v_t \quad v_t \sim iid N(0, I)$$

where $S = \text{Cov}(w_t, v_t)$, and NOT $\text{Cov}(sQ w_t, sR v_t)$.

NOTE: If it is easier to model in terms of Q and R , simply input the square root matrices $sQ = Q^{.5}$ and $sR = R^{.5}$.

Note that in either version, $sQ w_t$ has to be p-dimensional, but w_t does not, and $sR v_t$ has to be q-dimensional, but v_t does not.

Value

Time varying values are returned as arrays.

Xp	one-step-ahead prediction of the state
Pp	mean square prediction error
Xf	filter value of the state
Pf	mean square filter error
like	the negative of the log likelihood
innov	innovation series
sig	innovation covariances
Kn	last value of the gain, needed for smoothing

Note

If it is easier to model in terms of Q and R , simply input the square root matrices $sQ = Q^{.5}$ and $sR = R^{.5}$.

Author(s)

D.S. Stoffer

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

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The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[Ksmooth](#)

Examples

```
# generate some data
set.seed(1)
sQ = 1; sR = 3; n = 100
mu0 = 0; Sigma0 = 10; x0 = rnorm(1,mu0,Sigma0)
w = rnorm(n); v = rnorm(n)
x = c(x0 + sQ*w[1]); y = c(x[1] + sR*v[1]) # initialize
for (t in 2:n){
  x[t] = x[t-1] + sQ*w[t]
  y[t] = x[t] + sR*v[t]
}
# run and plot the filter
run = Kfilter(y, A=1, mu0, Sigma0, Phi=1, sQ, sR)

tsplot(cbind(x, y, Xf=run$Xf), spaghetti=TRUE, type='o', col=c(3,4,6), pch=c(NA,1,NA),
       addLegend=TRUE, location='topleft', lwd=c(2,1,2), margins=1)

# NOTE: CRAN tests need extra white space :( so margins=1 not necessary nor desired
```

Ksmooth

Quick Kalman Smoother

Description

Returns the smoother values for various linear state space models. The predicted and filtered values and the likelihood at the given parameter values are also returned (via `Kfilter`).

Usage

```
Ksmooth(y, A, mu0, Sigma0, Phi, sQ, sR, Ups = NULL, Gam = NULL,
       input = NULL, S = NULL, version = 1)
```

Arguments

y	data matrix (n × q), vector or time series, n = number of observations. Use NA or zero (0) for missing data.
A	can be constant or an array with dimension dim=c(q,p,n) if time varying (see details). Use NA or zero (0) for missing data.
mu0	initial state mean vector (p × 1)
Sigma0	initial state covariance matrix (p × p)
Phi	state transition matrix (p × p)
sQ	state error pre-matrix (see details)
sR	observation error pre-matrix (see details)
Ups	state input matrix (p × r); leave as NULL (default) if not needed
Gam	observation input matrix (q × r); leave as NULL (default) if not needed
input	NULL (default) if not needed or a matrix (n × r) of inputs having the same row dimension (n) as y
S	covariance matrix between state and observation errors; not necessary to specify if not needed and only used if version=2; see details
version	either 1 (default) or 2; version 2 allows for correlated errors

Details

The states x_t are p-dimensional, the data y_t are q-dimensional, and the inputs u_t are r-dimensional for $t = 1, \dots, n$. The initial state is $x_0 \sim N(\mu_0, \Sigma_0)$.

The measurement matrices A_t can be constant or time varying. If time varying, they should be entered as an array of dimension dim = c(q,p,n). Otherwise, just enter the constant value making sure it has the appropriate $q \times p$ dimension.

Version 1 (default): The general model is

$$\begin{aligned} x_t &= \Phi x_{t-1} + \Upsilon u_t + sQ w_t \quad w_t \sim iid N(0, I) \\ y_t &= A_t x_{t-1} + \Gamma u_t + sR v_t \quad v_t \sim iid N(0, I) \end{aligned}$$

where $w_t \perp v_t$. Consequently the state noise covariance matrix is $Q = sQ sQ'$ and the observation noise covariance matrix is $R = sR sR'$ and sQ, sR do not have to be square as long as everything is conformable. Notice the specification of the state and observation covariances has changed from the original scripts.

NOTE: If it is easier to model in terms of Q and R , simply input the square root matrices $sQ = Q^{.5}$ and $sR = R^{.5}$.

Version 2 (correlated errors): The general model is

$$\begin{aligned} x_{t+1} &= \Phi x_t + \Upsilon u_{t+1} + sQ w_t \quad w_t \sim iid N(0, I) \\ y_t &= A_t x_{t-1} + \Gamma u_t + sR v_t \quad v_t \sim iid N(0, I) \end{aligned}$$

where $S = \text{Cov}(w_t, v_t)$, and NOT $\text{Cov}(sQ w_t, sR v_t)$.

NOTE: If it is easier to model in terms of Q and R , simply input the square root matrices $sQ = Q^{.5}$ and $sR = R^{.5}$.

Note that in either version, $sQ w_t$ has to be p-dimensional, but w_t does not, and $sR v_t$ has to be q-dimensional, but v_t does not.

Value

Time varying values are returned as arrays.

Xs	state smoothers
Ps	smoother mean square error
X0n	initial mean smoother
P0n	initial smoother covariance
J0	initial value of the J matrix
J	the J matrices
Xp	state predictors
Pp	mean square prediction error
Xf	state filters
Pf	mean square filter error
like	negative of the log likelihood
innov	innovation series
sig	innovation covariances
Kn	the value of the last Gain

Author(s)

D.S. Stoffer

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

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The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[Kfilter](#)

Examples

```
# generate some data
set.seed(1)
sQ = 1; sR = 3; n = 100
mu0 = 0; Sigma0 = 10; x0 = rnorm(1,mu0,Sigma0)
w = rnorm(n); v = rnorm(n)
x = c(x0 + sQ*w[1]); y = c(x[1] + sR*v[1]) # initialize
for (t in 2:n){
  x[t] = x[t-1] + sQ*w[t]
```

```

    y[t] = x[t] + sR*v[t]
  }

# run and plot the smoother
run = Ksmooth(y, A=1, mu0, Sigma0, Phi=1, sQ, sR)

tsplot(cbind(x, y, Xs=run$Xs), spaghetti=TRUE, type='o', col=c(3,4,6), pch=c(NA,1,NA),
       addLegend=TRUE, location='bottomright', lwd=c(2,1,2), gg=TRUE, margins=1)

# NOTE: CRAN tests need extra white space :( so margins=1 not necessary nor desired

```

lag1.plot

Lag Plot - one time series

Description

Produces a grid of scatterplots of a series versus lagged values of the series.

Usage

```

lag1.plot(series, max.lag = 1, corr = TRUE, smooth = TRUE, col = gray(.1),
          bg = NA, lwl = 1, lwc = 2, bgl = NULL, ltcol = 1, box.col = NULL,
          cex = .9, gg = FALSE, location="topright", ...)

```

Arguments

series	the data
max.lag	maximum lag
corr	if TRUE, shows the autocorrelation value in a legend
smooth	if TRUE, adds a lowess fit to each scatterplot
col	color of points; default is gray(.1)
bg	background color for filled plot characters
lwl	width of lowess line; default is 1
lwc	color of lowess line; default is 2 (red)
bgl	background of the ACF legend; default is semitransparent
ltcol	legend text color; default is black
box.col	color of the border of the ACF legend; default matches type of plot
cex	size of points; default is .9
gg	if TRUE, will produce a gris-gris plot (gray graphic interior with white grid lines); the default is FALSE. The grammar of astsa is voodoo
location	the location of the ACF legend with options "bottomright", "bottom", "bottomleft", "left", "topleft", "top", "topright" (the default), "right" and "center".
...	additional graphical arguments

Author(s)

D.S. Stoffer

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[lag2.plot](#)

Examples

```
lag1.plot(log(varve), max.lag=9, lwl=3, lwc=6, col=5, location='topleft')
```

```
lag1.plot(soi, 12, cex=1, pch=19, col=astsa.col(4, .3), gg=TRUE)
```

lag2.plot

Lag Plot - two time series

Description

Produces a grid of scatterplots of one series versus another lagged. The first named series is the one that gets lagged.

Usage

```
lag2.plot(series1, series2, max.lag = 0, corr = TRUE, smooth = TRUE, col = gray(.1),
          bg = NA, lwl = 1, lwc = 2, bgl = NULL, ltcol = 1, box.col = NULL, cex = .9,
          gg = FALSE, location="topright", ...)
```

Arguments

series1	first series (the one that gets lagged)
series2	second series
max.lag	maximum number of lags
corr	if TRUE, shows the cross-correlation value in a legend
smooth	if TRUE, adds a lowess fit to each scatterplot
col	color of points; default is gray(.1)
bg	background color for filled plot characters

lwl	width of lowess line; default is 1
lwc	color of lowess line; default is 2 (red)
bg1	background of the ACF legend; default is semitransparent
ltcol	legend text color; default is black
box.col	color of the border of the ACF legend; default matches type of plot
cex	size of points; default is .9
gg	if TRUE, will produce a gris-gris plot (gray graphic interior with white grid lines); the default is FALSE. The grammar of astsa is voodoo
location	the location of the CCF legend with options "bottomright", "bottom", "bottomleft", "left", "topleft", "top", "topright" (the default), "right" and "center".
...	additional graphical parameters

Author(s)

D.S. Stoffer

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[lag1.plot](#)

Examples

```
lag2.plot(Hare, Lynx, max.lag=5, lwl=2, lwc=3, cex=1.5, pch=24, bg='orange')
lag2.plot(soi, rec, 8, cex=1.1, pch=19, col=5, lwl=2, location="bottomleft")
lag2.plot(diff(lead), diff(sales), 5, corr=FALSE, gg=TRUE, pch=8, col=6, lwc=4)
```

LagReg

*Lagged Regression***Description**

Performs lagged regression as discussed in Chapter 4.

Usage

```
LagReg(input, output, L = c(3, 3), M = 40, threshold = 0,
       inverse = FALSE)
```

Arguments

input	input series
output	output series
L	degree of smoothing; see spans in the help file for spec.pgram.
M	must be even; number of terms used in the lagged regression
threshold	the cut-off used to set small (in absolute value) regression coefficients equal to zero
inverse	if TRUE, will fit a forward-lagged regression

Details

For a bivariate series, input is the input series and output is the output series. The degree of smoothing for the spectral estimate is given by L; see spans in the help file for spec.pgram. The number of terms used in the lagged regression approximation is given by M, which must be even. The threshold value is the cut-off used to set small (in absolute value) regression coefficients equal to zero (it is easiest to run LagReg twice, once with the default threshold of zero, and then again after inspecting the resulting coefficients and the corresponding values of the CCF). Setting inverse=TRUE will fit a forward-lagged regression; the default is to run a backward-lagged regression. The script is based on code that was contributed by Professor Doug Wiens, Department of Mathematical and Statistical Sciences, University of Alberta.

Value

Graphs of the estimated impulse response function, the CCF, and the output with the predicted values superimposed.

beta	Estimated coefficients
fit	The output series, the fitted values, and the residuals

Note

See Chapter 4 of the text for an example.

Author(s)

D.S. Stoffer

References

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In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

 lap

LA Pollution-Mortality Study

Description

LA Pollution-Mortality Study (1970-1979), weekly data.

Format

The format is: mts [1:508, 1:11]

Details

columns are time series	with names
(1) Total Mortality	tmort
(2) Respiratory Mortality	rmort
(3) Cardiovascular Mortality	cmort
(4) Temperature	tempr
(5) Relative Humidity	rh
(6) Carbon Monoxide	co
(7) Sulfur Dioxide	so2
(8) Nitrogen Dioxide	no2
(9) Hydrocarbons	hycarb
(10) Ozone	o3
(11) Particulates	part

Note

Details may be found in <http://www.sungpark.net/ShumwayAzariPawitan88.pdf>

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[lap.xts](#)

lap.xts

LA Pollution-Mortality Study: Sampled Daily

Description

Original data from a study of the effects of pollution and weather on mortality, LA, 1970-1979. These are 3652 daily observations for the 10 year period. The data set is an xts object indexed by Date.

Format

The format is: An xts object on 1970-01-01 / 1979-12-31 containing:

Data: double [3652, 11]

Columns: Tmort, Rmort, Cmort, Temp, Rhumid, CO, SO2, NO2, HC, Ozone, Part

Index: Date [3652] (TZ: "UTC")

Details

columns are time series	with names
(1) Total Mortality	Tmort
(2) Respiratory Mortality	Rmort
(3) Cardiovascular Mortality	Cmort
(4) Temperature	Temp
(5) Relative Humidity	Rhumid
(6) Carbon Monoxide	CO
(7) Sulfur Dioxide	SO2
(8) Nitrogen Dioxide	NO2
(9) Hydrocarbons	HC
(10) Ozone	Ozone
(11) Particulates	Part

Note

These are the original data from <https://github.com/DStoffer/dsstofer.github.io/blob/main/files/LAP.pdf>.

The weekly data in `lap` were taken from this data set last century. The details, however, were never entirely made clear and it's too late to get them now. It is easy to pull out the weekly averages from this data set, and how to do so is given in the Examples section below; the resulting data set will be slightly different than `lap`. The names for this data set are different from `lap`, the main difference is these names have capitals.

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

`lap`

Examples

```
## Not run:

library(xts) # assumes package has been installed

plot(lap.xts$Cmort, col=4)

lapw = apply.weekly(lap.xts, FUN=colMeans) # get weekly averages

plot(lapw[,c('Cmort', 'Temp', 'Part')], col=astsa.col(2:4, .7), main=NA)
addLegend(col=2:4, lty=1, lwd=2, ncol=3, bty="white")

sarima(lapw$Cmort, 0,1,1, no.constant=TRUE) # fit ARIMA(0,1,1) to weekly Cmort

## End(Not run)
```

lead

Leading Indicator

Description

Leading indicator, 150 months; taken from Box and Jenkins (1970).

Usage

```
data(lead)
```

Format

The format is: Time-Series [1:150] from 1 to 150: 10.01 10.07 10.32 9.75 10.33 ...

Details

This is also the R time series `BJsales.lead`: The sales time series `BJsales` and leading indicator `BJsales.lead` each contain 150 observations. The objects are of class "ts".

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[sales](#)

Lynx

Canadian Lynx

Description

This is one of the classic studies of predator-prey interactions, the 90-year data set is the number, in thousands, of lynx pelts purchased by the Hudson's Bay Company of Canada. While this is an indirect measure of predation, the assumption is that there is a direct relationship between the number of pelts collected and the number of hare and lynx in the wild.

Format

The format is: Time-Series [1:91] from 1845 to 1935: 30.1 45.1 49.1 39.5 21.2 ...

Note

The data are in units of one thousand. This data set pairs with [Hare](#) and is NOT the same as [lynx](#).

Source

From Odum's "Fundamentals of Ecology", p. 191. Additional information at <http://people.whitman.edu/~hundredr/courses/M250F03/M250.html> scroll down to: Chapter 6, Difference Equations

NB: For some reason, there is not a secure link to the site above.

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[Hare](#)

matrixpwr

Powers of a Square Matrix

Description

matrixpwr computes powers of a square matrix including negative powers for nonsingular matrices.

%% is a more intuitive interface as an operator.

Usage

```
matrixpwr(A, power)
```

```
A %% power
```

Arguments

A	a square matrix
power	single numeric

Details

Raises matrix to the specified power. The matrix must be square and if power < 0, the matrix must be nonsingular.

Note that %% is defined as "`%%`" <- function(A, power) matrixpwr(A, power)

If power = 0, the identity matrix is returned.

Value

Returns matrix raised to the given power.

Author(s)

D.S. Stoffer

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

Examples

```
# 2-state Markov transition matrix to steady state
( P = matrix(c(.7,.4,.3,.6), 2) )
P %%% 50

# surround with parentheses if used in an expression
c(.2, .8) %*% (P %%% 50)

# Inverse square root
var(econ5) %^% -.5
```

MEI

Multivariate El Nino/Southern Oscillation Index (version 1)

Description

Bimonthly MEI values, starting with Dec1949/Jan1950 through Oct/Nov2019. All values are normalized for each bimonthly season so that the 44 values from 1950 to 1993 have an average of zero and a standard deviation of 1. Larger values correspond to warmer temperatures (unlike soi and ENSO).

Format

The format is: Time-Series [1:827] from 1950 to 2019: -1.03 -1.13 -1.28 -1.07 -1.43 ...

Details

For full details, see <https://psl.noaa.gov/enso/mei.old/mei.html>. Multivariate ENSO Index (MEI) is a combined score on the six main observed variables over the tropical Pacific. These six variables are: sea-level pressure (P), zonal (U) and meridional (V) components of the surface wind, sea surface temperature (S), surface air temperature (A), and total cloudiness fraction of the sky (C). These observations have been collected and published in ICOADS for many years. The MEI is computed separately for each of twelve sliding bi-monthly seasons (Dec/Jan, Jan/Feb,..., Nov/Dec). After spatially filtering the individual fields into clusters, the MEI is calculated as the first unrotated Principal Component (PC) of all six observed fields combined. This is accomplished by normalizing the total variance of each field first, and then performing the extraction of the first PC on the covariance matrix of the combined fields. In order to keep the MEI comparable, all seasonal values are standardized with respect to each season and to the 1950-93 reference period.

Source

<https://psl.noaa.gov/enso/mei.old/table.html>

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).
The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.
In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.
The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[soi](#), [ENSO](#)

Months	<i>Month Labels</i>
--------	---------------------

Description

Provides labels for the (English) months of the year to be used in plotting monthly time series.

Format

The format is: chr [1:12] "J" "F" "M" "A" "M" "J" "J" "A" "S" "O" "N" "D"

Note

Hi Kids. The months of the year in English are:
January, February, March, April, May, June, July, August, September, October, November, December.

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

Examples

```
sAR = sarima.sim(sar=.9, S=12, n=36)
tsplot(sAR, type='c')
points(sAR, pch=Months, cex=1.1, font=4, col=1:4)
```

mvspec

Univariate and Multivariate Spectral Estimation

Description

This is spec.pgram with a number changes and written so you can easily extract the estimate of the multivariate spectral matrix as fxx. The bandwidth calculation has been changed to the more practical definition given in the text and this can be used to replace spec.pgram.

Usage

```
mvspec(x, spans = NULL, kernel = NULL, taper = 0, pad = 0, fast = TRUE,
       demean = FALSE, detrend = TRUE, lowess = FALSE, log = 'n', plot = TRUE,
       gg = FALSE, type = NULL, na.action = na.fail, nxm = 2, nym = 1,
       main = NULL, xlab = NULL, cex.main = NULL, ci = .95, ci.col = 4, plot.type, ...)
```

Arguments

x	univariate or multivariate time series (i.e., the p columns of x are time series)
spans	vector of odd integers giving the widths of modified Daniell smoothers to be used to smooth the periodogram
kernel	alternatively, a kernel smoother of class tskernel
taper	specifies the proportion of data to taper using a split cosine bell taper (.5 specifies a full taper)
pad	proportion of data to pad (zeros are added to the end of the series to increase its length by the proportion pad)
fast	logical; if TRUE, pad the series to a highly composite length
demean	if TRUE, series is demeaned first
detrend	if TRUE, series is detrended first (unless demean is TRUE)

lowess	if TRUE and detrend TRUE (and demean FALSE), series is detrended using lowess first
log	if log='y', spectra plotted on a log scale; otherwise a log scale is not used
plot	plot the estimated spectra
gg	if TRUE, will produce a gris-gris plot (gray graphic interior with white grid lines); the default is FALSE. The grammar of astsa is voodoo
type	type of plot to be drawn, defaults to lines (see par)
na.action	how to handle missing values
nxm, nym	the number of minor tick mark divisions on x-axis, y-axis; the default is one minor tick on the x-axis and none on the y-axis
main	title of the graphics; if NULL (default), a totally awesome title is generated dude, but if NA there will be no gnarly title and the top margin will be used for the plot
xlab	label for frequency axis; if NULL (default), a totally awesome label is generated for your viewing pleasure
cex.main	magnification for main title; default is 1.
ci	confidence level if one is drawn.
ci.col	color of the confidence interval if one is drawn.
plot.type	plot type for multivariate time series; leave blank or specify 'marginal' if a marginal plot of the spectra is desired, or set plot.type='coh' for coherency plot(s), or plot.type='phase' for phase plot(s). .
...	additional graphical arguments.

Details

This is built off of `spec.pgram` from the `stats` package with a few changes in the defaults and written so you can easily extract the estimate of the multivariate spectral matrix as `fxx`.

The default for the plot is NOT to plot on a log scale and the graphic will have a grid. Overall, the graphics have been improved.

The bandwidth calculation has been changed to the more practical definition given in the text, $(L_h/n.used) * frequency(x)$. Also, the bandwidth is not displayed in the graphic, but is returned.

Although initially meant to be used to easily obtain multivariate (mv) spectral (spec) estimates, this script can be used for univariate time series as a replacement for `spec.pgram`.

Note that the script does not taper by default (`taper=0`); this forces the user to do "conscious tapering".

In the multivariate case (more than 2 series), if "coherency" or "phase" plots are desired, the result is a grid of plots. There is a "scale" factor that can be set to prevent the labels from getting too small if there are many series. The default is `scale=1`, and to expand the labels by 10% for example, set `scale=1.1`. Also, if there are many series, having minor tick marks can make the graphic look crowded; in this case, set `minor=FALSE`. Finally, the plots include a type of legend that shows the axes with their corresponding labels (by default). To turn the legend off, include `addLegend=FALSE` in the call. Again, these considerations only work in the multivariate case (more than 2 series).

Value

All results are returned invisibly.

If plot is TRUE and smoothing is used, the bandwidth, degrees of freedom, and taper amount are printed.

An object of class "spec", which is a list containing at least the following components:

fxx	spectral matrix estimates; an array of dimensions $\text{dim} = c(p, p, \text{nfreq})$.
freq	vector of frequencies at which the spectral density is estimated.
spec	vector (for univariate series) or matrix (for multivariate series) of estimates of the spectral density at frequencies corresponding to freq.
details	matrix with columns: frequency, period, spectral ordinate(s)
coh	NULL for univariate series. For multivariate time series, a matrix containing the squared coherency between different series. Column $i + (j - 1) * (j - 2)/2$ of coh contains the squared coherency between columns i and j of x , where $i < j$.
phase	NULL for univariate series. For multivariate time series a matrix containing the cross-spectrum phase between different series. The format is the same as coh.
Lh	Number of frequencies (approximate) used in the band.
n.used	Sample length used for the FFT
df	Degrees of freedom (may be approximate) associated with the spectral estimate.
bandwidth	Bandwidth (may be approximate) associated with the spectral estimate.
method	The method used to calculate the spectrum.

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

Examples

```
# real raw periodogram
mvspec(soi)
mvspec(soi, log='y') # on a log scale

# smooth and some details printed
mvspec(soi, spans=c(7,7), taper=.1)$details[1:45,]

# bivariate example
deth = cbind(mdeaths, fdeaths) # two R data sets, male/female monthly deaths ...
tsplot(deth, type='b', col=c(4,6), spaghetti=TRUE, pch=c('M','F'), addLegend=TRUE)
dog = mvspec(deth, spans=c(3,3), taper=.1)
dog$fxx[,1:5] # look at a few spectral matrix estimates
```

```

dog$bandwidth # bandwidth with time unit = year
dog$df        # degrees of freedom
mvspec(deth, spans=c(3,3), taper=.1, plot.type='coh') # coherence

# multivariate example
mvspec(diff(log(econ5)), spans=c(5,5), col=5, lwd=2, ci=NA, gg=TRUE, minor=FALSE,
        plot.type='coh')

mvspec(diff(log(econ5)), spans=c(5,5), col=5, lwd=2, ci=NA, gg=TRUE, minor=FALSE,
        plot.type='coh', addLegend=FALSE)

```

nyse

Returns of the New York Stock Exchange

Description

Returns of the New York Stock Exchange (NYSE) from February 2, 1984 to December 31, 1991.

Format

The format is: Time-Series [1:2000] from 1 to 2000: 0.00335 -0.01418 -0.01673 0.00229 -0.01692
...

Note

Various packages have data sets called nyse. Consequently, it may be best to specify this data set as `nyse = astsa::nyse` to avoid conflicts.

Source

S+GARCH module - Version 1.1 Release 2: 1998

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

oil*Crude oil, WTI spot price FOB*

Description

Crude oil, WTI spot price FOB (in dollars per barrel), weekly data from 2000 to mid-2010.

Format

The format is: Time-Series [1:545] from 2000 to 2010: 26.2 26.1 26.3 24.9 26.3 ...

Details

pairs with the series gas

Source

Data were obtained from the URL: www.eia.doe.gov/dnav/pet/pet_pri_spt_s1_w.htm

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[gas](#)

part*Particulate levels from the LA pollution study*

Description

Particulate series corresponding to cmor t from the LA pollution study.

Format

The format is: Time-Series [1:508] from 1970 to 1980: 72.7 49.6 55.7 55.2 66 ...

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[lap](#), [lap.xts](#)

PLT

Platelet Levels

Description

PLT: Measurements made for 91 days on the three variables, log(white blood count) [WBC], log(platelet) [PLT] and hematocrit [HCT]. Missing data code is 0 (zero).

Usage

```
data(PLT)
```

Format

The format is: Time-Series [1:91] from 1 to 91: 4.47 4.33 4.09 4.6 4.41 ...

Details

See Examples 6.1 and 6.9 for more details.

Source

Jones, R.H. (1984). Fitting multivariate models to unequally spaced data. In *Time Series Analysis of Irregularly Observed Data*, pp. 158-188. E. Parzen, ed. Lecture Notes in Statistics, 25, New York: Springer-Verlag.

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also[blood](#), [HCT](#), [WBC](#)

polio

*Poliomyelitis cases in US***Description**

Monthly time series of poliomyelitis cases reported to the U.S. Centers for Disease Control for the years 1970 to 1983, 168 observations.

Format

The format is: Time-Series [1:168] from 1970 to 1984: 0 1 0 0 1 3 9 2 3 5 ...

Details

The data were originally modelled by Zeger (1988) “A Regression Model for Time Series of Counts,” *Biometrika*, 75, 822-835.

Source

Data taken from the `gamlss.data` package; see <https://www.gamlss.com/>.

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

Examples

```
tsplot(polio, type='s')
```

`polyMul`*Multiplication of Two Polynomials*

Description

Multiplication of two polynomials.

Usage

```
polyMul(p, q)
```

Arguments

p	coefficients of first polynomial
q	coefficients of second polynomial

Details

inputs are vectors of coefficients a, b, c, ..., in order of power $ax^0 + bx^1 + cx^2 + \dots$

Value

coefficients of the product in order of power

Author(s)

D.S. Stoffer

Source

based on code from the polynom package <https://CRAN.R-project.org/package=polynom>

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

Examples

```
a = 1:3 # 1 + 2x + 3x^2
b = 1:2 # 1 + 2x
polyMul(a, b)
# [1] 1 4 7 6
# 1 + 4x + 7x^2 + 6x^3
```

pre.white

*Cross-Correlation Analysis With Automatic Prewhitening***Description**

Performs a cross-correlation analysis on two series after prewhitening the first series and filtering the second series accordingly.

Usage

```
pre.white(series1, series2, diff = FALSE, max.lag = NULL, main = NULL,
          order.max = NULL, plot = TRUE, ...)
```

Arguments

series1, series2	univariate time series
diff	(logical or integer) should the series be differenced prior to the analysis and if more than first order, by how much
max.lag	maximum lag for which to plot the CCF - if NULL, a suitable number is chosen (see details)
main	plot title - if NULL, uses series1 name appended by .w for whitened and series2 name appended by .f for filtered
order.max	maximum order of model to fit (see details)
plot	should the sample CCF be plotted
...	additional graphic arguments

Details

The first series is prewhitened by fitting a long AR based on AIC and the second series is filtered appropriately. Then a cross-correlation analysis is performed via [ccf2](#). If differencing is specified, both series are differenced the same way prior to the prewhitening. The resulting series are returned invisibly.

The default is no differencing. Differences of order 1 can be set by entering `diff = TRUE` or `diff = 1`. If it is necessary to use higher orders, then enter a positive integer (this is rare).

The maximum lag (`max.lag`) in the CCF graphic defaults (if NULL) to the smaller of 50 and 20% of the sample size.

The maximum order (`order.max`) for fitting the AR via AIC defaults (if NULL) to the minimum of 30 and 15% of the number of observations.

Value

Returns the sample CCF graphic using the prewhitened series unless `plot = FALSE`. The prewhitened series are returned invisibly.

Author(s)

D.S. Stoffer

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

Examples

```
pre.white(cmort, part, diff=TRUE, col=4)
```

prodn

Monthly Federal Reserve Board Production Index

Description

Monthly Federal Reserve Board Production Index (1948-1978, n = 372 months).

Usage

```
data(prodn)
```

Format

The format is: Time-Series [1:372] from 1948 to 1979: 40.6 41.1 40.5 40.1 40.4 41.2 39.3 41.6 42.3 43.2 ...

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

qinfl	<i>Quarterly Inflation</i>
-------	----------------------------

Description

Quarterly inflation rate in the Consumer Price Index from 1953-I to 1980-II, n = 110 observations.

Format

The format is: Time-Series [1:110] from 1953 to 1980: 1.673 3.173 0.492 -0.327 -0.333 ...

Details

pairs with qintr (interest rate)

Source

Newbold, P. and T. Bos (1985). *Stochastic Parameter Regression Models*. Beverly Hills: Sage.

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[qintr](#)

qintr	<i>Quarterly Interest Rate</i>
-------	--------------------------------

Description

Quarterly interest rate recorded for Treasury bills from 1953-I to 1980-II, n = 110 observations.

Format

The format is: Time-Series [1:110] from 1953 to 1980: 1.98 2.15 1.96 1.47 1.06 ...

Details

pairs with qinfl (inflation)

Source

Newbold, P. and T. Bos (1985). *Stochastic Parameter Regression Models*. Beverly Hills: Sage.

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[qinfl](#)

QQnorm

Normal Quantile-Quantile Plot

Description

Produces a normal QQ plot with a line of equality and a confidence band (by default) of the input data. This is basically a prettier version of qqnorm from the stats package.

Usage

```
QQnorm(xdata, col = c(4, 6), ylab = "Sample Quantiles", xlab = "Theoretical Quantiles",
       main = "Normal Q-Q Plot", ylim = NULL, ci = TRUE, qqlwd = 1, ...)
```

Arguments

xdata	the data. If a matrix, the data are collapsed.
col	vector of 2, first is point color, second is line color (default is blue-4 and magenta-6).
ylab	y-axis label (default is 'Sample Quantiles').
xlab	x-axis label (default is 'Theoretical Quantiles').
main	plot title (default is 'Normal Q-Q Plot')
ylim	limits on y-axis (default is the most beautiful limits ever).
ci	if TRUE (default) draws pointwise 99.99% CIs as a band. If FALSE or 0, no CI is drawn. Alternately, enter a percentage (e.g., either ci=95 or ci=.95 will work).
qqlwd	line width of the qqline (default is 1).
...	other graphical parameters sent to tsplot .

Details

If you want a graphic to check normality of your data in `xdata`, just enter `QQnorm(xdata)` and sit back and enjoy the beauty of this script (you may want to wear sunglasses).

For confidence levels, various values are allowed. For example, 95% limits can be obtained as `ci=95` or `ci=.95`, both of which are conventional. However, `ci=5`, or `ci=.05` will also work for 95% intervals (so you can not go below 50%). If you ask for a confidence level of 100% or larger, you will get the default without a warning and maybe you are unconventional.

Author(s)

D.S. Stoffer

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

Examples

```
QQnorm(log(varve))
```

rec

Recruitment (number of new fish index)

Description

Recruitment (index of the number of new fish) for a period of 453 months ranging over the years 1950-1987. Recruitment is loosely defined as an indicator of new members of a population to the first life stage at which natural mortality stabilizes near adult levels.

Usage

```
data(rec)
```

Format

The format is: Time-Series [1:453] from 1950 to 1988: 68.6 68.6 68.6 68.6 68.6 ...

Details

can pair with `soi` (Southern Oscillation Index)

Source

Data furnished by Dr. Roy Mendelssohn of the Pacific Fisheries Environmental Laboratory, NOAA (personal communication). Further discussion of the concept of Recruitment may be found here: derekogle.com/fishR/examples/oldFishRVignettes/StockRecruit.pdf

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

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See Also

[soi](#)

sales

Sales

Description

Sales, 150 months; taken from Box and Jenkins (1970).

Format

The format is: Time-Series [1:150] from 1 to 150: 200 200 199 199 199 ...

Details

This is also the R data set `BJsales`: The sales time series `BJsales` and leading indicator `BJsales.lead` each contain 150 observations. The objects are of class "ts".

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[lead](#)

salmon	<i>Monthly export price of salmon</i>
--------	---------------------------------------

Description

Farm Bred Norwegian Salmon, export price, US Dollars per Kilogram

Format

The format is: Time-Series [1:166] from September 2003 to June 2017: 2.88 3.16 2.96 3.12 3.23 3.32 3.45 3.61 3.48 3.21 ...

Source

<https://www.indexmundi.com/commodities/>

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

salt	<i>Salt Profiles</i>
------	----------------------

Description

Salt profiles taken over a spatial grid set out on an agricultural field, 64 rows at 17-ft spacing.

Usage

```
data(salt)
```

Format

The format is: Time-Series [1:64] from 1 to 64: 6 6 6 3 3 3 4 4 4 1.5 ...

Details

pairs with `saltemp`, temperature profiles on the same grid

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[saltemp](#)

saltemp

Temperature Profiles

Description

Temperature profiles over a spatial grid set out on an agricultural field, 64 rows at 17-ft spacing.

Usage

```
data(saltemp)
```

Format

The format is: Time-Series [1:64] from 1 to 64: 5.98 6.54 6.78 6.34 6.96 6.51 6.72 7.44 7.74 6.85
...

Details

pairs with `salt`, salt profiles on the same grid

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[salt](#)

 sarima

Fit ARIMA Models

Description

Fits ARIMA models (with diagnostics) in a short command. It can also be used to perform regression with autocorrelated errors.

Usage

```
sarima(xdata, p, d, q, P = 0, D = 0, Q = 0, S = -1,
       details = TRUE, xreg = NULL, Model = TRUE,
       fixed = NULL, tol = sqrt(.Machine$double.eps),
       no.constant = FALSE, col, ...)
```

Arguments

xdata	univariate time series
p	AR order
d	difference order
q	MA order
P	SAR order; use only for seasonal models
D	seasonal difference; use only for seasonal models
Q	SMA order; use only for seasonal models
S	seasonal period; use only for seasonal models
details	if FALSE, turns off the diagnostic plot and the output from the nonlinear optimization routine, which is <code>optim</code> . The default is TRUE.
xreg	Optionally, a vector or matrix of external regressors, which must have the same number of rows as xdata.
Model	if TRUE (default), the model orders are printed on the diagnostic plot.
fixed	optional numeric vector of the same length as the total number of parameters. If supplied, only parameters corresponding to NA entries will be estimated.
tol	controls the relative tolerance (<code>reltol</code> in <code>optim</code>) used to assess convergence. The default is <code>sqrt(.Machine\$double.eps)</code> , the R default.
no.constant	controls whether or not <code>sarima</code> includes a constant in the model. In particular, if there is no differencing (<code>d = 0</code> and <code>D = 0</code>) you get the mean estimate. If there is differencing of order one (either <code>d = 1</code> or <code>D = 1</code> , but not both), a constant term is included in the model. These two conditions may be overridden (i.e., no constant will be included in the model) by setting this to TRUE; e.g., <code>sarima(x, 1, 1, 0, no.constant=TRUE)</code> . Otherwise, no constant or mean term is included in the model. If regressors are included (via <code>xreg</code>), this is ignored.
col	color of diagnostic plots; default is 1 (black)
...	additional graphical arguments

Details

If your time series is in `x` and you want to fit an $\text{ARIMA}(p,d,q)$ model to the data, the basic call is `sarima(x,p,d,q)`. As of version 2.3, the orders do not have to be specified if they are zero. For example, `sarima(x, p=1)` is the same as `sarima(x, 1, 0, 0)`.

To fit a seasonal ARIMA model, the basic call is `sarima(x,p,d,q,P,D,Q,S)`. For example, `sarima(x, 2, 1, 0, 0, 1, 1, 12)` will fit a seasonal $\text{ARIMA}(2, 1, 0) \times (0, 1, 1)_{12}$ model to the series in `x`. The orders do not have to be specified if they are zero; e.g., `sarima(x, d=1, q=1, D=1, Q=1, S=4)` works.

The results are the parameter estimates, standard errors, AIC, AICc, BIC and diagnostics. The difference between the information criteria given by `sarima()` and `arima()` is that they differ by a scaling factor of the effective sample size.

Value

A t-table, the estimated noise variance, and AIC, AICc, BIC are printed. The following are returned invisibly as a list:

<code>fit</code>	[[1]] an object of class <code>Arima</code> with more information than you need
<code>sigma2</code>	[[2]] the estimate of the noise variance
<code>degrees_of_freedom</code>	[[3]] error degrees of freedom
<code>t.table</code>	[[4]] a little t-table with two-sided p-values
<code>ICs</code>	[[5]] AIC - AICc - BIC

Missing Data

Yes it's ok if input as NA and the observations are vector or ts objects (meaning equally spaced).

Source

This is an enhancement of `arima` from the `stats` package.

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[sarima.for](#), [sarima.sim](#)

Examples

```
# easy to use
sarima(rec, 2,0,0) # data, p, d, and q
# redux - minimal output
sarima(rec, p=2, details=FALSE)

# fun for the whole family
dog = sarima(log(AirPassengers), 0,1,1, 0,1,1,12, details=FALSE)
# dog[[1]] has most of the results ...
tsplot(resid(dog[[1]]))

# fixed parameters
x = sarima.sim( ar=c(0,-.9), n=200 ) + 50
sarima(x, p=2, fixed=c(0,NA,NA)) # phi1 fixed, phi2 and mean free

# regression with autocorrelated errors
sarima(log(cpg), p=1, xreg=time(cpg))

# missing data (color, gris-gris, and pch, added for fun)
sarima(ar1miss, p=1, col=4, gg=TRUE, pch=19)
```

sarima.for

ARIMA Forecasting

Description

ARIMA forecasting.

Usage

```
sarima.for(xdata, n.ahead, p, d, q, P=0, D=0, Q=0, S=-1, tol = sqrt(.Machine$double.eps),
  no.constant = FALSE, plot = TRUE, plot.all = FALSE, ylab = NULL, xreg = NULL,
  newxreg = NULL, fixed = NULL, pcol = 2, pch = 1, ...)
```

Arguments

xdata	univariate time series
n.ahead	forecast horizon (number of periods)
p	AR order
d	difference order
q	MA order
P	SAR order; use only for seasonal models
D	seasonal difference; use only for seasonal models
Q	SMA order; use only for seasonal models

<code>S</code>	seasonal period; use only for seasonal models
<code>tol</code>	controls the relative tolerance (<code>reltol</code>) used to assess convergence. The default is <code>sqrt(.Machine\$double.eps)</code> , the R default.
<code>no.constant</code>	controls whether or not a constant is included in the model. If <code>no.constant=TRUE</code> , no constant is included in the model. See sarima for more details.
<code>plot</code>	if <code>TRUE</code> (default) the data (or some of it) and the forecasts and bounds are plotted
<code>plot.all</code>	if <code>TRUE</code> , all the data are plotted in the graphic; otherwise, only the last 100 observations are plotted in the graphic.
<code>ylab</code>	if <code>NULL</code> (default), the y-axis label is the name of the series.
<code>xreg</code>	Optionally, a vector or matrix of external regressors, which must have the same number of rows as the series. If this is used, <code>newxreg</code> MUST be specified.
<code>newxreg</code>	New values of <code>xreg</code> to be used for prediction. Must have at least <code>n.ahead</code> rows.
<code>fixed</code>	optional numeric vector of the same length as the total number of parameters. If supplied, only parameters corresponding to NA entries will be estimated.
<code>pcol</code>	color of the predictions in the graphic.
<code>pch</code>	plot character for the graphic. If <code>plot.all=TRUE</code> , this only applies to the forecasts.
<code>...</code>	additional graphical arguments

Details

For example, `sarima.for(x, 5, 1, 0, 1)` or `sarima.for(x, 5, p=1, q=1)` will forecast five time points ahead for an ARMA(1,1) fit to `x`. The output prints the forecasts and the standard errors of the forecasts, and supplies a graphic of the forecast with +/- 1 and 2 prediction error bounds.

Value

<code>pred</code>	the forecasts
<code>se</code>	the prediction (standard) errors

Missing Data

Yes it's ok if input as NA and the observations are vector or `ts` objects (meaning equally spaced). In this case, the graphic includes a line with points. Otherwise, lone observations would not be visible.

Note

If `plot.all=TRUE`, the data are displayed as a line only unless there are missing observations; see the Missing Data section. Points (and more) can be added to the graphic as long as the device stays open. For example:

```
sarima.for(gtemp_land, 10, d=1, q=1, plot.all=TRUE, pch=19)
points(gtemp_land, pch=20, col=4)
abline(v=2024, col=6, lty=5)
text(2000, 2.2, "it's getting hot in here", font=2, col=6, srt=45)
```


References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[sarima](#)

Examples

```
sarima.for(gtemp_both, n.ahead=6, d=1, q=1, col=6, pcol=4, gg=TRUE)

# with regressors
nummy = length(soi)
n.ahead = 24
nureg = time(soi)[nummy] + seq(1,n.ahead)/12
sarima.for(soi, n.ahead, 2,0,0, 2,0,0,12, xreg=time(soi), newxreg=nureg)

# missing data
sarima.for(ar1miss, n.ahead=5, p=1, pch=19)
```

sarima.sim

ARIMA Simulation

Description

Simulate data from (seasonal) ARIMA models.

Usage

```
sarima.sim(ar = NULL, d = 0, ma = NULL, sar = NULL, D = 0, sma = NULL, S = NULL,
           n = 500, rand.gen = rnorm, innov = NULL, burnin = NA, t0 = 0, ...)
```

Arguments

ar	coefficients of AR component (does not have to be specified)
d	order of regular difference (does not have to be specified)
ma	coefficients of MA component (does not have to be specified)
sar	coefficients of SAR component (does not have to be specified)
D	order of seasonal difference (does not have to be specified)
sma	coefficients of SMA component (does not have to be specified)

<code>S</code>	seasonal period (does not have to be specified)
<code>n</code>	desired sample size (defaults to 500)
<code>rand.gen</code>	optional; a function to generate the innovations (defaults to normal)
<code>innov</code>	an optional times series of innovations. If not provided, <code>rand.gen</code> is used.
<code>burnin</code>	length of burn-in (a non-negative integer). If NA, or not a positive integer, or a vector, a reasonable value is selected.
<code>t0</code>	start time (defaults to 0)
<code>...</code>	additional arguments applied to the innovations. For <code>rand.gen</code> , the standard deviation of the innovations generated by <code>rnorm</code> can be specified by <code>sd</code> or the mean by <code>mean</code> (see details and examples). In addition, <code>rand.gen</code> may be overridden using a preset sequence of innovations specifying <code>innov</code> (see details and examples).

Details

Will generate a time series of length `n` from the specified SARIMA model using simplified input.

The use of the term `mean` in `...` refers to the generation of normal innovations. For example, `sarima.sim(ar=.9, mean=5)` will generate data using $N(5,1)$ or $5+N(0,1)$ innovations, so that the constant in the model is 5 and the mean of the AR model is $5/(1-.9) = 50$. In `sarima.sim(ma=.9, mean=5)`, however, the model mean is 5 (the constant). Also, a random walk with drift = .1 can be generated by `sarima.sim(d=1, mean=.1, burnin=0)`, which is equivalent to `cumsum(rnorm(500, mean=.1))`. The same story goes if `sd` is specified; i.e., it's applied to the innovations. Because anything specified in `...` refers to the innovations, a simpler way to generate a non-zero mean is to add the value outside the call; see the examples.

If `innov` is used to input the innovations and override `rand.gen`, be sure that `length(innov)` is at least `n + burnin`. If the criterion is not met, the script will return less than the desired number of values and a warning will be given.

Value

A time series of length `n` from the specified SARIMA model with the specified frequency if the model is seasonal and start time `t0`.

Note

The model autoregressive polynomial ('AR side' = AR x SAR) is checked for causality and the model moving average polynomial ('MA side' = MA x SMA) is checked invertibility. The script stops and reports an error at the first violation of causality or invertibility; i.e., it will not report multiple errors.

Author(s)

D.S. Stoffer

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

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In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

Examples

```
## AR(2) with mean 50 [n = 500 is default]
y = sarima.sim(ar=c(1.5,-.75)) + 50
tsplot(y)

## ARIMA(0,1,1) with drift ['mean' refers to the innovations]
tsplot(sarima.sim(ma=-.8, d=1, mean=.1))

## SAR(1) example from text
set.seed(666) # not that 666
sAR = sarima.sim(sar=.9, S=12, n=36)
tsplot(sAR, type='c')
points(sAR, pch=Months, cex=1.1, font=4, col=1:4)

## SARIMA(0,1,1)x(0,1,1)_12 - B&J's favorite
set.seed(101010)
tsplot(sarima.sim(d=1, ma=-.4, D=1, sma=-.6, S=12, n=120))

## infinite variance t-errors
tsplot(sarima.sim(ar=.9, rand.gen=function(n, ...) rt(n, df=2) ))

## use your own innovations
dog = rexp(150, rate=.5)*sign(runif(150,-1,1))
tsplot(sarima.sim(n=100, ar=.99, innov=dog, burnin=50))

## generate seasonal data but no P, D or Q - you will receive
## a message to make sure that you wanted to do this on purpose:
tsplot(sarima.sim(ar=c(1.5,-.75), n=144, S=12), ylab='doggy', xaxt='n')
mtext(seq(0,144,12), side=1, line=.5, at=0:12)
```

Description

Draws a scatterplot with histograms in the margins.

Usage

```
scatter.hist(x, y, xlab = NULL, ylab = NULL, title = NULL, pt.size = 1,
             hist.col = gray(0.82), pt.col = gray(0.1, 0.25), pch = 19,
             reset.par = TRUE, ...)
```

Arguments

<code>x</code>	vector of x-values
<code>y</code>	corresponding vector of y-values
<code>xlab</code>	x-axis label (defaults to name of <code>x</code>)
<code>ylab</code>	y-axis label (defaults to name of <code>y</code>)
<code>title</code>	plot title (optional)
<code>pt.size</code>	size of points in scatterplot
<code>hist.col</code>	color for histograms
<code>pt.col</code>	color of points in scatterplot
<code>pch</code>	scatterplot point character
<code>reset.par</code>	reset graphics - default is TRUE; set to FALSE to add on to scatterplot
<code>...</code>	additional graphical parameters sent to <code>Grid</code>

Author(s)

D.S. Stoffer

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

Examples

```
scatter.hist(tempr, cmort, hist.col=astsa.col(5,.4), pt.col=5, pt.size=1.5, reset=FALSE)
lines(lowess(tempr, cmort), col=6)
```

```
scatter.hist(diff(log(econ5[, 'gnp'])), diff(log(econ5[, 'unemp'])), nxm=5, pt.size=2,
             col=astsa.col(6,.1), pt.col=astsa.col(4, alpha=.6, wheel=TRUE, num=25), hist='green3')
```

Description

Performs signal extraction and optimal filtering as discussed in Chapter 4.

Usage

```
SigExtract(series, L = c(3, 3), M = 50, max.freq = 0.05, col = 4)
```

Arguments

series	univariate time series to be filtered
L	degree of smoothing (may be a vector); see spans in spec.pgram for more details
M	number of terms used in the lagged regression approximation
max.freq	truncation frequency, which must be larger than 1/M
col	color of the main graphs

Details

The basic function of the script, and the default setting, is to remove frequencies above 1/20 (and, in particular, the seasonal frequency of 1 cycle every 12 time points). The sampling frequency of the time series is set to unity prior to the analysis.

See Example 4.29 for an example.

Value

Returns plots of (1) the original and filtered series, (2) the estimated spectra of each series, (3) the filter coefficients and the desired and attained frequency response function. The filtered series is returned invisibly.

Note

The script is based on code that was contributed by Professor Doug Wiens, Department of Mathematical and Statistical Sciences, University of Alberta.

Author(s)

D.S. Stoffer

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

sleep1

Sleep State and Movement Data - Group 1

Description

Sleep-state and number of movements of infants taken from a study on the effects of prenatal exposure to alcohol. This is Group 1 where the mothers did not drink alcohol during pregnancy.

Format

List of 12 (by subjects) `'data.frame'`: 120 obs. of 3 variables: `.. min : int [1:120] minute (1 to 120)`
`.. state: int [1:120] sleep state 1 to 6 with NA missing (see details)` `.. mvmnt: int [1:120] number of movements`

Details

Per minute sleep state, for approximately 120 minutes, is categorized into one of six possible states, non-REM: NR1 [1] to NR4 [4], and REM [5], or AWAKE [6]. NA means no state is recorded for that minute (if there, it occurs at end of the session). Group 1 (this group) is from mothers who abstained from drinking during pregnancy. In addition, the number of movements per minute are listed.

Source

Stoffer, D. S., Scher, M. S., Richardson, G. A., Day, N. L., Coble, P. A. (1988). A Walsh-Fourier Analysis of the Effects of Moderate Maternal Alcohol Consumption on Neonatal Sleep-State Cycling. *Journal of the American Statistical Association*, 83(404), 954-963. <https://doi.org/10.2307/2290119>

Stoffer, D. S. (1990). Multivariate Walsh-Fourier Analysis. *Journal of Time Series Analysis*, 11(1), 57-73. <https://doi.org/10.1111/j.1467-9892.1990.tb00042.x>

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also[sleep2](#)**Examples**

```
## Not run:

# spectral analysis
x = dna2vector(sleep1[[1]]$state[1:115], alphabet=c('1','2','3','4','5')) # never awake
specenv(x, spans=c(3,3))
abline(v=1/60, lty=2, col=8)

## End(Not run)
```

sleep2

*Sleep State and Movement Data - Group 2***Description**

Sleep-state and number of movements of infants taken from a study on the effects of prenatal exposure to alcohol. This is Group 2 where the mothers drank alcohol in moderation during pregnancy.

Format

List of 12 (by subjects) : 'data.frame': 120 obs. of 3 variables: .. min : int [1:120] minute (1 to 120)
 .. state: int [1:120] sleep state 1 to 6 with NA missing (see details) .. mvmnt: int [1:120] number of
 movements

Details

Per minute sleep state, for approximately 120 minutes, is categorized into one of six possible states, non-REM: NR1 [1] to NR4 [4], and REM [5], or AWAKE [6]. NA means no state is recorded for that minute (if there, it occurs at end of the session). Group 2 (this group) is from mothers who drank alcohol in moderation during pregnancy. In addition, the number of movements per minute are listed.

Source

Stoffer, D. S., Scher, M. S., Richardson, G. A., Day, N. L., Coble, P. A. (1988). A Walsh-Fourier Analysis of the Effects of Moderate Maternal Alcohol Consumption on Neonatal Sleep-State Cycling. *Journal of the American Statistical Association*, 83(404), 954-963. <https://doi.org/10.2307/2290119>

Stoffer, D. S. (1990). Multivariate Walsh-Fourier Analysis. *Journal of Time Series Analysis*, 11(1), 57-73. <https://doi.org/10.1111/j.1467-9892.1990.tb00042.x>

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[sleep1](#)

Examples

```
## Not run:

# spectral analysis
x = dna2vector(sleep1[[1]]$state[1:115], alphabet=c('1','2','3','4','5')) # never awake
specenv(x, spans=c(3,3))
abline(v=1/60, lty=2, col=8)

## End(Not run)
```

so2

SO2 levels from the LA pollution study

Description

Sulfur dioxide levels from the LA pollution study

Format

The format is: Time-Series [1:508] from 1970 to 1980: 3.37 2.59 3.29 3.04 3.39 2.57 2.35 3.38 1.5 2.56 ...

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[lap](#)

soi*Southern Oscillation Index*

Description

Southern Oscillation Index (SOI) for a period of 453 months ranging over the years 1950-1987.

Format

The format is: Time-Series [1:453] from 1950 to 1988: 0.377 0.246 0.311 0.104 -0.016 0.235 0.137 0.191 -0.016 0.29 ...

Details

pairs with rec (Recruitment)

Source

Data furnished by Dr. Roy Mendelsohn of the Pacific Fisheries Environmental Laboratory, NOAA (personal communication).

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[rec](#), [ENSO](#)

soiltemp*Spatial Grid of Surface Soil Temperatures*

Description

A 64 by 36 matrix of surface soil temperatures.

Format

The format is: num [1:64, 1:36] 6.7 8.9 5 6.6 6.1 7 6.5 8.2 6.7 6.6 ...

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

sp500.gr

Returns of the S&P 500

Description

Daily growth rate of the S&P 500 from 2001 though 2011.

Format

The format is: Time Series; Start = c(2001, 2); End = c(2011, 209); Frequency = 252

Source

Douc, Moulines, & Stoffer (2014). *Nonlinear Time Series: Theory, Methods and Applications with R Examples*. CRC Press. ISBN: <9781466502253>

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

sp500w

*Weekly Growth Rate of the Standard and Poor's 500***Description**

Weekly closing returns of the SP 500 from 2003 to September, 2012.

Format

An 'xts' object on 2003-01-03 to 2012-09-28; Indexed by objects of class: [Date] TZ: UTC

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

spec.ic

*Estimate Spectral Density of a Time Series from AR Fit***Description**

Fits an AR model to data and computes (and by default plots) the spectral density of the fitted model based on AIC (default) or BIC.

Usage

```
spec.ic(xdata, BIC=FALSE, order.max=NULL, main=NULL, plot=TRUE, detrend=TRUE,
        lowess=FALSE, method=NULL, cex.main=NULL, xlab=NULL, ...)
```

Arguments

xdata	a univariate time series.
BIC	if TRUE, fit is based on BIC. If FALSE (default), fit is based on AIC.
order.max	maximum order of model to fit. Defaults (if NULL) to the minimum of 100 and 10% of the number of observations.
main	plot title. Defaults to name of series, method and chosen order.
plot	if TRUE (default) produces a graphic of the estimated AR spectrum.
detrend	if TRUE (default), detrends the data first. If FALSE, the series is demeaned.
lowess	if TRUE, detrends using lowess. Default is FALSE.

method	method of estimation - a character string specifying the method to fit the model chosen from the following: "yule-walker", "burg", "ols", "mle", "yw". Defaults to "yule-walker".
cex.main	magnification for main title; default is 1.
xlab	label for frequency axis; if NULL (default), a totally awesome label is generated for your viewing pleasure.
...	additional graphical arguments.

Details

Uses `ar` to fit the best AR model based on pseudo AIC or BIC. Using `method='mle'` will be slow. The minimum centered AIC and BIC values and the spectral and frequency ordinates are returned silently.

Value

[[1]]	Matrix with columns: ORDER, AIC, BIC
[[2]]	Matrix with columns: freq, spec

Author(s)

D.S. Stoffer

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[ar](#), [spec.ar](#)

Examples

```
## Not run:
# AIC
spec.ic(soi)
spec.ic(sunspotz, method='burg', col=4)

# BIC after detrending on log scale
dog <- spec.ic(soi, BIC=TRUE, detrend=TRUE, log='y')

# plot AIC and BIC (dog from previous example)
tsplot(dog[[1]][,1], dog[[1]][,2:3], col=2*1:2, type='o', pch=1:2, xlab='order',
```

```

    spag=TRUE, addLegend=TRUE)

## End(Not run)

```

specenv

Spectral Envelope

Description

Computes the spectral envelope of categorical-valued or real-valued time series.

Usage

```

specenv(xdata, section = NULL, spans = NULL, kernel = NULL, taper = 0,
        significance = 1e-04, plot = TRUE, ylim = NULL, real = FALSE, ...)

```

Arguments

xdata	For categorical-valued sequences, a matrix with rows that are indicators of the categories represented by the columns, possibly a sequence converted using dna2vector . For real-valued sequences, a matrix with at least two columns that are various transformations of the data.
section	of the form start:end where start < end are positive integers; specifies the section used in the analysis - default is the entire sequence.
spans	specify smoothing used in mvspec.
kernel	specify kernel to be used in mvspec.
taper	specify amount of tapering to be used in mvspec.
significance	significance threshold exhibited in plot - default is .0001; set to NA to cancel
plot	if TRUE (default) a graphic of the spectral envelope is produced
ylim	limits of the spectral envelope axis; if NULL (default), a suitable range is calculated.
real	FALSE (default) for categorical-valued sequences and TRUE for real-valued sequences.
...	other graphical parameters.

Details

Calculates the spectral envelope for categorical-valued series as discussed in https://www.stat.pitt.edu/stoffer/dss_files/spenv.pdf and summarized in <https://doi.org/10.1214/ss/1009212816>.

Alternately, calculates the spectral envelope for real-valued series as discussed in [https://doi.org/10.1016/S0378-3758\(96\)00044-4](https://doi.org/10.1016/S0378-3758(96)00044-4).

These concepts are also presented (with examples) in Section 7.9 (Chapter 7) of Time Series Analysis and Its Applications: With R Examples: <https://www.stat.pitt.edu/stoffer/tsa4/>.

For categorical-valued series, the input `xdata` must be a matrix of indicators which is perhaps a sequence preprocessed using `dna2vector`.

For real-valued series, the input `xdata` should be a matrix whose columns are various transformations of the univariate series.

The script does not detrend the data prior to estimating spectra. If this is an issue, then detrend the data prior to using this script.

Value

By default, will produce a graph of the spectral envelope and an approximate significance threshold. A matrix containing: frequency, spectral envelope ordinates, and (1) the scalings of the categories in the order of the categories in the alphabet or (2) the coefficients of the transformations, is returned invisibly.

Author(s)

D.S. Stoffer

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[dna2vector](#)

Examples

```
## Not run:
# a DNA sequence
data = bnrfl1ebv
xdata = dna2vector(data)
u = specenv(xdata, section=1:1000, spans=c(7,7))
head(u) # scalings are for A, C, G, and last one T=0 always

# a real-valued series (nyse returns)
x = astsa::nyse
xdata = cbind(x, abs(x), x^2)
u = specenv(xdata, real=TRUE, spans=c(3,3))
# plot optimal transform at freq = .001
beta = u[2, 3:5]
b = beta/beta[2] # makes abs(x) coef=1
gopt = function(x) { b[1]*x+b[2]*abs(x)+b[3]*x^2 }
curve(gopt, -.2, .2, col=4, lwd=2, panel.first=Grid())
```

```
g2 = function(x) { b[2]*abs(x) } # corresponding to |x|
curve(g2, -.2,.2, add=TRUE, col=6)

## End(Not run)
```

speech

*Speech Recording***Description**

A small .1 second (1000 points) sample of recorded speech for the phrase "aaa...hhh".

Format

The format is: Time-Series [1:1020] from 1 to 1020: 1814 1556 1442 1416 1352 ...

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

ssm

*State Space Model***Description**

Fits a simple univariate state space model to data. The parameters are estimated (the state regression parameter may be fixed). State predictions, filters, and smoothers and corresponding error variances are evaluated at the estimates. The sample size must be at least 20.

Usage

```
ssm(y, A, phi, alpha, sigw, sigv, fixphi = FALSE)
```

Arguments

y	data
A	measurement value (fixed constant)
phi	initial value of phi, may be fixed
alpha	initial value for alpha
sigw	initial value for sigma[w]
sigv	initial value for sigma[v]
fixphi	if TRUE, the phi parameter is fixed

Details

The script works for a specific univariate state space model,

$$x_t = \alpha + \phi x_{t-1} + w_t \quad \text{and} \quad y_t = Ax_t + v_t.$$

The initial state conditions use a default calculation and cannot be specified. The parameter estimates are printed and the script returns the state predictors and smoothers. The regression parameter ϕ may be fixed.

Value

At the MLEs, these are returned invisibly:

Xp	time series - state prediction, x_t^{t-1}
Pp	corresponding MSPEs, P_t^{t-1}
Xf	time series - state filter, x_t^t
Pf	corresponding MSEs, P_t^t
Xs	time series - state smoother, x_t^n
Ps	corresponding MSEs, P_t^n

Author(s)

D.S. Stoffer

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

Examples

```
## Not run:

u = ssm(gtemp_land, A=1, alpha=.01, phi=1, sigw=.05, sigv=.15, fixphi=TRUE)
tsplot(gtemp_land, type='o', col=4)
lines(u$Xs, col=6, lwd=2)

## End(Not run)
```

star

Variable Star

Description

The magnitude of a star taken at midnight for 600 consecutive days. The data are taken from the classic text, *The Calculus of Observations, a Treatise on Numerical Mathematics*, by E.T. Whittaker and G. Robinson, (1923, Blackie and Son, Ltd.).

Format

The format is: Time-Series [1:600] from 1 to 600: 25 28 31 32 33 33 32 ...

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

stoch.reg

Frequency Domain Stochastic Regression

Description

Performs frequency domain stochastic regression discussed in Chapter 7.

Usage

```
stoch.reg(xdata, cols.full, cols.red=NULL, alpha, L, M, plot.which, col.resp=NULL, ...)
```

Arguments

xdata	data matrix with the last column being the response variable
cols.full	specify columns of data matrix that are in the full model
cols.red	specify columns of data matrix that are in the reduced model (use NULL if there are no inputs in the reduced model)
alpha	test size; number between 0 and 1
L	odd integer specifying degree of smoothing
M	number (integer) of points in the discretization of the integral

<code>plot.which</code>	coh or F.stat, to plot either the squared-coherencies or the F-statistics, respectively
<code>col.resp</code>	specify column of the response variable if it is not the last column of the data matrix
<code>...</code>	additional graphic arguments

Details

This function computes the spectral matrix, F statistics and coherencies, and plots them. Returned as well are the coefficients in the impulse response function.

Enter, as the argument to this function, the full data matrix, and then the labels of the columns of input series in the "full" and "reduced" regression models - enter NULL if there are no inputs under the reduced model.

If the response variable is the LAST column of the data matrix, it need not be specified. Otherwise specify which column holds the responses as `col.resp`.

Other inputs are alpha (test size), L (smoothing), M (number of points in the discretization of the integral) and `plot.which` = "coh" or "F", to plot either the coherencies or the F statistics.

Value

<code>power.full</code>	spectrum under the full model
<code>power.red</code>	spectrum under the reduced model
<code>Betahat</code>	regression parameter estimates
<code>eF</code>	pointwise (by frequency) F-tests
<code>coh</code>	coherency

Note

See Example 7.1 of the text. The script is based on code that was contributed by Professor Doug Wiens, Department of Mathematical and Statistical Sciences, University of Alberta.

Author(s)

D.S. Stoffer

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

sunspotz

*Biannual Sunspot Numbers***Description**

Biannual smoothed (12-month moving average) number of sunspots from June 1749 to December 1978; $n = 459$. The "z" on the end is to distinguish this series from the one included with R (called sunspots).

Format

The format is: Time Series: Start = c(1749, 1) End = c(1978, 1) Frequency = 2

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

SV.mcmc

*Fit Bayesian Stochastic Volatility Model***Description**

Fits a stochastic volatility model to a univariate time series of returns.

Usage

```
SV.mcmc(y, nmcmc = 1000, burnin = 100, init = NULL, hyper = NULL, tuning = NULL,
        sigma_MH = NULL, npart = NULL, mcmseed = NULL)
```

Arguments

y	single time series of returns
nmcmc	number of iterations for the MCMC procedure
burnin	number of iterations to discard for the MCMC procedure
init	initial values of (phi, sigma, beta) - default is c(0.9, 0.5, .1)
hyper	hyperparameters for bivariate normal distribution of (phi, sigma); user inputs (mu_phi, mu_q, sigma_phi, sigma_q, rho) - default is c(0.9, 0.5, 0.075, 0.3, -0.25)

tuning	tuning parameter - default is .03
sigma_MH	covariance matrix used for random walk Metropolis; it will be scaled by tuning in the script - default is <code>matrix(c(1,-.25,-.25,1), nrow=2, ncol=2)</code>
npart	number of particles used in particle filter - default is 10
mcmseed	seed for mcmc - default is 90210

Details

The log-volatility process is x_t and the returns are y_t . The SV model is

$$x_t = \phi x_{t-1} + \sigma w_t \quad y_t = \beta \exp\left\{\frac{1}{2}x_t\right\} \epsilon_t$$

where w_t and ϵ_t are independent standard normal white noise.

The model is fit using a technique described in the paper listed below (in the Source section) where the state parameters (ϕ, σ) are sampled simultaneously with a bivariate normal prior specified in the arguments `init` and `hyper`.

Two graphics are returned: (1) the three parameter traces with the posterior mean highlighted, their ACFs [with effective sample sizes (ESS)], and their histograms with the .025, .5, and .975 quantiles displayed, and (2) the log-volatility posterior mean along with corresponding .95 credible intervals.

Value

Returned invisibly:

phi	vector of sampled state AR parameter
sigma	vector of sampled state error std deviation
beta	vector of sampled observation error scale
log.vol	matrix of sampled log-volatility
options	values of the input arguments

Note

Except for the data, all the other inputs have defaults. The time to run and the acceptance rate are returned at the end of the analysis. The acceptance rate should be around 30% and this is easily adjusted using the tuning parameter.

Author(s)

D.S. Stoffer

Source

Gong & Stoffer (2021). A note on efficient fitting of stochastic volatility models. *Journal of Time Series Analysis*, 42(2), 186-200. <https://github.com/nickpoison/Stochastic-Volatility-Models>

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[SV.mle](#)

Examples

```
## Not run:
#-- A minimal example --##
myrun <- SV.mcmc(sp500w) # results in object myrun - don't forget it

str(myrun) # an easy way to see the default input options

## End(Not run)
```

SV.mle

Stochastic Volatility Model with Feedback via MLE

Description

Fits a stochastic volatility model with feedback (optional) to a univariate time series of returns via quasi-MLE.

Usage

```
SV.mle(returns, gamma = 0, phi = 0.95, sQ = 0.1, alpha = NULL, sR0 = 1, mu1 = -3, sR1 = 2,
      rho = NULL, feedback = FALSE)
```

Arguments

returns	single time series of returns
gamma	feedback coefficient - included if feedback=TRUE (does not have to be specified)
phi	initial value of the log-volatility AR parameter (does not have to be specified)
sQ	initial value of the standard deviation of log-volatility noise (does not have to be specified)
alpha	initial value of the log-returns ² constant parameter (does not have to be specified)

sR0	initial value of the log-returns ² normal mixture standard deviation parameter (component 0 - does not have to be specified)
mu1	initial value of the log-returns ² normal mixture mean parameter (component 1 - does not have to be specified)
sR1	initial value of the log-returns ² normal mixture standard deviation parameter (component 1 - does not have to be specified)
rho	correlation between the state noise and observation noise (so called "leverage"). If feedback=TRUE this will be included if given a proper numerical value; if NULL (default) it is not included because it is often not significant when the feedback coefficient is included.
feedback	if TRUE feedback is included in the model; default is FALSE.

Details

The returns are r_t (input this). The log-volatility process is x_t and $y_t = \log r_t^2$.

If feedback=TRUE, the model is

$$x_{t+1} = \gamma r_t + \phi x_t + \sigma w_t \quad y_t = \alpha + x_t + \eta_t$$

where w_t is standard normal noise. The observation error η_t is a mixture of two normals, $N(0, \sigma_0^2)$ and $N(\mu_1, \sigma_1^2)$. The state and observation noise can be correlated if ρ is given a value between -1 and 1.

If feedback=FALSE, γ and ρ are not included in the model.

Value

Returned invisibly:

PredLogVol	one-step-ahead predicted log-volatility
RMSPE	corresponding root MSPE
Coefficients	table of estimates and estimated standard errors

In addition to the one step ahead predicted log-volatility, corresponding root MSPE, and table of estimates returned invisibly, the estimates and SEs are printed and a graph of (1) the data with the predicted log-volatility, and (2) the normal mixture are displayed in one graphic.

Author(s)

D.S. Stoffer

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also[SV.mcmc](#)**Examples**

```
## Not run:  
  
SV.mle(sp500.gr, feedback=TRUE)  
  
SV.mle(nyse)  
  
## End(Not run)
```

temp*Temperatures from the LA pollution study*

Description

Temperature series corresponding to cmort from the LA pollution study.

Format

The format is: Time-Series [1:508] from 1970 to 1980: 72.4 67.2 62.9 72.5 74.2 ...

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also[lap](#), [lap.xts](#)

test.linear

*Test Linearity of a Time Series via Normalized Bispectrum***Description**

Produces a plot of the tail probabilities of a normalized bispectrum of a series under the assumption the model is a linear process with iid innovations.

Usage

```
test.linear(series, color = TRUE, detrend = FALSE, main = NULL)
```

Arguments

series	the time series (univariate only)
color	if FALSE, the graphic is produced in gray scale
detrend	if TRUE, the series is detrended first
main	if NULL (default), a very nice title is chosen for the plot

Value

prob	matrix of tail probabilities - returned invisibly
------	---

Note

The null hypothesis is that the data are from a linear process with i.i.d. innovations. Under the null hypothesis, the bispectrum is constant over all frequencies. Chi-squared test statistics are formed in blocks to measure departures from the null hypothesis and the corresponding p-values are displayed in a graphic and returned invisibly. Details are in Hinich, M. and Wolinsky, M. (2005). Normalizing bispectra. *Journal of Statistical Planning and Inference*, 130, 405–411.

Author(s)

D.S. Stoffer

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

Examples

```
## Not run:
test.linear(nyse) # :(
test.linear(soi)  # :)

## End(Not run)
```

trend	<i>Estimate Trend</i>
-------	-----------------------

Description

Estimates the trend (polynomial or lowess) of a time series and returns a graphic of the series with the trend and error bounds (returned invisibly) superimposed.

Usage

```
trend(series, order = 1, lowess = FALSE, lowspan = .75, robust = TRUE,
      col = c(4, 6), ylab = NULL, ci = TRUE, results = FALSE, ...)
```

Arguments

series	The time series to be analyzed (univariate only).
order	Order of the polynomial used to estimate the trend with a linear default (order=1) unless lowess is TRUE.
lowess	If TRUE, loess from the stats package is used to fit the trend. The default is FALSE.
lowspan	The smoother span used for lowess.
robust	If TRUE (default), the lowess fit is robust.
col	Vector of two colors for the graphic, first the color of the data (default is blue [4]) and second the color of the trend (default is magenta [6]). Both the data and trend line will be the same color if only one value is given.
ylab	Label for the vertical axis (default is the name of the series).
ci	If TRUE (default), pointwise 95% confidence intervals are drawn.
results	For polynomial regression, if TRUE, will print a summary (using ttable) of the regression results.
...	Other graphical parameters.

Details

Produces a graphic of the time series with the trend and a .95 pointwise confidence interval superimposed. The trend estimate and the error bounds are returned invisibly.

Value

Produces a graphic and returns the trend estimate fit and error bounds lwr and upr invisibly (see details) and with the same time series attributes as the input series.

Author(s)

D.S. Stoffer

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[detrend](#)

Examples

```
## Not run:

par(mfrow=2:1)
trend(soi)
trend(soi, lowess=TRUE)

## End(Not run)
```

tspairs

Scatterplot Matrix for Time Series

Description

Produces a matrix of scatterplots with the time series (or a histogram) plotted on the diagonal.

Usage

```
tspairs(x, main = NA, pt.col = astsa.col(4, 0.6), pt.size = 1.1, lab.size = 1.25,
        title.size = 1.5, scale = 1, corr = TRUE, smooth = TRUE, lw1 = 1, lwc = 2,
        gg = FALSE, hist.diag = TRUE, col.diag = 4, location='topright', ...)
```

Arguments

<code>x</code>	multiple time series; use <code>ts.intersect</code> to include lagged values.
<code>main</code>	title (default is no title).
<code>pt.col</code>	point color.
<code>pt.size</code>	point size.
<code>lab.size</code>	label size.
<code>title.size</code>	title size.
<code>scale</code>	multiplier for the overall character expansion (<code>cex</code>).
<code>corr</code>	if TRUE (default), the correlations are shown in the scatterplots.
<code>smooth</code>	if TRUE (default), a lowess fit is displayed in the scatterplots.
<code>lwl</code>	width of the lowess line.
<code>lwc</code>	color of the lowess line.
<code>gg</code>	if TRUE, will produce a gris-gris plot (gray graphic interior with white grid lines); the default is FALSE. The grammar of <code>astsa</code> is <code>voodoo</code> .
<code>hist.diag</code>	if TRUE (default), will plot histograms on the diagonal; if FALSE, time plots of the series are displayed instead.
<code>col.diag</code>	color for the diagonal plots.
<code>location</code>	the location of the ACF legend with options "bottomright", "bottom", "bottomleft", "left", "topleft", "top", "topright" (the default), "right" and "center".
<code>...</code>	additional graphic parameters.

Value

Returns a matrix of scatterplots with time plots or histograms on the diagonal.

Note

Use `ts.intersect` if lagged variables are included, otherwise `cbind` will work.

Author(s)

D.S. Stoffer

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[lag1.plot](#), [lag2.plot](#)

Examples

```
## Not run:

tspairs(diff(log(econ5)), col.diag=6, pt.size=1.5, lwl=2, gg=TRUE, las=0)

tspairs(ts.intersect(cmort,tempr,partL4=lag(part,-4)), hist=FALSE, pch=9, scale=1.1)

## End(Not run)
```

tsplot	<i>Time Series Plot</i>
--------	-------------------------

Description

Produces a nice plot of univariate or multiple time series in one easy line.

Usage

```
tsplot(x, y=NULL, main=NULL, ylab=NULL, xlab='Time', type=NULL, margins=.25,
       ncolm=1, byrow=TRUE, nx=NULL, ny=nx, minor=TRUE, nxm=2, nym=1,
       xm.grid=TRUE, ym.grid=TRUE, col=1, gg=FALSE, spaghetti=FALSE, pch=NULL,
       lty=1, lwd=1, mgpp=0, topper=NULL, addLegend=FALSE, location='topright',
       boxit=TRUE, horiz=FALSE, legend=NULL, llwd=NULL, scale=1, ...)
```

Arguments

x, y	time series to be plotted; if both present, x will be the time index.
main	add a plot title - the default is no title.
ylab	y-axis label - the default is the name of the ts object.
xlab	x-axis label - the default is 'Time'.
type	type of plot - the default is line.
margins	inches to add (or subtract) to the margins. Input one value to apply to all margins or a vector of length 4 to add (or subtract) to the (bottom, left, top, right) margins.
ncolm	for multiple time series, the number of columns to plot.
byrow	for multiple time series - if TRUE (default), plot series row wise; if FALSE, plot series column wise.
nx, ny	number of major cells of the grid in x and y direction. When NULL, as per default, the grid aligns with the tick marks on the corresponding default axis (i.e., tickmarks as computed by axTicks). When NA, no grid lines are drawn in the corresponding direction.

minor, nxm, nym	if minor=TRUE, the number of minor tick marks on x-axis, y-axis. minor=FALSE removes both or set either to 0 or 1 to remove. The default is one minor tick on the x-axis and none on the y-axis.
xm.grid, ym.grid	if TRUE (default), adds grid lines at minor x-axis, y-axis ticks.
col	line color(s), can be a vector for multiple time series.
gg	if TRUE, will produce a gris-gris plot (gray graphic interior with white grid lines); the default is FALSE. The grammar of astsa is voodoo; see https://www.youtube.com/watch?v=b4J8VrprGE
spaghetti	if TRUE, will produce a spaghetti plot (all series on same plot).
pch	plot symbols (default is 1, circle); can be a vector for multiple plots.
lty	line type (default is 1, solid line); can be a vector for multiple plots.
lwd	line width (default is 1); can be a vector for multiple plots.
mgpp	this is used to adjust (add to) the mpg graphics parameters settings (?par), which are c(1.6, .6, 0) here; the R default is c(3, 1, 0). This will be helpful in moving an axis label farther from the axis if necessary.
topper	non-negative value to add to the top outer margin; if NULL (default) a suitable value is chosen
addLegend	if TRUE and spaghetti=TRUE, will add a simple legend. If more details are needed, leave this set to FALSE (the default) and use legend directly.
location	if addLegend=TRUE, the location of the legend with options "bottomright", "bottom", "bottomleft", "left", "topleft", "top", "topright" (the default), "right" and "center".
boxit	if TRUE (default), the legend is in a box; if FALSE, no box is drawn.
horiz	if addLegend=TRUE, should the legend be horizontal (default is FALSE - vertical).
legend	if NULL (default), the legend uses names of each time series; otherwise, use to customize legend.
llwd	line width for the legend if different from the plotted lines.
scale	for multiple series, the scale for character expansion (cex = .9*scale).
...	other graphical parameters; see par .

Value

Produces a graphic and returns it invisibly so it can be saved in an R variable with the ability to replay it; see [recordPlot](#).

Note

A legend can be added using addLegend=TRUE for spaghetti plots only. Spaghetti plots work if spaghetti=TRUE and there is more than one series being plotted.

Author(s)

D.S. Stoffer

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

Examples

```
## Not run:

# minimal
tsplot(soi)
# prettified
tsplot(soi, col=4, main="Southern Oscillation Index")

# gris-gris multiple plot
tsplot(climhyd, ncolm=2, gg=TRUE, col=2:7, lwd=2)

# spaghetti (and store it in an object - ?recordPlot for details)
x <- replicate(100, cumsum(rcauchy(1000))/1:1000)
u <- tsplot(x, col=1:8, main='No LLN For You', spaghetti=TRUE)
u # plot on demand

# quick legend with spaghetti
tsplot(cbind(Mortality=cmort, Pollution=part), col=5:6, gg=TRUE, spaghetti=TRUE, addLegend=TRUE)
# ... and a little more control
tsplot(cbind(gtemp_land, gtemp_ocean), spaghetti=TRUE, lwd=2, col=astsa.col(c(4,2),.7),
  ylab="\u00B0C", main="Global Temperature Anomalies", addLegend=TRUE, location='topleft',
  horiz=TRUE, legend=c("Land Surface", "Sea Surface"))

# compare these
par(mfrow=2:1)
tsplot(1:453, soi, ylab='SOI', xlab='Month')
# now recklessly add to the margins and add to mgp to get to the default
tsplot(1:453, soi, ylab='SOI', xlab='Month', margins=c(2,3,4,5), las=1, mgpp=c(1.4,.4,0))

# cex (and scale)
par(mfrow=c(3,1)) # cex gets small with mf plots
  tsplot(cmort); tsplot(tempr); tsplot(part)
par(mfrow=c(3,1), cex=.9) # so fix it
  tsplot(cmort); tsplot(tempr); tsplot(part)

# it's rescaled here for your pleasure
tsplot(cbind(big=rnorm(100), bad=rnorm(100), john=rnorm(100))) # default scale=1
tsplot(cbind(big=rnorm(100), bad=rnorm(100), john=rnorm(100)), scale=1.5) # big

## End(Not run)
```

ttable	<i>t-table summary for an lm object</i>
--------	---

Description

Works like `summary` for an `lm` object but adds AIC, AICc, and BIC to the output with the option to display VIFs.

Usage

```
ttable(obj, digits = 4, vif = FALSE, ...)
```

Arguments

<code>obj</code>	an object of class "lm" as a result of a call to lm .
<code>digits</code>	the (approximate) number of significant digits to use when printing.
<code>vif</code>	if TRUE, variance inflation factors are printed if applicable; default is FALSE.
<code>...</code>	further arguments passed to or from other methods.

Details

Produces a t-table for an `lm` object much like [print.summary.lm](#) with added information including AIC, AICc, BIC, and VIF (if requested and if applicable), to the output. The output is rounded and there are no significance stars. In fact, there are no stars at all.

TO REPEAT THE WARNING ON USING 'lm' FOR TIME SERIES: Considerable care must be taken:

- Include `na.action = NULL` in the `lm` call to avoid stripping the time series attributes from the variables before the regression is done.
- If any lagged or differenced variables are used in 'lm', the series must be lined up first. In this case, prepare a data frame using `ts.intersect(..., dframe = TRUE)`; e.g.,

```
mydata = ts.intersect(M = cmort, P4 = lag(part,-4), dframe=TRUE)
fit     = lm(M ~ P4, na.action=NULL, data=mydata)
ttable(fit)
```

Value

Prints a typical t-table with additional information as mentioned in the details. The p-values are two-sided. Also silently returns the same values as described in [summary.lm](#).

Author(s)

D.S. Stoffer

Source

Built using `print.summary.lm` from the 'stats' package.

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[summary.lm](#), [print.summary.lm](#), [lm](#)

Examples

```
fit1 = lm(cmort~ time(cmort) + tempr + I(tempr^2))
ttable(fit1, vif=TRUE)
# if you center `tempr`, the squared term doesn't change
temp = tempr - mean(tempr)
fit2 = lm(cmort~ time(cmort) + temp + I(temp^2))
ttable(fit2, vif=TRUE)
```

unemp

U.S. Unemployment

Description

Monthly U.S. Unemployment series (1948-1978, $n = 372$)

Usage

```
data(unemp)
```

Format

The format is: Time-Series [1:372] from 1948 to 1979: 235 281 265 241 201 ...

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[UnempRate](#)

UnempRate	<i>U.S. Unemployment Rate</i>
-----------	-------------------------------

Description

Monthly U.S. unemployment rate in percent unemployed (Jan, 1948 - Nov, 2016, n = 827)

Format

The format is: Time-Series [1:827] from 1948 to 2017: 4 4.7 4.5 4 3.4 3.9 3.9 3.6 3.4 2.9 ...

Source

<https://data.bls.gov/timeseries/LNU04000000/>

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[unemp](#)

USpop

U.S. Population - 1900 to 2010

Description

U.S. Population by official census, every ten years from 1900 to 2010.

Format

The format is: Time-Series [1:12] from 1900 to 2010: 76 92 106 123 132 ...

Details

The census from 2020 is not included in this data set because, by many accounts, it was a nightmare (<https://www.npr.org/2022/01/15/1073338121/2020-census-interference-trump>) due to the COVID-19 pandemic coupled with the fact that the Census Bureau is in the Department of Commerce, and its head is appointed by and reports directly to the POTUS, who at the time was DJ tRump: "Historians rank Trump among worst presidents in US history ... " (<https://www.businessinsider.com/historia>

The data with the 2020 census is in [USpop20](#). Note that the two data files differ a bit, presumably because they are revised ad infinitum. Both data sets were obtained from the Census Bureau's website.

Source

<https://www.census.gov/>

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[USpop20](#)

USpop20

*U.S. Population - 1900 to 2020***Description**

U.S. Population by official census, every ten years from 1900 to 2020, in millions.

Format

The format is: Time-Series [1:13] from 1900 to 2020: 76.2 92.2 106 122.8 132.2 ...

Details

The census from 2020 is included in this data set, but the results from 2020 are questionable. By many accounts, it was a nightmare (<https://www.npr.org/2022/01/15/1073338121/2020-census-interference-trump>) due to the COVID-19 pandemic coupled with the fact that the Census Bureau is in the Department of Commerce, and its head is appointed by and reports directly to the POTUS, who at the time was DJ tRump: "Historians rank Trump among worst presidents in US history ... " (<https://www.businessinsider.com/historians-rank-trump-among-worst-presidents-us-history-c-span-2020>)

The data without the 2020 census is in [USpop](#). Note that the two data files differ a bit, presumably because they are revised ad infinitum. Both data sets were obtained from the Census Bureau's website.

Source

<https://www.census.gov/>

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[USpop](#)

Examples

```
## Not run:

# This is from the 2nd edition of Time Series: A Data Analysis Approach ...
t = time(USpop20) - 1960
reg = lm( USpop20~ poly(t, 10, raw=TRUE) )
```

```

b = as.vector(coef(reg))
g = function(s){b[1] + b[2]*s + b[3]*s^2 + b[4]*s^3 + b[5]*s^4 + b[6]*s^5 + b[7]*s^6 +
               b[8]*s^7 + b[9]*s^8 + b[10]*s^9 + b[11]*s^10 }
t = 1900:2044
tsplot(t, g(t-1960), ylab="Population", xlab='Year',cex.main=1, col=4,
       main="U.S. Population by Official Census")
points(time(USpop20), USpop20, pch=21, bg=rainbow(13), cex=1.25)
mtext(bquote('\u00D7'~10^6), side=2, line=1.5, adj=1, cex=.8)

## End(Not run)

```

varve

Annual Varve Series

Description

Sedimentary deposits from one location in Massachusetts for 634 years, beginning nearly 12,000 years ago.

Format

The format is: Time-Series [1:634] from 1 to 634: 26.3 27.4 42.3 58.3 20.6 ...

Source

Antevs, E. (1928). The Last Glaciation with Special Reference to the Ice Retreat in Northeastern North America. *Amer. Geog. Soc. Res. Series*, No. 17, 292p.

Shumway and Verosub (1992). State space modeling of paleoclimatic time series. *Proceedings of the 5th International Meeting on Statistical Climatology*, June, 1992, University of Toronto, 139p.

References

You can find demonstrations of `astsa` capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

WBC*White Blood Cell Levels*

Description

WBC: Measurements made for 91 days on the three variables, log(white blood count) [WBC], log(platelet) [PLT] and hematocrit [HCT]. Missing data code is 0 (zero).

Format

The format is: Time-Series [1:91] from 1 to 91: 2.33 1.89 2.08 1.82 1.82 ...

Details

See Examples 6.1 and 6.9 for more details.

Source

Jones, R.H. (1984). Fitting multivariate models to unequally spaced data. In *Time Series Analysis of Irregularly Observed Data*, pp. 158-188. E. Parzen, ed. Lecture Notes in Statistics, 25, New York: Springer-Verlag.

References

You can find demonstrations of astsa capabilities at [FUN WITH ASTSA](#).

The most recent version of the package can be found at <https://github.com/nickpoison/astsa/>.

In addition, the News and ChangeLog files are at <https://github.com/nickpoison/astsa/blob/master/NEWS.md>.

The webpages for the texts and some help on using R for time series analysis can be found at <https://nickpoison.github.io/>.

See Also

[blood](#), [HCT](#), [PLT](#)

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