

ITSM-R Reference Manual

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1 Introduction

ITSM-R is an R package for analyzing and forecasting univariate time series data. The intended audience is students using the textbook *Introduction to Time Series and Forecasting* by Peter J. Brockwell and Richard A. Davis. The textbook includes a student version of the Windows-based program ITSM 2000. The initials ITSM stand for Interactive Time Series Modeling. ITSM-R provides a subset of the functionality found in ITSM 2000.

The following example shows ITSM-R at work forecasting a time series.

```
library(itsmr)           # Load ITSM-R
plotc(wine)              # Plot the data
M = c("log","season",12,"trend",1) # Model the data
e = Resid(wine,M)        # Obtain residuals
test(e)                  # Check for stationarity
a = arma(e,p=1,q=1)      # Model the residuals
ee = Resid(wine,M,a)     # Obtain secondary residuals
test(ee)                 # Check for white noise
forecast(wine,M,a)       # Forecast future values
```

Note the use of `Resid` with a capital R to compute residuals. ITSM-R uses `Resid` instead of `resid` to avoid a name conflict warning when the library is loaded.

1.1 Time series analysis in a nutshell

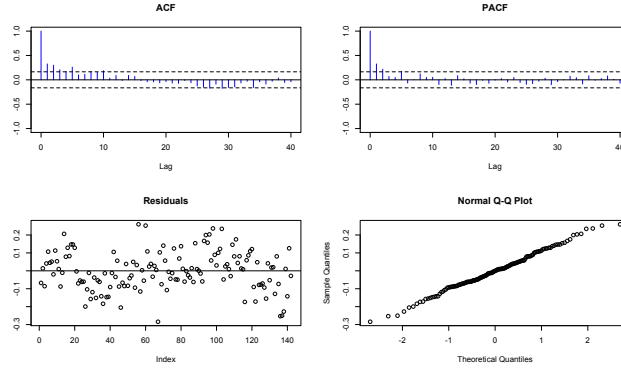
It turns out that you do most of the work. You have to analyze the data and determine a data model M and ARMA model a that yield white noise for residuals.

$$\text{Time series data} \longrightarrow M \longrightarrow a \longrightarrow \text{White noise}$$

What the software does is invert M and a to transform zero into a forecast. (Recall that zero is the best predictor of noise.)

$$0 \longrightarrow a^{-1} \longrightarrow M^{-1} \longrightarrow \text{Forecast}$$

Model M should yield residuals that are a stationary time series. Basically this means that any trend in the data is eliminated. (Seasonality in the data can be removed by either M or a .) The `test` function is used to determine if the output of M is stationary. The following is the result of `test(e)` in the above example.



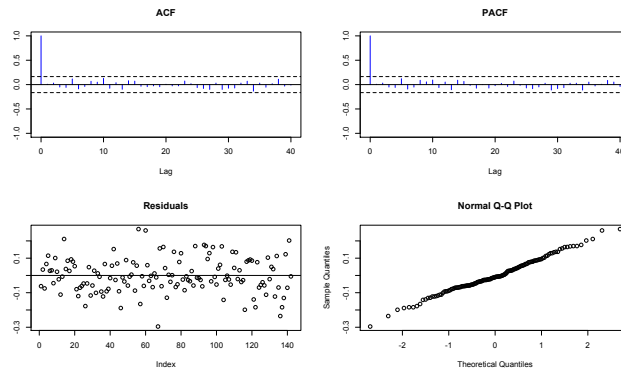
Null hypothesis: Residuals are iid noise.

Test	Distribution	Statistic	p-value
Ljung-Box Q	$Q \sim \text{chisq}(20)$	71.57	0 *
McLeod-Li Q	$Q \sim \text{chisq}(20)$	12.07	0.9138
Turning points T	$(T-93.3)/5 \sim N(0,1)$	93	0.9468
Diff signs S	$(S-70.5)/3.5 \sim N(0,1)$	70	0.8848
Rank P	$(P-5005.5)/283.5 \sim N(0,1)$	5136	0.6453

As seen above, the ACF and PACF plots decay rapidly indicating stationarity. After M is determined, the residuals of M are modeled with an ARMA model a . In some circumstances the following table can be useful for choosing p and q from the plots of `test(e)`.

ACF	PACF	Model
Decays	Zero for $h > p$	$\text{AR}(p)$
Zero for $h > q$	Decays	$\text{MA}(q)$
Decays	Decays	$\text{ARMA}(p, q)$

The ARMA model a should yield residuals that resemble white noise. To check for white noise, the `test` function is used a second time. The following is the result of `test(ee)` in the above example.



Null hypothesis: Residuals are iid noise.

Test	Distribution	Statistic	p-value
Ljung-Box Q	$Q \sim \text{chisq}(20)$	13.3	0.8642
McLeod-Li Q	$Q \sim \text{chisq}(20)$	16.96	0.6556

Turning points T	(T-93.3)/5 ~ N(0,1)	93	0.9468
Diff signs S	(S-70.5)/3.5 ~ N(0,1)	70	0.8848
Rank P	(P-5005.5)/283.5 ~ N(0,1)	4999	0.9817

The vertical lines in the ACF and PACF plots are below the noise threshold (the dashed horizontal lines) for lags greater than zero which is good. In addition, the p -values in the table all fail to reject the null hypothesis of iid noise. Recall that a p -value is the probability that the null hypothesis is true.

1.2 White Noise Variance

ITSM-R includes five methods for estimating ARMA parameters. They are Yule-Walker, Burg, Hannan-Rissanen, maximum likelihood, and the innovations method. For all estimation methods, ITSM-R uses the following formula to estimate white noise variance (Brockwell and Davis, p. 164).

$$\hat{\sigma}^2 = \frac{1}{n} \sum_{t=1}^n \frac{(X_t - \hat{X}_t)^2}{r_{t-1}}$$

The $X_t - \hat{X}_t$ are innovations (Brockwell and Davis, p. 71). Residuals are defined as follows (Brockwell and Davis, p. 164).

$$\hat{W}_t = \frac{X_t - \hat{X}_t}{\sqrt{r_{t-1}}}$$

Thus $\hat{\sigma}^2$ is the average of the squared residuals.

The innovations algorithm (Brockwell and Davis, p. 73) is used to compute \hat{X}_t and r_{t-1} . Note that the algorithm produces mean squared errors $v_{t-1} = E(X_t - \hat{X}_t)^2$ for $\kappa(i, j) = E(X_i X_j)$ given in equation (2.5.24). However, ITSM-R uses $\kappa(i, j) = E(W_i W_j)$ given in equation (3.3.3). For the covariance in (3.3.3) the innovations algorithm produces $r_{t-1} = E(W_t - \hat{W}_t)^2$ instead of v_{t-1} . The relationship between v_{t-1} and r_{t-1} is given in equation (3.3.8) as

$$v_{t-1} = E(X_t - \hat{X}_t)^2 = \sigma^2 E(W_t - \hat{W}_t)^2 = \sigma^2 r_{t-1}$$

where σ^2 is white noise variance.

It should be noted that γ_X in (3.3.3) is the autocovariance of the ARMA model, not of the data. Then by equation (3.2.3) it can be seen that σ^{-2} in (3.3.3) cancels with σ^2 in γ_X . Hence the innovations algorithm does not depend on white noise variance at all. After the innovations are computed, white noise variance is estimated using the above formula for $\hat{\sigma}^2$.

$$\gamma_X(h) = E(X_{t+h} X_t) = \sigma^2 \sum_{j=0}^{\infty} \psi_j \psi_{j+|h|} \quad (3.2.3)$$

$$m = \max(p, q) \quad (3.3.2)$$

$$\kappa(i, j) = \begin{cases} \sigma^{-2}\gamma_X(i - j), & 1 \leq i, j \leq m \\ \sigma^{-2} \left[\gamma_X(i - j) - \sum_{r=1}^p \phi_r \gamma_X(r - |i - j|) \right], & \min(i, j) \leq m < \max(i, j) \leq 2m, \\ \sum_{r=0}^q \theta_r \theta_{r+|i-j|}, & \min(i, j) > m, \\ 0 & \text{otherwise.} \end{cases} \quad (3.3.3)$$

Because all estimation methods use the above formula for $\hat{\sigma}^2$, variance estimates computed by ITSM-R are different from what is normally expected. For example, the Yule-Walker estimation function `yw` returns a white noise variance estimate that differs from the traditional Yule-Walker algorithm.

Since variance estimates are computed uniformly, model AICCs can always be compared directly, even when different methods are used to estimate model parameters. For example, it is perfectly valid to compare the AICC of a Yule-Walker model to that of a maximum likelihood model.

1.3 ARMA Coefficients

ARMA coefficient vectors are ordered such that the vector index corresponds to the lag of the coefficient. For example, the model

$$X_t - \phi_1 X_{t-1} - \phi_2 X_{t-2} = Z_t + \theta_1 Z_{t-1} + \theta_2 Z_{t-2}$$

with

$$\phi_1 = 1/2$$

$$\phi_2 = 1/3$$

$$\theta_1 = 1/4$$

$$\theta_2 = 1/5$$

corresponds to the following R vectors.

```
phi = c(1/2, 1/3)
theta = c(1/4, 1/5)
```

The equivalent model

$$X_t = \phi_1 X_{t-1} + \phi_2 X_{t-2} + Z_t + \theta_1 Z_{t-1} + \theta_2 Z_{t-2}$$

makes plain the sign convention of the AR coefficients.

1.4 ARIMA Models

ARIMA(p,d,q) models are specified by differencing d times with a lag of one each time. For example, the following R code specifies an ARIMA(1,2,3) model for data set `quant`.

```
M = c("diff",1,"diff",1)
e = Resid(quant,M)
a = arma(e,1,3)
```

1.5 Data Sets

ITSM-R includes the following data sets that are featured in *Introduction to Time Series and Forecasting*. Each data set is an ordinary vector (not a time series object).

Name	Obs.	Description
<code>airpass</code>	144	Number of international airline passengers, 1949 to 1960
<code>deaths</code>	72	USA accidental deaths, 1973 to 1978
<code>dowj</code>	78	Dow Jones utilities index, August 28 to December 18, 1972
<code>lake</code>	98	Level of Lake Huron, 1875 to 1972
<code>strikes</code>	30	USA union strikes, 1951 to 1980
<code>Sunspots</code>	100	Number of sunspots, 1770 to 1869
<code>wine</code>	142	Australian red wine sales, January 1980 to October 1991

2 Functions

2.1 aacvf

Computes the autocovariance of an ARMA model.

`aacvf(a,h)` $\begin{cases} \text{a} & \text{ARMA model} \\ \text{h} & \text{Maximum lag} \end{cases}$

The ARMA model is a list with the following components.

`$phi` AR coefficients $[1:p] = \phi_1, \dots, \phi_p$
`$theta` MA coefficients $[1:q] = \theta_1, \dots, \theta_q$
`$sigma2` White noise variance σ^2

Returns a vector of length `h+1` to accomodate lag 0 at index 1.

The following example is from p. 103 of *Introduction to Time Series and Forecasting*.

```
R> a = specify(ar=c(1,-0.24),ma=c(0.4,0.2,0.1))
R> aacvf(a,3)
[1] 7.171327 6.441393 5.060274 3.614340
```

2.2 acvf

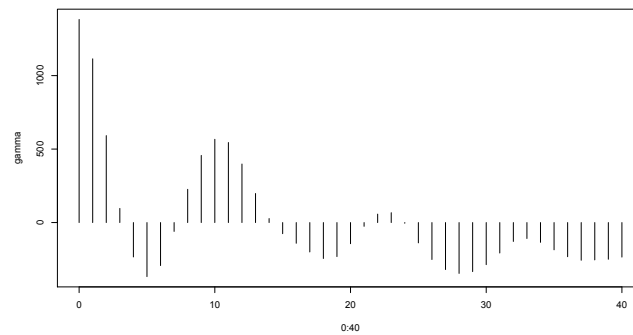
Computes the autocovariance of time series data.

`acvf(x,h=40)` $\begin{cases} \text{x} & \text{Time series data} \\ \text{h} & \text{Maximum lag} \end{cases}$

Returns a vector of length `h+1` to accommodate lag 0 at index 1.

Example:

```
R> gamma = acvf(Sunspots)
R> plot(0:40,gamma,type="h")
```



2.3 ar.inf

Returns the $\text{AR}(\infty)$ coefficients $\pi_0, \pi_1, \dots, \pi_n$ where $\pi_0 = 1$.

`ar.inf(a,n)` $\begin{cases} \text{a} & \text{ARMA model} \\ \text{n} & \text{Order} \end{cases}$

A vector of length `n+1` is returned to accommodate π_0 at index 1.

Example:

```
R> a = yw(Sunspots,2)
R> ar.inf(a,10)
[1] 1.0000000 -1.3175005 0.6341215 0.0000000 0.0000000 0.0000000
[7] 0.0000000 0.0000000 0.0000000 0.0000000 0.0000000
```

For invertible ARMA processes, $\text{AR}(\infty)$ is the polynomial $\pi(z)$ such that

$$\pi(z) = \frac{\phi(z)}{\theta(z)} = 1 + \pi_1 z + \pi_2 z^2 + \dots$$

The corresponding autoregression is

$$Z_t = \pi(B)X_t = X_t + \pi_1 X_{t-1} + \pi_2 X_{t-2} + \dots$$

The coefficients π_j are calculated recursively using the following formula. (See p. 86 of *Introduction to Time Series and Forecasting*.)

$$\pi_j = -\phi_j - \sum_{k=1}^q \theta_k \pi_{j-k} \quad j = 1, 2, \dots$$

2.4 arar

Predict future values of a time series using the ARAR algorithm.

`arar(x,h=10,opt=2)` $\left\{ \begin{array}{ll} \text{x} & \text{Observed data} \\ \text{h} & \text{Steps ahead} \\ \text{opt} & \text{Display option} \end{array} \right.$

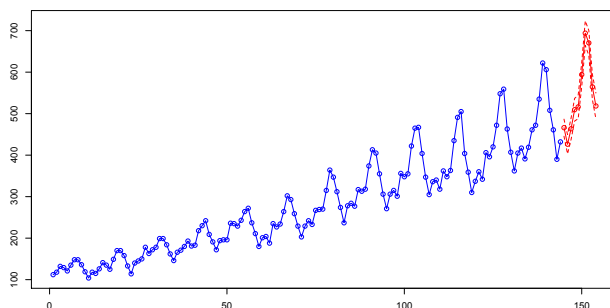
The display options are 0 for silent, 1 for tabulate, 2 for plot and tabulate. Returns the following list invisibly.

`$pred` Predicted values
`$se` Standard errors
`$l` Lower bounds
`$u` Upper bounds

Example:

```
R> arar(airpass)
Optimal lags 1 2 9 10
Optimal coeffs 0.5247184 0.2735903 0.2129203 -0.316453
WN Variance 110.1074
Filter 1 -0.5247184 -0.2735903 0 0 0 0 0 0 -0.2129203 0.316453 0
-1.114253 0.5846688 0.3048487 0 0 0 0 0 0 0.237247 -0.3526086
```

Step	Prediction	sqrt(MSE)	Lower Bound	Upper Bound
1	466.1915	10.49321	445.6248	486.7582
2	426.3592	11.85003	403.1331	449.5853
3	463.614	13.17574	437.7895	489.4384
4	509.5108	13.93231	482.2035	536.8182
5	516.2016	14.48202	487.8169	544.5864
6	594.0837	14.85609	564.9658	623.2017
7	693.9735	15.12129	664.3358	723.6112
8	670.4816	15.30835	640.4772	700.4859
9	564.4617	15.44148	534.1964	594.727
10	518.5135	15.93834	487.2743	549.7526



2.5 arma

Returns an ARMA model using maximum likelihood to estimate the AR and MA coefficients.

`arma(x,p=0,q=0)` $\left\{ \begin{array}{l} \text{x} \text{ Time series data} \\ \text{p} \text{ AR order} \\ \text{q} \text{ MA order} \end{array} \right.$

The R function `arma` is called to estimate ϕ and θ . The innovations algorithm is used to estimate the white noise variance σ^2 . The resulting ARMA model is a list with the following components.

<code>\$phi</code>	AR coefficients $[1:p] = \phi_1, \dots, \phi_p$
<code>\$theta</code>	MA coefficients $[1:q] = \theta_1, \dots, \theta_q$
<code>\$sigma2</code>	White noise variance σ^2
<code>\$aicc</code>	Akaike information criterion corrected
<code>\$se.phi</code>	Standard errors for the AR coefficients
<code>\$se.theta</code>	Standard errors for the MA coefficients

The signs of the coefficients correspond to the following model.

$$X_t = \sum_{j=1}^p \phi_j X_{t-j} + Z_t + \sum_{k=1}^q \theta_k Z_{t-k}$$

The following example estimates ARMA(1,0) parameters for a stationary transformation of the Dow Jones data.

```
R> M = c("diff",1)
R> e = Resid(dowj,M)
R> a = arma(e,1,0)
R> a
$phi
[1] 0.4478187

$theta
[1] 0

$sigma2
[1] 0.1455080

$aicc
[1] 74.48316

$se.phi
[1] 0.01105692
```

```
$se.theta  
[1] 0
```

2.6 autofit

Uses maximum likelihood to determine the best ARMA model given a range of models. The `autofit` function tries all combinations of argument sequences and returns the model with the lowest AICC. This is a wrapper function, the R function `arima` does the actual estimation.

`autofit(x,p=0:5,q=0:5)` $\left\{ \begin{array}{l} \text{x} \text{ Time series data} \\ \text{p} \text{ AR order} \\ \text{q} \text{ MA order} \end{array} \right.$

Returns a list with the following components.

<code>\$phi</code>	AR coefficients $[1:p] = \phi_1, \dots, \phi_p$
<code>\$theta</code>	MA coefficients $[1:q] = \theta_1, \dots, \theta_q$
<code>\$sigma2</code>	White noise variance σ^2
<code>\$aicc</code>	Akaike information criterion corrected
<code>\$se.phi</code>	Standard errors for the AR coefficients
<code>\$se.theta</code>	Standard errors for the MA coefficients

The signs of the coefficients correspond to the following model.

$$X_t = \sum_{j=1}^p \phi_j X_{t-j} + Z_t + \sum_{k=1}^q \theta_k Z_{t-k}$$

The following example shows that an ARMA(1,1) model has the lowest AICC for the Lake Huron data.

```
R> autofit(lake)
$phi
[1] 0.7448993

$theta
[1] 0.3205891

$sigma2
[1] 0.4750447

$aicc
[1] 212.7675

$se.phi
[1] 0.006029624

$se.theta
[1] 0.01288894
```

2.7 burg

Estimates AR coefficients using the Burg method.

$\text{burg}(\mathbf{x}, p) \quad \begin{cases} \mathbf{x} & \text{Time series data} \\ p & \text{AR order} \end{cases}$

Returns an ARMA model with the following components.

<code>\$phi</code>	AR coefficients $[1:p] = \phi_1, \dots, \phi_p$
<code>\$theta</code>	0
<code>\$sigma2</code>	White noise variance σ^2
<code>\$aicc</code>	Akaike information criterion corrected
<code>\$se.phi</code>	Standard errors for AR coefficients
<code>\$se.theta</code>	0

Example:

```
R> burg(lake,1)
```

```
$phi  
[1] 0.8388953
```

```
$theta  
[1] 0
```

```
$sigma2  
[1] 0.5096105
```

```
$aicc  
[1] 217.3922
```

```
$se.phi  
[1] 0.003023007
```

```
$se.theta  
[1] 0
```

2.8 check

Check for causality and invertibility of an ARMA model.

`check(a)` { a ARMA model

The ARMA model is a list with the following components.

`$phi` AR coefficients $[1:p] = \phi_1, \dots, \phi_p$
`$theta` MA coefficients $[1:q] = \theta_1, \dots, \theta_q$

Example:

```
R> a = specify(ar=c(0,0,0.99))
R> check(a)
Causal
Invertible
```

2.9 forecast

Predict future values of a time series.

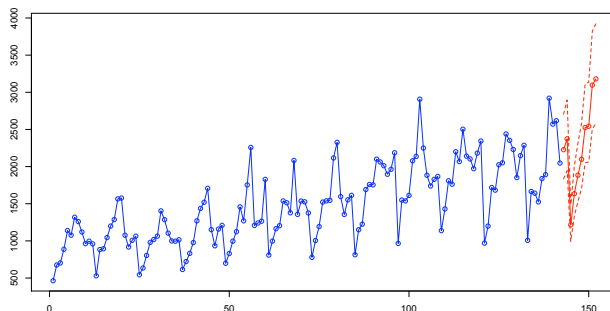
`forecast(x,M,a,h=10,opt=2,alpha=0.05)` $\left\{ \begin{array}{ll} \text{x} & \text{Time series data} \\ \text{M} & \text{Data model} \\ \text{a} & \text{ARMA model} \\ \text{h} & \text{Steps ahead} \\ \text{opt} & \text{Display option} \\ \text{alpha} & \text{Level of significance} \end{array} \right.$

The data model M can be NULL for none. The display options are 0 for none, 1 for tabulate, 2 for plot and tabulate. See below for details about the data model.

Example:

```
R> M = c("log","season",12,"trend",1)
R> e = Resid(wine,M)
R> a = arma(e,1,1)
R> forecast(wine,M,a)
```

Step	Prediction	Lower Bound	Upper Bound
1	2227.556	1834.156	2705.334
2	2374.062	1946.145	2896.069
3	1216.429	993.925	1488.743
4	1634.838	1332.583	2005.651
5	1883.996	1532.926	2315.467
6	2097.927	1704.717	2581.833
7	2524.942	2049.658	3110.437
8	2542.990	2062.775	3135.001
9	3096.280	2510.185	3819.219
10	3180.418	2577.324	3924.636



The data model M is a vector of function names. The functions are applied to the data in left to right order. There are five functions from which to choose.

<code>diff</code>	Difference the data
<code>hr</code>	Subtract harmonic components
<code>log</code>	Take the log of the data
<code>season</code>	Subtract seasonal component
<code>trend</code>	Subtract trend component

A function name may be followed by one or more arguments.

`diff` has a single argument, the lag.

`hr` has one or more arguments, each specifying the number of observations per harmonic period.

`log` has no arguments.

`season` has one argument, the number of observations per season.

`trend` has one argument, the polynomial order of the trend, (1 for linear, 2 for quadratic, etc.)

A data model is built up by concatenating the function names and arguments. For example, the following vector takes the log of the data, then subtracts a seasonal component of period twelve then subtracts a linear trend component.

```
R> M = c("log","season",12,"trend",1)
```

At the end of the data model there is an implied subtraction of the mean operation. Hence the resulting time series always has zero mean.

2.10 hannan

Estimates ARMA coefficients using the Hannan-Rissanen algorithm. It is required that $q > 0$.

$\text{hannan}(x, p, q) \begin{cases} x & \text{Time series data} \\ p & \text{AR order} \\ q & \text{MA order} \end{cases}$

Returns a list with the following components.

<code>\$phi</code>	AR coefficients $[1:p] = \phi_1, \dots, \phi_p$
<code>\$theta</code>	MA coefficients $[1:q] = \theta_1, \dots, \theta_q$
<code>\$sigma2</code>	White noise variance σ^2
<code>\$aicc</code>	Akaike information criterion corrected
<code>\$se.phi</code>	Standard errors for AR coefficients
<code>\$se.theta</code>	Standard errors for MA coefficients

Example:

```
R> hannan(lake, 1, 1)
```

```
$phi
```

```
[1] 0.6960772
```

```
$theta
```

```
[1] 0.3787969
```

```
$sigma2
```

```
[1] 0.477358
```

```
$aicc
```

```
[1] 213.183
```

```
$se.phi
```

```
[1] 0.07800321
```

```
$se.theta
```

```
[1] 0.1465265
```

2.11 hr

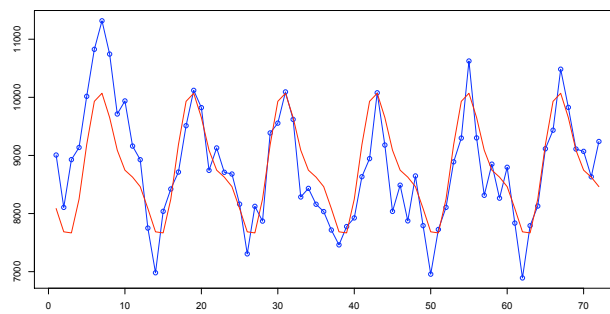
Returns the sum of harmonic components of time series data.

$$\text{hr}(x,d) \quad \begin{cases} x & \text{Time series data} \\ d & \text{Vector of harmonic periods} \end{cases}$$

Example:

```
R> y = hr(deaths,c(12,6))
```

```
R> plotc(deaths,y)
```



2.12 ia

Calculates MA coefficients using the innovations algorithm.

`ia(x,q,m=17)` $\left\{ \begin{array}{l} \text{x} \text{ Time series data} \\ \text{q} \text{ MA order} \\ \text{m} \text{ Recursion level} \end{array} \right.$

Returns a list with the following components.

<code>\$phi</code>	0
<code>\$theta</code>	MA coefficients $[1:q] = \theta_1, \dots, \theta_q$
<code>\$sigma2</code>	White noise variance σ^2
<code>\$aicc</code>	Akaike information criterion corrected
<code>\$se.phi</code>	0
<code>\$se.theta</code>	Standard errors for MA coefficients

Normally `m` should be set to the default 17 even when fewer MA coefficients are required.

The following example generates results that match *Introduction to Time Series and Forecasting* p. 154.

```
R> y = diff(dowj)
R> a = ia(y,17)
R> round(a$theta,4)
[1] 0.4269 0.2704 0.1183 0.1589 0.1355 0.1568 0.1284 -0.0060
[9] 0.0148 -0.0017 0.1974 -0.0463 0.2023 0.1285 -0.0213 -0.2575
[17] 0.0760
R> round(a$theta/a$se/1.96,4)
[1] 1.9114 1.1133 0.4727 0.6314 0.5331 0.6127 0.4969 -0.0231
[9] 0.0568 -0.0064 0.7594 -0.1757 0.7666 0.4801 -0.0792 -0.9563
[17] 0.2760
```

The formula for the standard error of the j th coefficient is given on p. 152 of *Introduction to Time Series and Forecasting* as

$$\sigma_j = n^{-1/2} \left[\sum_{i=0}^{j-1} \hat{\theta}_{mi}^2 \right]^{1/2}$$

where $\hat{\theta}_{m0} = 1$. Hence the standard error for $\hat{\theta}_1$ is $\sigma_1 = n^{-1/2}$.

2.13 `ma.inf`

Returns the $\text{MA}(\infty)$ coefficients $\psi_0, \psi_1, \dots, \psi_n$ where $\psi_0 = 1$.

`ma.inf(m,n)` $\begin{cases} \mathbf{a} & \text{ARMA model} \\ \mathbf{n} & \text{Order} \end{cases}$

A vector of length $\mathbf{n}+1$ is returned to accommodate ψ_0 at index 1.

Example:

```
R> a = yw(Sunspots,2)
R> ma.inf(a,10)
[1] 1.00000000 1.31750053 1.10168617 0.61601672 0.11299949
[6] -0.24175256 -0.39016452 -0.36074148 -0.22786538 -0.07145884
[11] 0.05034727
```

For causal ARMA processes, $\text{MA}(\infty)$ is the polynomial $\psi(z)$ such that

$$\psi(z) = \frac{\theta(z)}{\phi(z)} = 1 + \psi_1 z + \psi_2 z^2 + \dots$$

The corresponding moving average is

$$X_t = \psi(B)Z_t = Z_t + \psi_1 Z_{t-1} + \psi_2 Z_{t-2} + \dots$$

The coefficients ψ_j are calculated recursively using the following formula. (See p. 85 of *Introduction to Time Series and Forecasting*.)

$$\psi_j = \theta_j + \sum_{k=1}^p \phi_k \psi_{j-k} \quad j = 1, 2, \dots$$

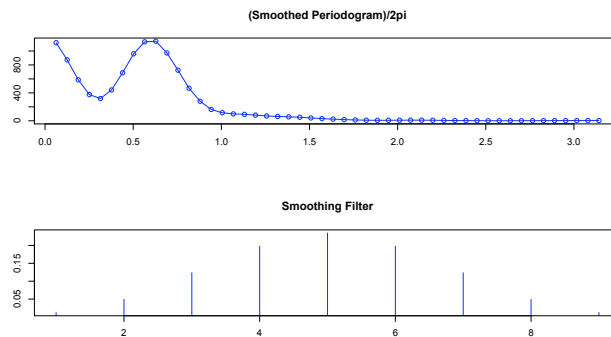
2.14 periodogram

Plots a periodogram.

`periodogram(x,q=0,opt=2)` $\left\{ \begin{array}{ll} x & \text{Time series data} \\ q & \text{MA filter order} \\ \text{opt} & \text{Plot option} \end{array} \right.$

The periodogram vector divided by 2π is returned invisibly. The plot options are 0 for no plot, 1 to plot only the periodogram, and 2 to plot both the periodogram and the filter coefficients. The filter `q` can be a vector in which case the overall filter is the composition of MA filters of the designated orders.

```
R> periodogram(Sunspots,c(1,1,1,1))
Filter  0.01234568 0.04938272 0.1234568 0.1975309 0.2345679 0.1975309 0.1234568
        0.04938272 0.01234568
```



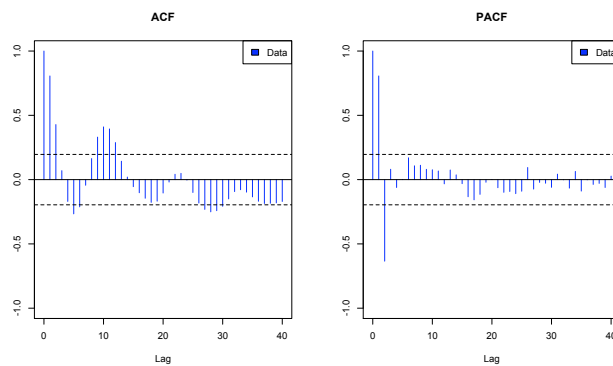
2.15 plota

Plots ACF and PACF for time series data and/or ARMA model.

`plota(u,v=NULL,h=40)` $\begin{cases} u, v & \text{Time series data and/or ARMA model, in either order} \\ h & \text{Maximum lag} \end{cases}$

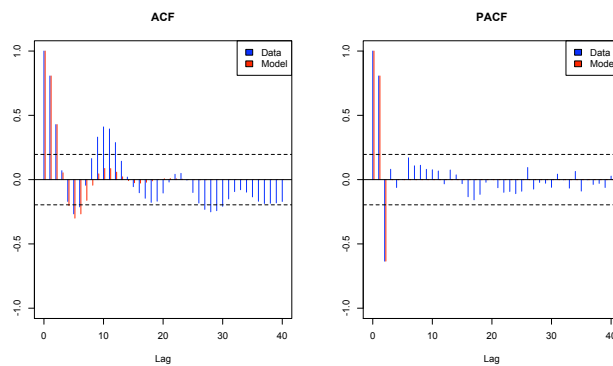
Example:

```
R> plota(Sunspots)
```



```
R> a = yw(Sunspots,2)
```

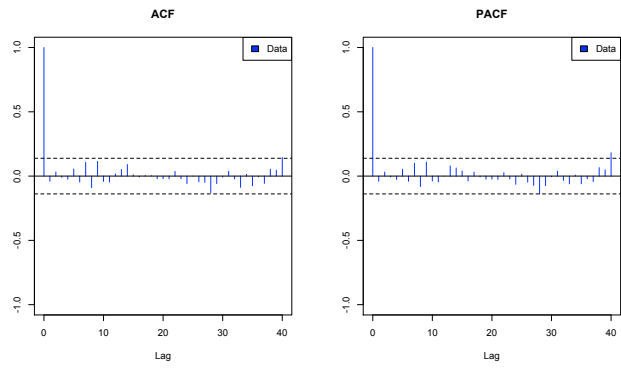
```
R> plota(Sunspots,a)
```



The following example demonstrates the utility of the $\pm 1.96/\sqrt{n}$ bounds as a test for noise.

```
R> noise = rnorm(200)
```

```
R> plota(noise)
```



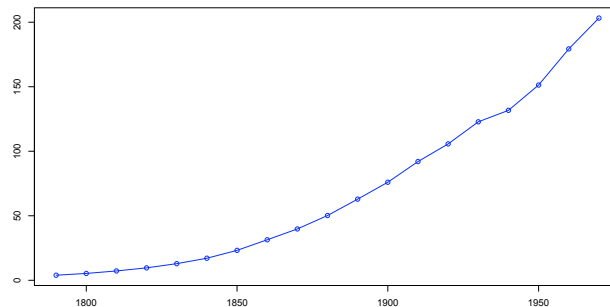
2.16 plotc

Plots one or two time series in color.

`plotc(y1,y2=NULL)` $\begin{cases} y1 & \text{Blue line with knots} \\ y2 & \text{Red line without knots} \end{cases}$

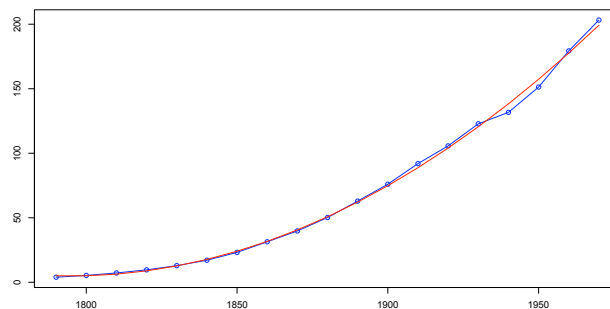
Example:

```
R> plotc(uspop)
```



```
R> y = trend(uspop,2)
```

```
R> plotc(uspop,y)
```



2.17 plots

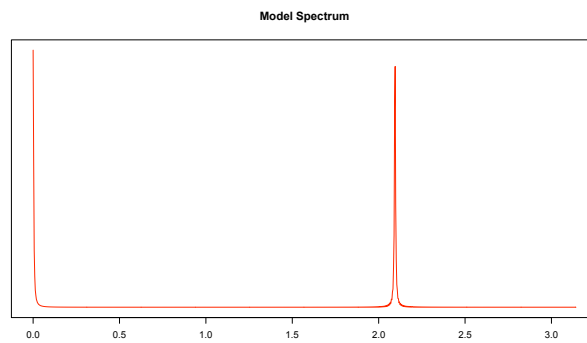
Plots the spectrum of data or an ARMA model.

`plots(u)` { `u` Time series data or ARMA model

Example:

```
R> a = specify(ar=c(0,0,0.99))
```

```
R> plots(a)
```



2.18 Resid

Returns the residuals of a time series model.

$$\text{Resid}(x, M=\text{NULL}, a=\text{NULL}) \quad \left\{ \begin{array}{ll} x & \text{Time series data} \\ M & \text{Data model} \\ a & \text{ARMA model} \end{array} \right.$$

Either **M** or **a** can be **NULL** for none. See below for details about the data model. The returned residuals always have zero mean.

In the following example, **Resid** and **test** are used to check for stationarity of the transformed data. Then they are used again to check that the overall model reduces the data to white noise.

```
R> M = c("log","season",12,"trend",1)
R> e = Resid(airpass,M)
R> test(e)
R> a = arma(e,1,0)
R> ee = Resid(airpass,M,a)
R> test(ee)
```

The data model **M** is a vector of function names. The functions are applied to the data in left to right order. There are five functions from which to choose.

diff	Difference the data
hr	Subtract harmonic components
log	Take the log of the data
season	Subtract seasonal component
trend	Subtract trend component

A function name may be followed by one or more arguments.

diff has a single argument, the lag.

hr has one or more arguments, each specifying the number of observations per harmonic period.

log has no arguments.

season has one argument, the number of observations per season.

trend has one argument, the polynomial order of the trend, (1 for linear, 2 for quadratic, etc.)

A data model is built up by concatenating the function names and arguments. For example, the following vector takes the log of the data, then subtracts a seasonal component of period twelve then subtracts a linear trend component.

```
> M = c("log","season",12,"trend",1)
```

At the end of the data model there is an implied subtraction of the mean operation. Hence the resulting time series always has zero mean.

2.19 season

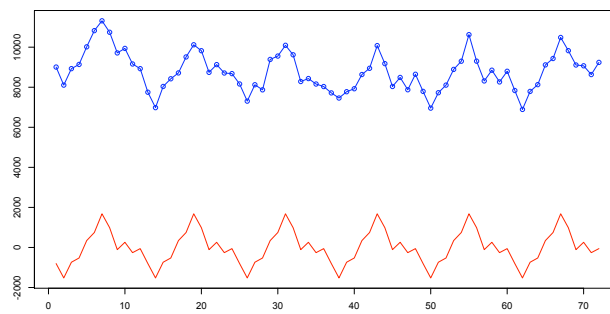
Returns the seasonal component of time series data.

`season(x,d)` $\begin{cases} x & \text{Time series data} \\ d & \text{Number of observations per season} \end{cases}$

Example:

```
R> s = season(deaths,12)
```

```
R> plotc(deaths,s)
```



2.20 `sim`

Generate synthetic observations for an ARMA model.

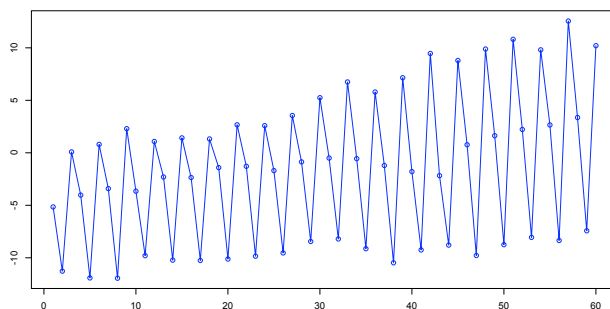
`sim(a,n)` $\begin{cases} \text{a} & \text{ARMA model} \\ \text{n} & \text{Number of synthetic observations required} \end{cases}$

The ARMA model is a list with the following components.

<code>\$phi</code>	AR coefficients ϕ_1, \dots, ϕ_p
<code>\$theta</code>	MA coefficients $\theta_1, \dots, \theta_q$
<code>\$sigma2</code>	White noise variance σ^2

Example:

```
R> a = specify(ar=c(0,0,0.99))
R> x = sim(a,60)
R> plotc(x)
```



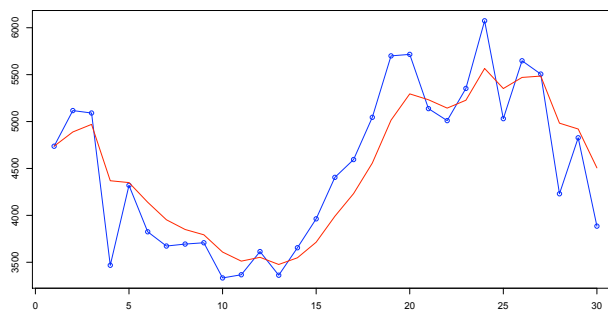
2.21 smooth.exp

Applies an exponential smoothing filter to data and returns the result.

`smooth.exp(x,a)` $\begin{cases} x & \text{Time series data} \\ a & 0 \text{ to } 1, \text{ zero is maximum smoothness.} \end{cases}$

Example:

```
R> y = smooth.exp(strikes,0.4)
R> plotc(strikes,y)
```



2.22 smooth.fft

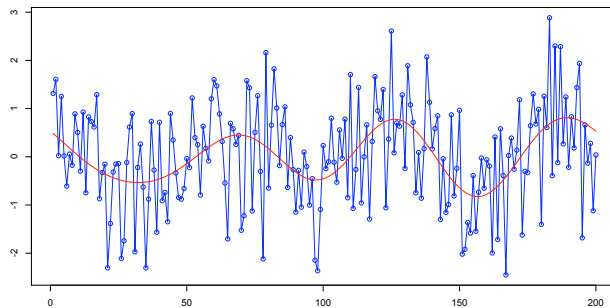
Applies a low pass filter to data and returns the result.

`smooth.fft(x,c)` $\begin{cases} \mathbf{x} & \text{Time series data} \\ \mathbf{c} & \text{Cut-off freq. 0-1} \end{cases}$

The cut-off frequency is specified as a fraction. For example, $c = 0.25$ passes the lowest 25% of the frequency spectrum.

Example:

```
R> y = smooth.fft(signal,0.035)
R> plotc(signal,y)
```



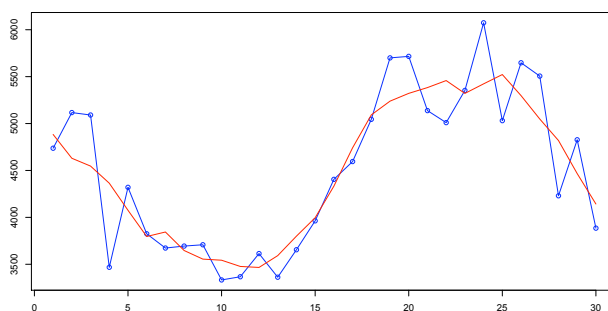
2.23 smooth.ma

Applies a moving average filter to data and returns the result.

`smooth.ma(x,q)` $\begin{cases} \mathbf{x} & \text{Time series data} \\ \mathbf{q} & \text{Filter order} \end{cases}$

Example:

```
R> y = smooth.ma(strikes,2)
R> plotc(strikes,y)
```



From p. 25 of *Introduction to Time Series and Forecasting*,

$$Y_t = \frac{1}{2q+1} \sum_{j=-q}^q X_{t-j}$$

Hence for $q = 2$ the moving average filter is

$$Y_t = \frac{1}{5}(X_{t-2} + X_{t-1} + X_t + X_{t+1} + X_{t+2})$$

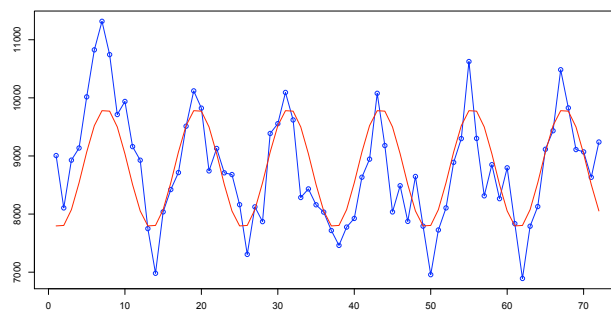
2.24 smooth.rank

Passes the mean and the k frequencies with the highest amplitude. The remainder of the spectrum is filtered out.

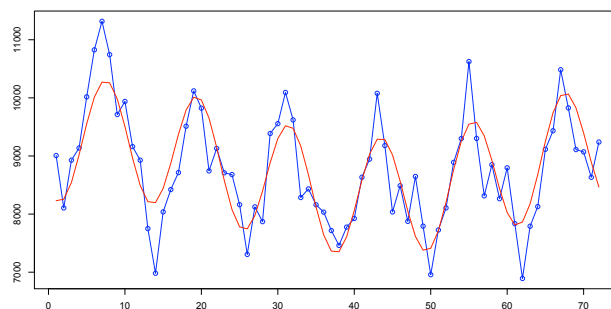
`smooth.rank(x,k)` $\begin{cases} x & \text{Time series data} \\ k & \text{Rank} \end{cases}$

Example:

```
R> y = smooth.rank(deaths,1)
R> plot(deaths,y)
```



```
R> y = smooth.rank(deaths,2)
R> plot(deaths,y)
```



2.25 specify

Returns an ARMA model with the specified parameters.

`specify(ar=0,ma=0,sigma2=1)` $\left\{ \begin{array}{ll} \text{ar} & \text{AR coefficients } [1:p] = \phi_1, \dots, \phi_p \\ \text{ma} & \text{MA coefficients } [1:q] = \theta_1, \dots, \theta_q \\ \text{sigma2} & \text{White noise variance } \sigma^2 \end{array} \right.$

The signs of the coefficients correspond to the following model.

$$X_t = \phi_1 X_{t-1} + \phi_2 X_{t-2} + \dots + Z_t + \theta_1 Z_{t-1} + \theta_2 Z_{t-2} + \dots$$

Example:

```
R> specify(ar=c(0,0,0.99))
```

```
$phi
```

```
[1] 0.00 0.00 0.99
```

```
$theta
```

```
[1] 0
```

```
$sigma2
```

```
[1] 1
```

2.26 test

Test residuals for randomness.

`test(e)` { e Residuals

Example:

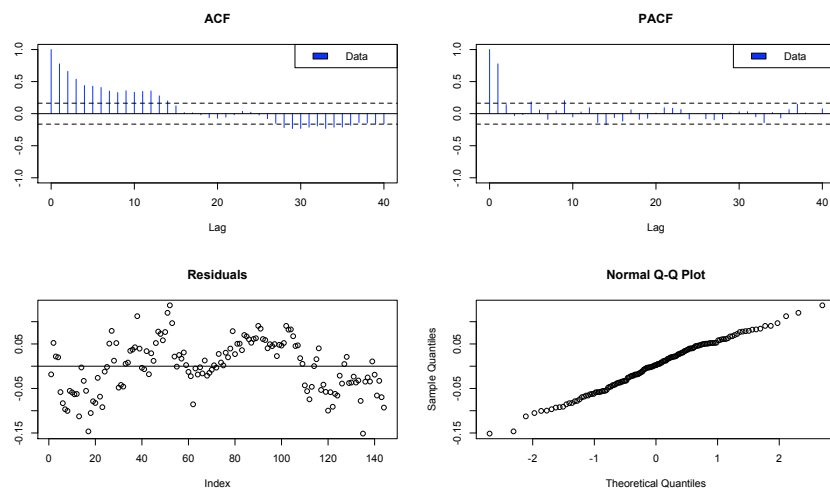
```
R> M = c("log","season",12,"trend",1)
```

```
R> e = Resid(airpass,M)
```

```
R> test(e)
```

Null hypothesis: Residuals are iid noise.

Test	Distribution	Statistic	p-value
Ljung-Box Q	$Q \sim \text{chisq}(20)$	412.43	0 *
McLeod-Li Q	$Q \sim \text{chisq}(20)$	41.29	0.0034 *
Turning points T	$(T-94.7)/5 \sim N(0,1)$	85	0.0545
Diff signs S	$(S-71.5)/3.5 \sim N(0,1)$	68	0.314
Rank P	$(P-5148)/289.5 \sim N(0,1)$	5187	0.8928



2.27 trend

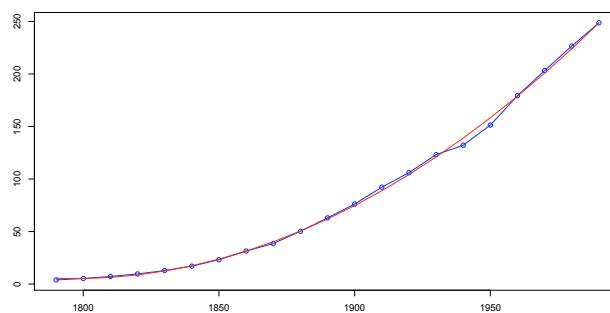
Returns the trend component of time series data.

`trend(x,p)` $\begin{cases} x & \text{Time series data} \\ n & \text{Polynomial order} \end{cases}$

Example:

```
R> y = trend(uspop,2)
```

```
R> plotc(uspop,y)
```



2.28 yw

Estimates AR coefficients using the Yule-Walker method.

$yw(x, p) \begin{cases} x & \text{Time series data} \\ p & \text{AR order} \end{cases}$

Returns an ARMA model with the following components.

<code>\$phi</code>	AR coefficients $[1:p] = \phi_1, \dots, \phi_p$
<code>\$theta</code>	0
<code>\$sigma2</code>	White noise variance σ^2
<code>\$aicc</code>	Akaike information criterion corrected
<code>\$se.phi</code>	Standard errors for AR coefficients
<code>\$se.theta</code>	0

Example:

```
R> yw(lake,1)
```

```
$phi  
[1] 0.8319112
```

```
$theta  
[1] 0
```

```
$sigma2  
[1] 0.5098608
```

```
$aicc  
[1] 217.4017
```

```
$se.phi  
[1] 0.003207539
```

```
$se.theta  
[1] 0
```

These are the relevant formulas from *Introduction to Time Series and Forecasting, Second Edition*.

$$\hat{\gamma}(h) = \frac{1}{n} \sum_{t=1}^{n-|h|} (X_{t+|h|} - \bar{X}_n) (X_t - \bar{X}_n) \quad (2.4.4)$$

$$\hat{\Gamma}_k = \begin{pmatrix} \hat{\gamma}(0) & \hat{\gamma}(1) & \cdots & \hat{\gamma}(k-1) \\ \hat{\gamma}(1) & \hat{\gamma}(0) & \cdots & \hat{\gamma}(k-2) \\ \vdots & \vdots & \ddots & \vdots \\ \hat{\gamma}(k-1) & \hat{\gamma}(k-2) & \cdots & \hat{\gamma}(0) \end{pmatrix} \quad (2.4.6)$$

$$\hat{R}_k = \frac{1}{\hat{\gamma}(0)} \hat{\Gamma}_k \quad (2.4.8)$$

$$\hat{\phi}_p = \hat{R}_p^{-1} \hat{\rho}_p \quad (5.1.7)$$

$$\hat{v}_p = \hat{\gamma}(0) \left(1 - \hat{\rho}_p' \hat{R}_p^{-1} \hat{\rho}_p \right) \quad (5.1.11)$$

where $\hat{\rho}_p = (\hat{\rho}(1), \dots, \hat{\rho}(p))' = \hat{\gamma}_p / \hat{\gamma}(0)$. The subscript p is the order of the AR model (i.e., number of AR coefficients). Note that (5.1.7) and (5.1.11) can also be written as

$$\hat{\phi}_p = \hat{\Gamma}_p^{-1} \hat{\gamma}_p$$

and

$$\hat{v}_p = \hat{\gamma}(0) - \hat{\gamma}_p' \hat{\phi}_p$$

From equation (5.1.13) the standard error for the j th AR coefficient is

$$SE_j = \sqrt{\frac{\hat{v}_{jj}}{n}}$$

where \hat{v}_{jj} is the j th diagonal element of $\hat{v}_p \hat{\Gamma}_p^{-1}$.

The R code follows immediately from the above equations.

```
gamma = acvf(x,p)
Gamma = toeplitz(gamma[1:p])
phi = solve(Gamma, gamma[2:(p+1)])
v = gamma[1] - drop(crossprod(gamma[2:(p+1)], phi))
V = v * solve(Gamma)
se.phi = sqrt(1/n*diag(V))
```

Recall that R indices are 1-based hence `gamma[1]` is $\hat{\gamma}(0)$. The function `crossprod` returns a 1-by-1 matrix which is then converted by `drop` to a scalar.