# Time Series: A First Course with Bootstrap Starter

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## Lesson 9-1: Weak Dependence

- We can quantify serial dependence through the ACVF.
- Recall entropy mixing (Paradigm 8.2.11) as another measure of serial dependence.

## Example 9.1.1. Slow Polynomial Decay – Long Memory

- Suppose that a time series has ACVF  $\gamma(h) = O(|h|^{-a})$  for  $0 < a \le 1$ .
- The notation means the ACVF is bounded by a constant times that rate.
- There is fairly slow decay in lag when a is small: this means persistence, or high association, across large time lags. This is called *long-range dependence* or *long memory*.

## Exercise 9.2. Long-Range Dependence

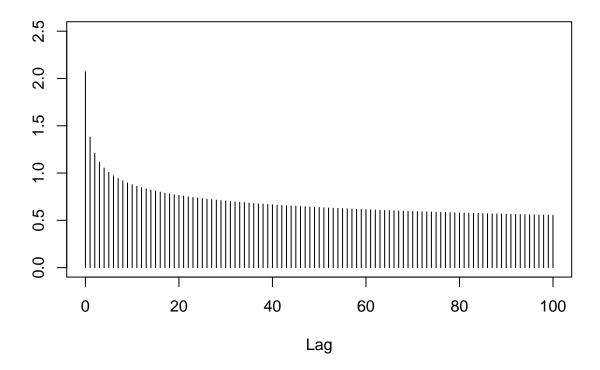
• An example of long-range dependence is given by the ACVF recursively defined via

$$\gamma(h+1) = \frac{h+d}{h+1-d}\,\gamma(h)$$

for  $d \in (0, 1/2)$ , where  $\Gamma$  denotes the gamma function, and  $\gamma(0) = \Gamma(1 - 2d)/\Gamma(1 - d)^2$ .

- This corresponds to  $\gamma(h) = O(|h|^{2d-1})$ , or the case of long memory with decay rate a = 1 2d.
- We plot with d = .4.

```
lags <- seq(1,100)
d <- .4
gamma.lm <- gamma(1-2*d)/(gamma(1-d))^2
for(i in 1:length(lags))
{
    gamma.new <- gamma.lm[i] * (i-1+d)/(i-d)
    gamma.lm <- c(gamma.lm,gamma.new)
}
plot(ts(gamma.lm,start=0),type="h",xlab="Lag",ylab="",ylim=c(0,2.5))</pre>
```



#### Example 9.1.3. Geometric Decay

- Suppose that a time series has ACVF  $\gamma(h) = O(r^{|h|})$  for 0 < r < 1.
- The notation means the ACVF is bounded by a constant times that rate.
- So there is a geometric, or exponential, rate of decay for the ACVF.
- This holds for ARMA processes by Proposition 5.8.3.

#### Example 9.1.4. MA(q) Correlation

- Suppose the a time series has ACVF  $\gamma(h)$  that is zero if |h| > q.
- Then this can be represented as an MA(q) process.
- This is an example of a q-dependent process:  $X_t$  and  $X_{t-h}$  are independent for any t when |h| > q.

## Lesson 9-2: Sample Mean

- We need to know the statistical properties of common time series estimators.
- We here focus on the sample mean, which estimates the population mean of a stationary time series.
- The sample mean is

$$\overline{X} = n^{-1} \sum_{t=1}^{n} X_t.$$

### Remark 9.2.1. The Long-Run Variance

• Suppose  $\{X_t\}$  is stationary and the autocovariance function (ACVF) is absolutely summable:

$$\sum_{h=-\infty}^{\infty} |\gamma(h)| < \infty.$$

• Then the long-run variance is defined as

$$\sum_{h=-\infty}^{\infty} \gamma(h).$$

- This is denoted by  $\sigma_{\infty}^2$ . Observe this is equal to G(1), i.e., setting z=1 in the AGF.
- The long-run variance can be zero: suppose  $\{X_t\}$  has a MA representation  $X_t = \psi(B)Z_t$ , so that

$$\sigma_{\infty}^2 = G(1) = \psi(1)^2 \sigma^2.$$

Then if  $\psi(1) = 0$ , the long-run variance is zero. For example, consider an MA(1) process with  $\theta(z) = 1 - z$ .

#### Proposition 9.2.2.

Suppose  $\{X_t\}$  is stationary with mean  $\mu$ , and ACVF that is absolutely summable. Then  $\overline{X}$  is unbiased for  $\mu$ , and its scaled variance tends to the long-run variance:

$$\operatorname{Var}[\sqrt{nX}] \to \sigma_{\infty}^2$$
.

#### Theorem 9.2.7.

- So long as the long-run variance is non-zero, we can establish a central limit theorem for the sample mean under the condition that the inputs (in the MA representation) are i.i.d.
- Suppose  $\{X_t\}$  has causal MA representation

$$X_t = \mu + \psi(B)Z_t,$$

where  $Z_t \sim i.i.d.(0, \sigma^2)$ . Suppose  $\sum_{j>0} j |\psi_j| < \infty$  and  $\psi(1) \neq 0$ . Then

$$\sqrt{n}(\overline{X} - \mu) \Rightarrow \mathcal{N}(0, \sigma_{\infty}^2).$$

#### Remark 9.2.8. Inference for the Mean

As an application, suppose we want to construct a confidence interval for the mean based on our estimator  $\overline{X}$ . To get the standard error, we need to know the long-run variance. This suggests the need to estimate all the autocovariances (however, that is not possible).

#### Exercise 9.13. Simulating an AR(1) Limiting Variance

• We simulate a Gaussian AR(1) process with parameter  $\phi_1 = .8$  and  $\sigma^2 = 1$ .

```
armapq.sim <- function(n,burn,ar.coefs,ma.coefs,innovar)
{
    p <- length(ar.coefs)
    q <- length(ma.coefs)
    z <- rnorm(n+burn+p+q,sd=sqrt(innovar))
    x <- filter(z,c(1,ma.coefs),method="convolution",sides=1)
    x <- x[(q+1):(q+n+burn+p)]
    y <- x[1:p]</pre>
```

```
for(t in (p+1):(p+n+burn))
{
    next.y <- sum(ar.coefs*y[(t-1):(t-p)]) + x[t]
    y <- c(y,next.y)
}
y <- y[(p+burn+1):(p+burn+n)]
return(y)
}
phi <- .8</pre>
```

```
• Based on sample sizes n = 50, 100, 200, we compute the sample mean.
   • Repeating over 10,000 simulations, we approximate the scaled variance of the sample mean, and see
     how close this is to the long-run variance.
# small sample
n <- 50
x.means <- NULL
for(i in 1:10000)
{
    x.sim <- armapq.sim(n,500,phi,NULL,1)</pre>
    x.means <- c(x.means,mean(x.sim))</pre>
}
y <- sqrt(n)*x.means
var(y)
## [1] 22.99445
# long-run variance for an AR(1)
(1-phi)^{-2}
## [1] 25
# moderate sample
n <- 100
x.means <- NULL
for(i in 1:10000)
    x.sim <- armapq.sim(n,500,phi,NULL,1)</pre>
    x.means <- c(x.means, mean(x.sim))</pre>
}
y <- sqrt(n)*x.means
var(y)
## [1] 23.15012
# long-run variance for an AR(1)
(1-phi)^{-2}
## [1] 25
# large sample
n <- 200
x.means <- NULL
for(i in 1:10000)
{
    x.sim <- armapq.sim(n,500,phi,NULL,1)</pre>
    x.means <- c(x.means,mean(x.sim))</pre>
```

```
}
y <- sqrt(n)*x.means
var(y)

## [1] 25.02043

# long-run variance for an AR(1)
(1-phi)^{-2}</pre>
```

## [1] 25

## Lesson 9-4: Serial Correlation

We want to estimate the ACVF: they are used in prediction, and are helpful in understanding serial dependence in the time series.

#### Remark 9.4.1. ACVF Estimator for Known Mean

- Suppose that the mean  $\mu$  of the stationary time series  $\{X_t\}$  was known, and we want to estimate  $\gamma(h)$ .
- This is the expectation of  $Y_t$ , where

$$Y_t = (X_{t+h} - \mu)(X_t - \mu).$$

- We can compute  $Y_1, \ldots, Y_{n-h}$  for any  $0 \le h < n$ , because  $\mu$  is known.
- The sample mean of these would be

$$\frac{1}{n-h}\sum_{t=1}^{n-h}Y_t = \frac{1}{n-h}\sum_{t=1}^{n-h}(X_{t+h} - \mu)(X_t - \mu).$$

This has expectation  $\gamma(h)$ .

- The long-run variance of this sample mean is based on the ACVF of  $\{Y_t\}$ , and has a complicated expression known as *Bartlett's Formula*.
- This suggests an estimator

$$\overline{\gamma}(h) = \frac{1}{n-h} \sum_{t=1}^{n-h} (X_{t+h} - \mu)(X_t - \mu),$$

which is unbiased for  $\gamma(h)$ .

#### Proposition 9.4.3. Bartlett's Formula

Suppose  $\{X_t\}$  is a causal linear time series with i.i.d. inputs  $\{Z_t\}$ , which have variance  $\sigma^2$  and kurtosis  $\eta$ . Then the long-run variance of  $\overline{\gamma}(h)$  is

$$\tau_{\infty}^{2} = \sum_{k=-\infty}^{\infty} \left( \gamma(k+h)\gamma(k-h) + \gamma(k)^{2} \right) + \gamma(h)^{2} (\eta - 3).$$

#### Remark 9.4.4. Limiting Variance of the ACVF Estimator

• Recall that the Fourier coefficients of a spectral density are denoted

$$\langle f \rangle_k = \frac{1}{2\pi} \int_{-\pi}^{\pi} f(\lambda) e^{i\lambda k} d\lambda.$$

• For a stationary Gaussian process with spectral density f, then Bartlett's Formula is

$$\tau_{\infty}^2 = \langle f^2 \rangle_{2k} + \langle f^2 \rangle_0.$$

• Example 1: suppose k=1 and the process is a white noise (with variance  $\sigma^2$ ). Then

$$\tau_{\infty}^2 = \langle \sigma^4 \rangle_2 + \langle \sigma^4 \rangle_0 = \sigma^4.$$

• Example 2: suppose k=1 and the process is a MA(1). So  $f(\lambda)=\sigma^2|1+\theta e^{-i\lambda}|^2$ , and

$$\tau_{\infty}^{2} = \sigma^{4} \langle |1 + \theta e^{-i\lambda}|^{4} \rangle_{2} + \sigma^{4} \langle |1 + \theta e^{-i\lambda}|^{2} \rangle_{0} = \sigma^{4} \left( 1 + 5\theta^{2} + \theta^{4} \right).$$

#### Remark 9.4.6. ACVF Estimator for Unknown Mean

- Since  $\mu$  is usually unknown, we can replace it by the sample mean  $\overline{X}$ .
- The resulting estimator is denoted  $\widetilde{\gamma}(h)$ .
- This estimator is asymptotically normal when the process  $\{X_t\}$  is m-dependent (i.e., random variables of at least lag m between them are independent).

## Lesson 9-5: Sample Autocovariance

• We renormalize the ACVF estimator via tapering.

#### Remark 9.5.2. Tapering the ACVF to Reduce Variability in High Lags

- The estimator  $\tilde{\gamma}(h)$  is approximately unbiased, but has high variance for large h.
- We can replace the n-h divisor by n, in order to decrease variance for larger h.
- This is like multiplying  $\tilde{\gamma}(h)$  by 1 h/n, which is an example of a taper.

#### Definition 9.5.3.

- An autocovariance taper is a bounded, even real-valued function  $\Lambda$  on [-1,1], such that  $\Lambda(0)=1$  and  $\Lambda(x)<1$ .
- We multiply the ACVF estimator at lag h by  $\Lambda(h/n)$ , where  $\Lambda(x) = 1 |x|$ .

#### Example 9.5.4. Bartlett Taper

- The Bartlett taper is  $\Lambda(x) = 1 |x|$ .
- The resulting estimator is

$$\widehat{\gamma}(h) = \Lambda(h/n)\widetilde{\gamma}(h) = \frac{1}{n} \sum_{t=1}^{n-|h|} (X_{t+|h|} - \overline{X})(X_t - \overline{X}).$$

- This is called the sample autocovariance function (sample ACVF)
- Under m-dependence, Corollary 9.5.8 establishes a central limit theorem for the sample ACVF:

$$\sqrt{n}(\widehat{\gamma}(h) - \gamma(h)) \Rightarrow \mathcal{N}(0, \tau_{\infty}^2).$$

## Exercise 9.28. Sample ACVF of ARMA(1,2) Process

• We load some functions needed to compute the true ACVF.

```
polymult <- function(a,b) {
bb <- c(b,rep(0,length(a)-1))
B <- toeplitz(bb)
B[lower.tri(B)] <- 0
aa <- rev(c(a,rep(0,length(b)-1)))
prod <- B %*% matrix(aa,length(aa),1)
return(rev(prod[,1]))</pre>
```

```
}
ARMAauto <- function(phi,theta,maxlag)
    p <- length(phi)</pre>
    q <- length(theta)</pre>
    gamMA <- polymult(c(1,theta),rev(c(1,theta)))</pre>
    gamMA \leftarrow gamMA[(q+1):(2*q+1)]
    if (p > 0)
         Amat <- matrix(0,nrow=(p+1),ncol=(2*p+1))</pre>
         for(i in 1:(p+1))
             Amat[i,i:(i+p)] <- c(-1*rev(phi),1)</pre>
         }
         Amat \leftarrow cbind(Amat[,(p+1)],as.matrix(Amat[,(p+2):(2*p+1)]) +
             t(matrix(apply(t(matrix(Amat[,1:p],p+1,p)),2,rev),p,p+1)))
         Bmat \leftarrow matrix(0,nrow=(q+1),ncol=(p+q+1))
         for(i in 1:(q+1))
         {
             Bmat[i,i:(i+p)] \leftarrow c(-1*rev(phi),1)
         Bmat <- t(matrix(apply(t(Bmat),2,rev),p+q+1,q+1))</pre>
         Bmat <- matrix(apply(Bmat,2,rev),q+1,p+q+1)</pre>
         Bmat <- Bmat[,1:(q+1)]</pre>
        Binv <- solve(Bmat)</pre>
         gamMix <- Binv %*% gamMA
         if (p <= q) { gamMix <- matrix(gamMix[1:(p+1),],p+1,1)</pre>
             } else gamMix <- matrix(c(gamMix,rep(0,(p-q))),p+1,1)</pre>
         gamARMA <- solve(Amat) %*% gamMix</pre>
    } else gamARMA <- gamMA[1]</pre>
    gamMA <- as.vector(gamMA)</pre>
    if (maxlag <= q) gamMA <- gamMA[1:(maxlag+1)] else gamMA <- c(gamMA,rep(0,(maxlag-q)))
    gamARMA <- as.vector(gamARMA)</pre>
    if (maxlag <= p) gamARMA <- gamARMA[1:(maxlag+1)] else {</pre>
    for(k in 1:(maxlag-p))
    {
         len <- length(gamARMA)</pre>
         acf <- gamMA[p+1+k]</pre>
         if (p > 0) acf <- acf + sum(phi*rev(gamARMA[(len-p+1):len]))</pre>
        gamARMA <- c(gamARMA,acf)</pre>
    } }
    return(gamARMA)
}
```

• We simulate a Gaussian ARMA(1,2) process of length n = 200.

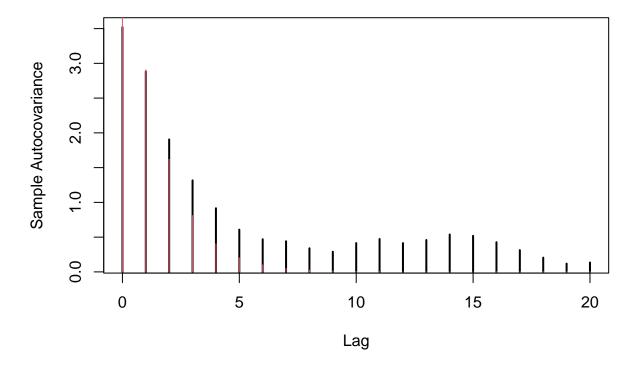
```
armapq.sim <- function(n,burn,ar.coefs,ma.coefs,innovar)
{
    p <- length(ar.coefs)
    q <- length(ma.coefs)
    z <- rnorm(n+burn+p+q,sd=sqrt(innovar))
    x <- filter(z,c(1,ma.coefs),method="convolution",sides=1)</pre>
```

```
x <- x[(q+1):(q+n+burn+p)]
y <- x[1:p]
for(t in (p+1):(p+n+burn))
{
    next.y <- sum(ar.coefs*y[(t-1):(t-p)]) + x[t]
    y <- c(y,next.y)
}
y <- y[(p+burn+1):(p+burn+n)]
return(y)
}

n <- 200
phi1 <- .5
theta1 <- 5/6
theta2 <- 1/6
x.sim <- armapq.sim(n,500,phi1,c(theta1,theta2),1)</pre>
```

- We construct and plot the sample acvf.
- We also overlay the true acvf in red.

```
y.sim <- x.sim - mean(x.sim)
x.acf <- mean(y.sim^2)
for(h in 1:20)
{
     x.acf <- c(x.acf,sum(y.sim[1:(n-h)]*y.sim[(h+1):n])/n)
}
gamma <- ARMAauto(phi1,c(theta1,theta2),21)
plot(ts(x.acf,start=0,frequency=1),xlab="Lag",ylab="Sample Autocovariance",type="h",lwd=2)
lines(ts(gamma,start=0,frequency=1),type="h",col=2)</pre>
```



# Lesson 9-6: Spectral Means

• We investigate statistics that are weighted sums of the periodogram.

## Proposition 9.6.1.

• The Fourier coefficients of the periodogram are the sample ACVF:

$$I(\lambda) = \sum_{k=-\infty}^{\infty} \widehat{\gamma}(k)e^{-i\lambda k},$$

and  $\langle I\rangle_k=\widehat{\gamma}(k).$  • So the sample ACVF is positive definite.

#### Definition 9.6.2.

A spectral mean is a functional of the spectral density f, of the form

$$\langle gf \rangle_0 = \frac{1}{2\pi} \int_{-\pi}^{\pi} g(\lambda) f(\lambda) d\lambda,$$

where g is a real-valued weighting function.

#### Remark 9.6.3. Spectral Mean Estimation

A spectral mean can be estimated by substituting the periodogram:

$$\langle gI\rangle_0 = \sum_{|h| < n} \langle g\rangle_h \widehat{\gamma}(h).$$

#### Theorem 9.6.6.

Suppose  $\{X_t\}$  is a linear time series with mean  $\mu$  and causal moving average representation, with inputs that have variance  $\sigma^2$  and kurtosis  $\eta$  (fourth moment divided by the square of the second moment). For sufficiently smooth g,

$$\sqrt{n} \left( \langle gI \rangle_0 - \langle gf \rangle_0 \right) \Rightarrow \mathcal{N} \left( 0, \langle g(g+g^{\sharp})f^2 \rangle_0 + (\eta-3)\langle gf \rangle_0^2 \right),$$

where  $g^{\sharp}(\lambda) = g(-\lambda)$ .

#### Remark 9.6.7. Autocovariance Limiting Variance

- If  $g(\lambda) = \cos(\lambda k)$ , then the spectral mean is  $\gamma(k)$ , and the estimate is  $\widehat{\gamma}(k)$ .
- The Bartlett Formula (Proposition 9.4.3) follows from Theorem 9.6.6.

#### Corollary 9.6.9. Ratio Statistics

Under same assumptions as Theorem 9.6.6, with smooth weighting functions a and b,

$$\sqrt{n} \left( \frac{\langle bI \rangle_0}{\langle aI \rangle_0} - \frac{\langle bf \rangle_0}{\langle af \rangle_0} \right) \Rightarrow \mathcal{N} \left( 0, \frac{\langle g(g+g^{\sharp})f^2 \rangle_0}{\langle af \rangle_0^2} \right),$$

where  $g = b - a\langle bf \rangle_0 / \langle af \rangle_0$ .

#### Remark 9.6.10. Bartlett's Formula for the Autocorrelations

- The sample autocorrelation is defined as  $\widehat{\rho}(k) = \widehat{\gamma}(k)/\widehat{\gamma}(0)$ .
- Applying Corollary 9.6.9 with  $b(\lambda) = \cos(\lambda k)$  and  $a \equiv 1$ , we find the asymptotic variance for the sample autocorrelation is

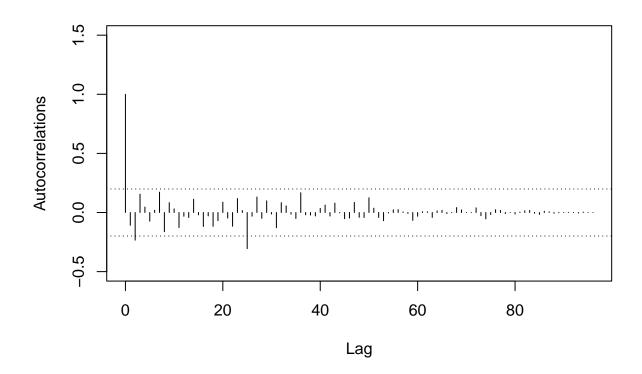
$$\frac{\langle f^2 \rangle_{2k}}{\langle f \rangle_0^2} + (1 + 2\rho(k)^2) \frac{\langle f^2 \rangle_0}{\langle f \rangle_0^2} - 4\rho(k) \frac{\langle f^2 \rangle_k}{\langle f \rangle_0^2}.$$

• For white noise, this equals 1.

#### Remark 9.6.12. Autocorrelations of Reduced Population Data

- Consider second differences of U.S. population data.
- We compute the sample autocorrelations, and test the hypothesis of zero serial correlation (i.e., white noise).
- For each lag  $k \ge 1$  (treated separately, ignoring multiple testing...) the asymptotic 95% critical values are  $\pm 1.96/\sqrt{n}$ .

```
pop <- read.table("USpop.dat")
pop <- ts(pop, start = 1901)
diffdiff.pop <- diff(diff(pop*10e-6))
data <- diffdiff.pop
n <- length(data)
acfs.sample <- NULL
mu.hat <- sum(data)/n
for(k in 0:(n-1))
{
        acf.sample <- sum((data[1:(n-k)]-mu.hat)*(data[(k+1):n]-mu.hat))/n
        acfs.sample <- c(acfs.sample,acf.sample)
}
plot(ts(acfs.sample/acfs.sample[1],start=0),xlab="Lag",ylab="Autocorrelations",ylim=c(-.5,1.5),type="h"
abline(h= 1.96/sqrt(n),lty=3)
abline(h= -1.96/sqrt(n),lty=3)</pre>
```



## Lesson 9-7: Periodogram

- We further investigate the periodogram.
- It can be viewed as a crude estimator of the spectral density.

#### Definition 9.7.2

• Given a sample of size n from a time series  $\{X_t\}$ , the centered DFT is

$$\widehat{X}(\lambda) = n^{-1/2} \sum_{t=1}^{n} (X_t - \overline{X}) e^{-i\lambda t}$$

for  $\lambda \in [-\pi, \pi]$ .

• So  $|\widehat{X}(\lambda)|^2 = I(\lambda)$ .

## Corollary 9.7.5

Suppose  $\{X_t\}$  is either *m*-dependent or is a linear time series with mean  $\mu$  and causal moving average representation. If  $\mu = 0$  then

$$\frac{\widetilde{I}(\lambda)}{f(\lambda)} \Rightarrow \begin{cases} \chi_1^2 & \text{if } |\lambda| = 0, \pi \\ .5\chi_2^2 & \text{if } |\lambda| \in (0, \pi). \end{cases}$$

If  $\mu \neq 0$ , we use the centered DFT and  $I(\lambda)/f(\lambda)$  has the same limit if  $\lambda \neq 0$ .

#### Remark 9.7.6. The Periodogram is Inconsistent

- Corollary 9.7.5 implies the periodogram is not consistent as an estimator of the spectral density.
- In contrast, Theorem 9.6.6 shows that weighted averages of the periodogram are consistent for spectral means.

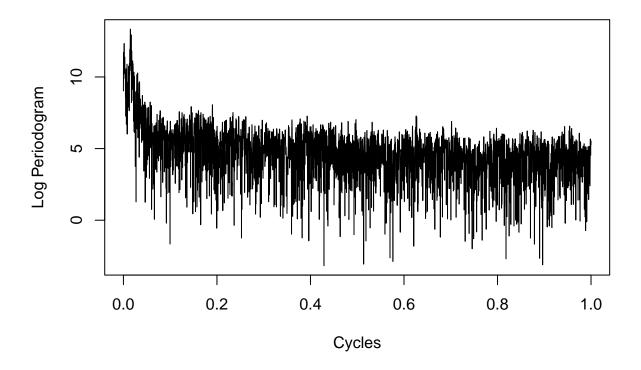
#### Fact 9.7.7. Independence of Periodogram Ordinates over Fourier Frequencies

- From Corollary 7.2.9, DFT ordinates are approximately uncorrelated.
- Because they are also asymptotically complex Gaussian (see book), the DFT ordinates are asymptotically independent.
- Hence  $I(\lambda_i)$  is asymptotically independent of  $I(\lambda_k)$  for  $j \neq k$ .

## Example 9.7.9. Periodogram of the Wolfer Sunspots

- We can plot the periodogram to identify cycles in a time series.
- We plot the periodogram of the Wolfer sunspots (recall Exercise 7.18).
- We omit the value I(0) = 0 so that we can plot in logs.
- The maximum occurs at  $\lambda = .047$ , corresponding to a period of 133.33 months, or 11.08 years.

```
wolfer <- read.table("wolfer.dat")</pre>
wolfer <- ts(wolfer[,1],start=1749,frequency=12)</pre>
data <- wolfer
n <- length(data)</pre>
acfs.sample <- NULL
mu.hat <- sum(data)/n
for(k in 0:(n-1))
{
    acf.sample \leftarrow sum((data[1:(n-k)]-mu.hat)*(data[(k+1):n]-mu.hat))/n
    acfs.sample <- c(acfs.sample,acf.sample)</pre>
}
grid <- 10000
lambda <- seq(0,grid)*pi/grid</pre>
pgram <- cos(0*lambda)*acfs.sample[1]
for(h in 1:(n-1))
{
    pgram <- pgram + 2*cos(h*lambda)*acfs.sample[h+1]
pgram <- ts(pgram[-1],start=0,frequency=grid)</pre>
plot(log(pgram),xlab="Cycles",ylab="Log Periodogram")
```



print(pi\*which.max(pgram)/grid)

## [1] 0.04712389

## Lesson 9-8: Spectral Density Estimation

• We estimate the spectral density by modifying the periodogram.

#### Paradigm 9.8.1. Smoothing the Periodogram

ullet We can average the periodogram over nearby Fourier frequencies: for integer m,

$$\widehat{f}(\omega) = \frac{1}{2m+1} \sum_{j=-m}^{m} I(\omega + \lambda_j).$$

- Recall nonparametric smoothing: higher m to reduce variance, but bias increases.
- More general smoothing of the periodogram:

$$\widehat{f}(\omega) = \frac{\sum_{\ell} W_n((\lambda_{\ell} - \omega)/m) I(\lambda_{\ell})}{\sum_{\ell} W_n((\lambda_{\ell} - \omega)/m)},$$

where  $W_n(x)$  is a kernel function, or spectral window, depending on n. The estimator also depends on the bandwidth m.

#### Paradigm 9.8.3. Tapering the ACVF

• Recall from Remark 9.5.2 that tapering reduces variability in ACVF estimation. We can insert tapered sample ACVF into the periodogram formula:

$$\widetilde{f}(\lambda) = \sum_{|h| \le d} \Lambda(h/d) \widehat{\gamma}(h) e^{-i\lambda h},$$

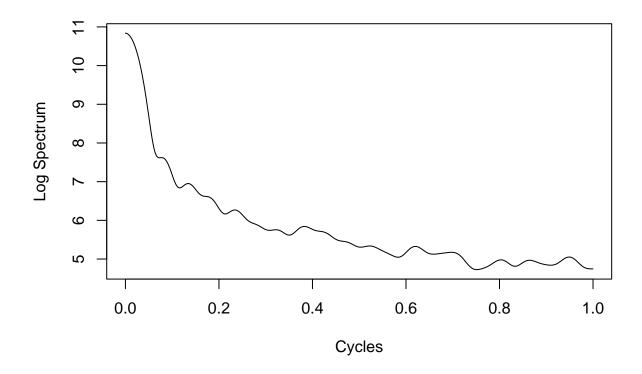
where  $\Lambda(x)$  is a taper (cf. Definition 9.5.3), and d is the bandwidth.

• We can rewrite  $\widetilde{f}$  as a spectral window estimator as well, where the spectral window is proportional to  $\sum_{|h| < d} \Lambda(h/d) e^{-i\lambda h}$ .

## Example 9.8.6. Bartlett Tapered Spectral Estimator

- We apply the tapered acvf spectral estimator of Paradigm 9.8.3, using the Bartlett taper (Example 9.5.4), to the Wolfer sunspot data.
- We choose the bandwidth  $d = 3n^{1/3} = 42$ , based on some asymptotic theory.
- We plot in logs, and so remove the frequency zero value.

```
wolfer <- read.table("wolfer.dat")</pre>
wolfer <- ts(wolfer[,1],start=1749,frequency=12)</pre>
data <- wolfer
n <- length(data)</pre>
d <- 3*floor(n^{1/3})
acfs.sample <- NULL</pre>
mu.hat <- sum(data)/n
for(k in 0:(n-1))
{
    acf.sample \leftarrow sum((data[1:(n-k)]-mu.hat)*(data[(k+1):n]-mu.hat))/n
    acfs.sample <- c(acfs.sample,acf.sample)</pre>
}
grid <- 10000
lambda <- seq(0,grid)*pi/grid</pre>
pgram <- cos(0*lambda)*acfs.sample[1]
for(h in 1:(n-1))
{
    pgram \leftarrow pgram + 2*(max(1-h/d,0))*cos(h*lambda)*acfs.sample[h+1]
pgram <- ts(pgram[-1], start=0, frequency=grid)</pre>
plot(log(pgram),xlab="Cycles",ylab="Log Spectrum")
```



## Lesson 9-9: Spectral Analysis

- We refine the analysis of spectral density estimation.
- We consider the class of flat-top tapers.

#### Paradigm 9.9.1. An Interesting Class of Tapers

- Consider a taper that takes the value 1 in a neighborhood of zero: this improves the bias.
- This suggests the definition  $\Lambda_{\infty}(x) = 1$  for  $|x| \leq c$ , for some  $c \in (0,1]$ . We call these flat-top tapers.
- Example: Bartlett taper (Example 9.5.4) is a limiting case with c = 0.
- Example: rectangular taper with c=1, where  $\Lambda(x)=1$  for  $|x|\leq 1$ , zero otherwise.
- Example: trapezoidal taper (depending on choice of c):

$$\Lambda_{\infty}(x) = \begin{cases} 1 & \text{if } |x| \le c \\ \frac{1-|x|}{1-c} & \text{if } c < |x| \le 1. \end{cases}$$

• Flat-top tapers correct bias by shifting the spectral estimate down, which can generate negative estimates. One can take the maximum with zero, so as to enforce a non-negative spectrum estimate.

#### Example 9.9.6. Application of a Trapezoidal Taper

- We apply a trapezoidal taper to the Wolfer sunspots, using c = 1/3 and d = 42.
- The spectral estimate has more variability, and is no longer positive (log of negative values is cut off).

```
wolfer <- read.table("wolfer.dat")
wolfer <- ts(wolfer[,1],start=1749,frequency=12)</pre>
```

```
data <- wolfer
n <- length(data)</pre>
d <- 3*floor(n^{1/3})</pre>
acfs.sample <- NULL</pre>
mu.hat <- sum(data)/n</pre>
for(k in 0:(n-1))
{
    acf.sample \leftarrow sum((data[1:(n-k)]-mu.hat)*(data[(k+1):n]-mu.hat))/n
    acfs.sample <- c(acfs.sample,acf.sample)</pre>
}
cutoff \leftarrow 1/3
trap <- function(h)</pre>
    val \leftarrow min(max((1 - h/d)/(1 - cutoff), 0), 1)
    return(val)
}
grid <- 10000
lambda <- seq(0,grid)*pi/grid</pre>
pgram <- cos(0*lambda)*acfs.sample[1]</pre>
for(h in 1:(n-1))
{
    pgram <- pgram + 2*trap(h)*cos(h*lambda)*acfs.sample[h+1]</pre>
}
pgram <- pmax(0,pgram)</pre>
pgram <- ts(pgram[-1],start=0,frequency=grid)</pre>
plot(log(pgram),xlab="Cycles",ylab="Log Spectrum",ylim=c(3,12))
```

