

Normalization of Power Spectral Density estimates.

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Abstract

A vast and deep pool of literature exists on the subject of spectral analysis; wading through it can obscure even the most fundamental concepts to the inexperienced practitioner. Appropriate interpretation of spectral analyses depends crucially on the normalization used, which is often this greatest source of confusion. Here we outline the normalization used by `rlpSpec`, namely the **single-sided power spectral density** (PSD). We briefly outline the background mathematics, present an example from scratch, and compare the results with the normalization used by the spectrum estimator included in the base distribution of R: `stats::spectrum`.

Contents

1	Background	2
1.1	Connection to the autocovariance function	3
1.2	Testing normalization	3
1.3	Summary of nomenclature	4
2	A from-scratch example: White noise.	4
3	Normalization used in <code>stats::spectrum</code>	9
4	Other PSD estimators	11
5	Session Info	11

1 Background

There can often be confusion about the different quantities used in spectral analysis, partly due to myriad nomenclature within the incredibly vast literature on the subject¹. Commonly one finds similarly sounding phrases, including “amplitude spectrum”, “energy spectral density”, “power”, “power spectra”, and even “spectra”. These all mean *something*, but are rarely equivalent, and are sometimes used improperly.

To clarify these terms, we will walk through some background and definitions, without overly complicating the discussion with proofs. Let us first define a single realization of a stochastic process in the time domain, $\mathcal{X}(t)$, sampled over a finite interval of time $(-T/2, T/2)$, denoted by $X_T(t)$. A realization of $\mathcal{X}(t)$ will have a Fourier transform:

$$\tilde{X}_T(f) = \mathcal{F}\{X_T(t)\} = \int_{-\infty}^{\infty} X_T(t)e^{-2\pi ift} dt = \int_{-T/2}^{T/2} \mathcal{X}(t)e^{-2\pi ift} dt \quad (1)$$

The **amplitude spectrum** is the modulus of \tilde{X}_T and the **phase spectrum** is the argument of \tilde{X}_T , although these are generally not informative for physical applications, if ever. The **energy spectral density** is found from \tilde{X}_T by finding the expectation of the squared amplitude spectrum:

$$E(f) = \mathcal{E}\{|\tilde{X}_T(f)|^2\} \quad (2)$$

There is a problem though, in that as T grows to infinity, so too does $E(f)$. We divide it by the interval length T to curb this growth, which gives us an expression for **power spectral density**:

$$\begin{aligned} S(f) &= \lim_{T \rightarrow \infty} T^{-1}E(f) \\ &= \lim_{T \rightarrow \infty} \mathcal{E} \left\{ \frac{1}{T} \left| \int_{-T/2}^{T/2} \mathcal{X}(t)e^{-2\pi ift} dt \right|^2 \right\} \end{aligned} \quad (3)$$

which is real, non-negative, and exists for all stationary processes with zero mean and finite variance.

Here is an important point to note regarding normalization: Equation (3) defines the **double-sided** PSD, because in the limit of T , the limits of integration are $\pm\infty$. If $\mathcal{X}(t)$ is real the power spectrum $S(f)$ is even; hence, we only need estimates for $f \geq 0$. The **single-sided** PSD is thus given by $2S(f)$ for $f \geq 0$. In many cases this sidedness distinction, as we will see, explains errant factors of two in PSD normalizations.

¹ A nice illustration of the type of confusion common in spectral analyses of confusion is found in this thread on R-help: <http://r.789695.n4.nabble.com/Re-How-do-I-normalize-a-PSD-td792902.html>

1.1 Connection to the autocovariance function

What is the connection between the PSD, defined in Equation (3), and the autocovariance function $\mathcal{R}(\tau)$?

From Equation (3) we see that $S(f)$ is obtained from products of $\mathcal{X}(t)$ with itself at any particular f , so it is related to the second-order moment of $\mathcal{X}(t)$ only; so too does the **autocovariance** (ACV) $\mathcal{R}(\tau)$:

$$\begin{aligned}\mathcal{R}(\tau) &= \text{Cov}(\mathcal{X}(\tau), \mathcal{X}(\tau)) \\ &= \mathcal{E}\{(\mathcal{X}(\tau) - \mathcal{E}\{\mathcal{X}(\tau)\})^2\}\end{aligned}\tag{4}$$

It may be surprising to note, as well, that $S(f)$ is simply the Fourier transform of $\mathcal{R}(\tau)$:

$$S(f) = \mathcal{F}\{\mathcal{R}(\tau)\} = \int_{-\infty}^{\infty} \mathcal{R}(\tau) e^{-2\pi i f \tau} d\tau\tag{5}$$

So, the functions $\mathcal{R}(\tau)$ and $S(f)$ exist as a transform pair. For real data, $\mathcal{R}(\tau)$ is always even, and always real. This implies that $S(f)$ is also a real and even function in f , which, because $S(f) \geq 0$, restricts the functions $\mathcal{R}(t)$ could possibly represent. Put another way, there are many examples of even functions having non-positive Fourier transforms (see Bracewell (2000)).

1.2 Testing normalization

We can use a property of the ACV function $\mathcal{R}(\tau)$ to test whether or not a PSD is properly normalized. To see this, we take the inverse Fourier transform of (5):

$$\mathcal{R}(t) = \int_{-\infty}^{\infty} S(f) e^{2\pi i f t} df\tag{6}$$

and find the ACV of a zero-mean process for zero lag. From (4) we have:

$$\mathcal{R}(0) = \mathcal{E}\{\mathcal{X}(t)^2\} = \mathcal{V}\{\mathcal{X}(t)\} = \sigma_{\mathcal{X}}^2\tag{7}$$

and by setting $t = 0$ in (6) we have the basis of our normalization test:

$$\sigma_{\mathcal{X}}^2 = \int_{-\infty}^{\infty} S(f) df\tag{8}$$

That is, the area under the power spectrum is the variance of the process. So, a straightforward way to test normalization is to compute the PSD for a realization of $\mathcal{X}(t)$ with known variance, and zero mean [e.g. $\mathcal{N}(0, \sigma^2)$]; and then calculate the integrated spectrum. For example, the single-sided PSD for a realization of a $\mathcal{N}(0, 1)$ process, sampled at 1 Hz, will be flat at 2 units²/Hz across the entire band $[0, 1/2]$, and will have an area equal to one.

1.3 Summary of nomenclature

In Table 1 we give a summary of some of the quantities we have reviewed.

Table 1: Top: Representers of various spectral quantities and their associated equations in the main text. Bottom: Normalizations for double and single sided PSD estimates to maintain process variance of $\sigma_{\mathcal{X}}^2$ and frequency ranges (f_N is the Nyquist frequency).

EXPRESSION	REPRESENTING	EQUIVALENT	EQUATION
$\mathcal{X}(t)$	stochastic process with variance $\sigma_{\mathcal{X}}^2$		
$X_T(t)$	single realization of $\mathcal{X}(t)$ over T		
$\mathcal{F}\{X_T(t)\}$	Fourier transform of $X_T(t)$	$\tilde{X}_T(f)$	1
$\sqrt{\Re^2 + \Im^2}$	amplitude spectrum of $X_T(t)$	$ \tilde{X}_T $ or $\text{mod}\{\tilde{X}_T\}$	
$\tan^{-1}(\Im/\Re)$	phase spectrum of $X_T(t)$	$\arg\{\tilde{X}_T\}$	
$ \tilde{X}_T ^2$	energy spectral density of $X_T(t)$	$E(f)$ or $\text{mod}\{\tilde{X} \star \tilde{X}^*\}$	2
$ \tilde{X}_T ^2/N$	power spectral density of $X_T(t)$	$S(f)$ or $\mathcal{E}\{ X ^2\}$	3
$\mathcal{F}^{-1}\{S(f)\}$	autocovariance of $X_T(t)$	$\text{Cov}(X_T(t), X_T(t))$	4 and 5
NORMALIZATIONS TO MAINTAIN $\sigma_{\mathcal{X}}^2$		FREQUENCY RANGE	
$S(f)$	double-sided	$[-f_N, f_N]$	
$2 S(f)$	single-sided	$[0, f_N]$	

2 A from-scratch example: White noise.

Without using the tools in `rlpSpec` we will build up an example using R commands, in order to highlight the topic of normalization.

First, generate a normally distributed series², and then find its Discrete Fourier Transform (DFT)³.

² Although a white noise process is not strictly bandlimited, we will use it to demonstrate differences in normalization.

³ A proper DFT is normalized by the length of the series; however, most DFT calculators (including `stats::fft`) eschew this normalization for efficiency's sake.

```

set.seed(1234)
N <- 1028
x <- rnorm(N, mean = 0, sd = 1)
xv <- var(x)
X <- fft(x)
class(X)

## [1] "complex"

length(X)

## [1] 1028

```

We can easily find the amplitude and phase response

```

Sa <- Mod(X)  # Amplitude spectrum
Sp <- Arg(X)  # Phase spectrum

```

followed by equivalent energy spectral density estimates⁴:

```

XC <- Conj(X)
all.equal(Se <- Sa^2, Se_2 <- Mod(XC * X), Se_2R <- Mod(X * XC))

## [1] TRUE

```

The single-sided power spectral density estimates follow once we have the Nyquist frequency, defined as half the sampling rate:

```

fsamp <- 1  # sampling freq, e.g. Hz
fNyq <- fsamp/2  # Nyquist freq
Nf <- N/2  # number of freqs
nyfreqs <- seq.int(from = 0, to = fNyq, length.out = Nf)
S <- Se[1:Nf] * 2/N  # Finally, the PSD!

```

To approximate the integrated spectrum in the case of a “flat” spectrum, we need an accurate measure of the first moment of the spectral values. The **median** estimator produces a biased estimate because the distribution of spectral values roughly follows a χ^2_ν distribution, where ν is the number of degrees of freedom and, for this distribution, the expectation of the

⁴ Note the equivalence between the complex conjugate based estimates.

first moment. To find this value, we will use a conjugate gradient minimization procedure to find the best-fitting χ^2 distribution. Our starting point will be the estimated mean value. We visualize the fit with a “Q-Q” plot, which shows PSD quantiles values as a function of χ^2 quantiles, using the optimized value of the number of degrees of freedom; this is shown in Figure 1.

```
# 0) Setup optimization function for dof, using conjugate gradients\\
# min L1 |PSD - Chi^2(dof)|
Chifit <- function(PSD) {
  optim(list(df = mean(PSD)), function(dof) {
    sum(log(PSD)) - sum(log(dchisq(PSD, dof)))
  }, method = "CG")
}
# 1) run optimization
Schi <- Chifit(S)
# Get 'df', the degrees of freedom
print(dof <- Schi$par[[1]])

## [1] 2.005

# compare with the mean and median
c(mSn <- mean(S), median(S))

## [1] 1.989 1.406
```

An estimate of the integrated spectrum should roughly equal the known variance. Figure 2 plots the PSD of our white noise series with the value of ν from the optimization, with which we can perform a variance-normalization test:

```
mSn <- dof
test_norm <- function(sval, nyq, xvar) {
  svar <- sval * nyq
  return(svar/xvar)
}
print(xv_1 <- test_norm(mSn, fNyq, xv))

## [1] 1.009

xv_2 <- sum(S)/Nf * fNyq/xv # an alternate test
all.equal(xv_1, xv_2)

## [1] "Mean relative difference: 0.007985"
```

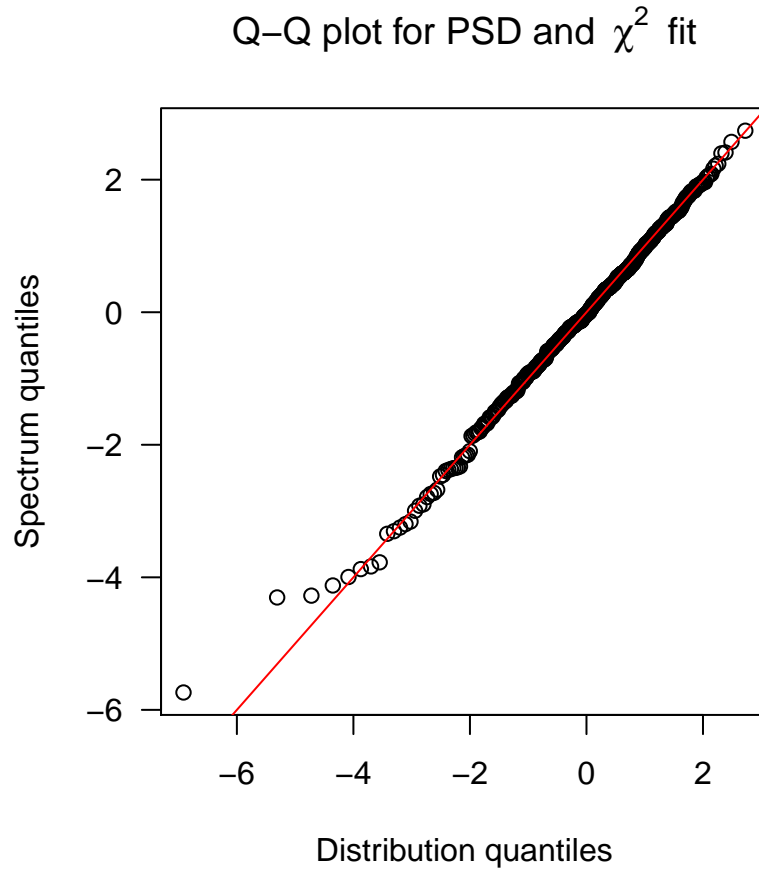


Figure 1: “Q-Q” plot of quantiles of our example PSD against theoretical χ^2_ν quantiles. The distribution is calculated with a value for degrees of freedom (ν) obtained from the L_1 -norm minimization procedure.

But what if the sampling frequency `fsamp` changes? An obvious change will be the actual Nyquist frequency, which means the variance-normalization test will fail if the PSD estimates are not re-scaled. We simply re-scale the frequencies and PSD with the sampling rate to obtain the properly-normalized spectra.

```
fsamp <- 20
fNyq <- fsamp/2
freqs <- fsamp * nyfreqs
```

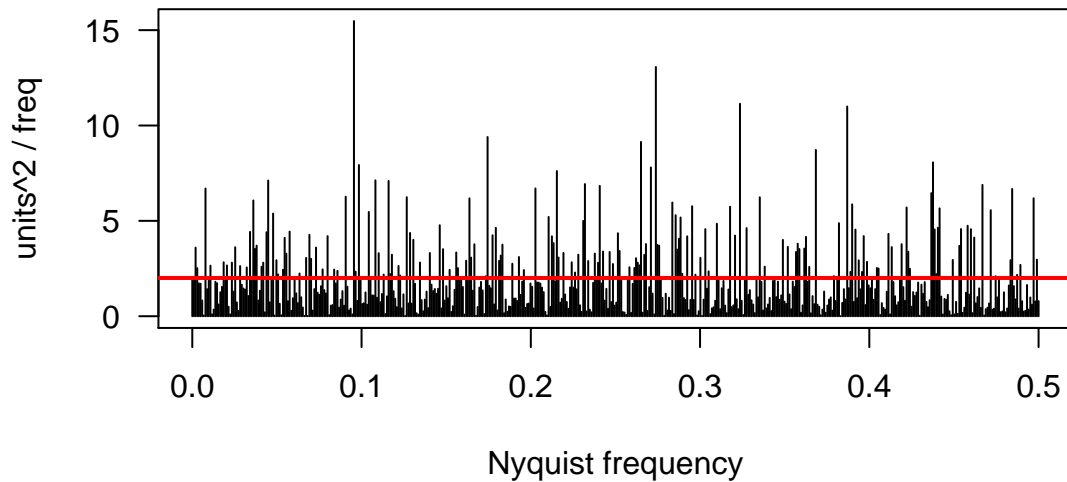


Figure 2: Power spectral density estimates for a single realization of a $\mathcal{N}(0, 1)$ process, in linear units. The horizontal line shows the expectation of the spectral from the χ^2 fit; this can be used to find the integrated spectrum and test normalization.

```
Snew <- S/fsamp
# Test variance crudely
mSn <- mean(Snew)
test_norm(mSn, fNyq, xv)

## [1] 1.001
```

In Figure 3 we plot the PSD with new normalization, and compare it to the previous normalization. Spectral values are shown as decibels (relative to 1 units²/frequency), using:

```
# decibel function
dB <- function(y) 10 * log10(y)
```

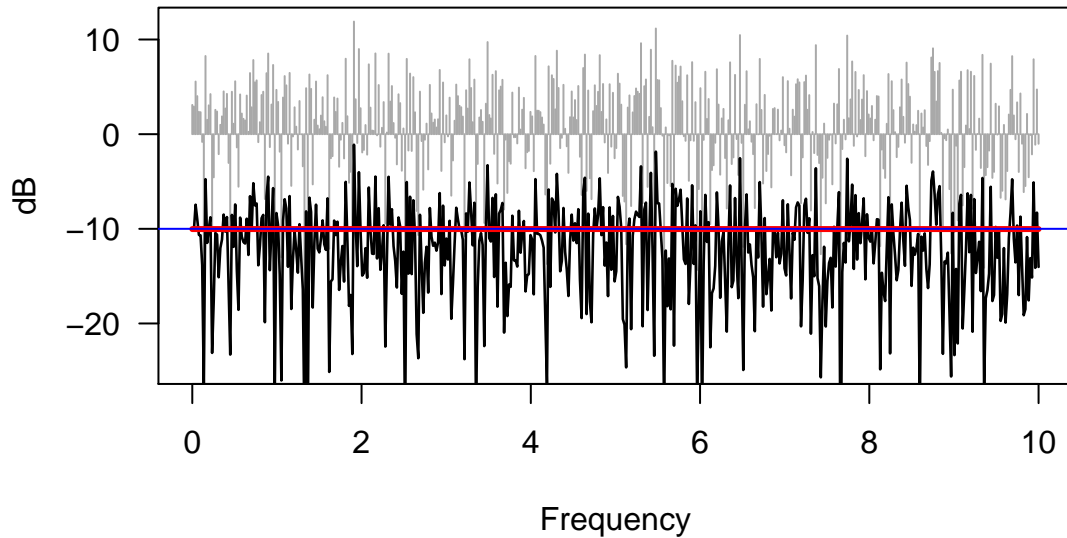



Figure 3: Rescaled PSD estimates for a single realization of a $\mathcal{N}(0, 1)$ process with a sampling rate of 20 s^{-1} rather than 1 s^{-1} as from before. The thick red line shows the mean (rescaled) spectral level, and the blue line shows the predicted mean value based on twice the sampling frequency.

3 Normalization used in `stats::spectrum`

The PSD estimator included in the core distribution of R is `stats::spectrum`, which calls either `stats::spec.ar` or `stats::spec.pgram` for cases of parametric and non-parametric estimation, respectively. For this discussion we compare to `spec.pgram`; the user can optionally apply a single cosine taper, and/or a smoothing kernel.

By default `spec.pgram` assumes the sampling frequency for the input series is 1, and normalizes accordingly; however, the sampling information may be specified by creating a `ts` object from the series prior to spectrum estimation:

```
fsamp <- 20
xt <- ts(x, frequency = fsamp)
pgram20 <- spec.pgram(xt, pad = 1, taper = 0, plot = FALSE)
```

```
pgram01 <- spec.pgram(ts(xt, frequency = 1), pad = 1, taper = 0, plot = FALSE)
```

We plot the two PSD estimates on the same scales, in Figure 4, utilizing the plot method for `spec` objects: `plot.spec`. We also show horizontal lines corresponding to the inverse of twice the sampling rate, which puts the spectra about a factor of 2 too low:

```
mSn/mean(pgram20$spec)
```

```
## [1] 2.004
```

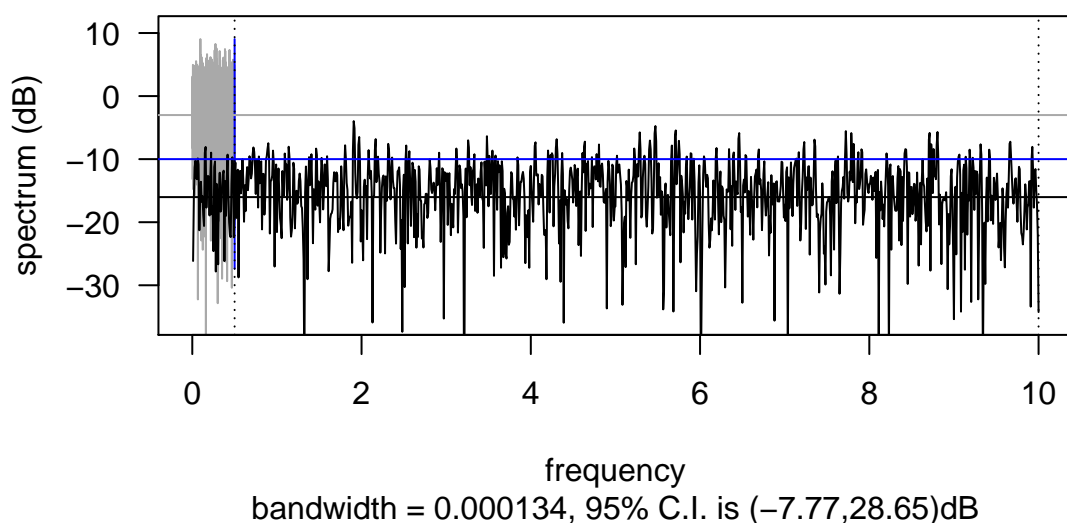


Figure 4: Power spectral densities from `spec.pgram` for the same data series. The grey curves show the PSD for a sampling rate of 1; whereas, the black curves show the PSD for a sampling rate of 20. The horizontal lines show levels corresponding to the inverse of twice the sampling rate (black and grey), and the expected spectral level for the 20 Hz sampling (blue). Vertical lines show the respective Nyquist frequencies.

Because the frequencies are clearly correct, this factor of two likely means the spectra will fail our simple variance-normalization test. They do fail, by a factor of two, again too low:

```
test_norm(mean(pgram01$spec), 0.5, xv)

## [1] 0.4994

test_norm(mean(pgram20$spec), 10, xv)

## [1] 0.4994
```

But why? This errant factor of two comes from the assumption of a double-sided spectrum, which is at odds with our definition of the single-sided spectrum by—you guessed it—a factor of two. We can illustrate this with the following example, where we compare the PSDs from `spec.pgram` for a real and complex series:

```
psd1 <- spec.pgram(x, plot = FALSE)
psd2 <- spec.pgram(xc <- complex(real = x, imag = x), plot = FALSE, demean = TRUE)
mx <- mean(Mod(x))
mxc <- mean(Mod(xc))
(mxc/mx)^2

## [1] 2

mean(psd2$spec/psd1$spec)

## [1] 2
```

Again, a factor of two. This means that unless we are interested in analyzing complex timeseries, we need only multiply by two to obtain properly normalized spectra from `spectrum`, assuming the sampling information is included in the series.

4 Other PSD estimators

The suite of extensions having similar functionality to `rlpSpec` is relatively limited; however, there are at least three which can produce sophisticated PSD estimates. We have summarized the available functions in Table 2 so far as we know⁵.

5 Session Info

⁵ As of this writing (Feb 2013), `sapa` appears to be orphaned.

Table 2: A comparison of power spectral density estimators in R, excluding extensions which only estimate raw-periodograms. Normalizations are shown as either “single” or “double” for either single- or double-sided spectra, and “various” if there are multiple, optional normalizations. A (*) denotes the default for a function having an option for either single or double.

FUNCTION	NAMESPACE	SINE M.T.?	ADAPTIVE?	NORM.	REFERENCE
mtapspec	RSEIS	YES	NO	various	Lees and Park (1995)
pspectrum	rlpSpec	YES	YES	single	Parker and Barbour (2013)
spectrum	stats	NO	NO	double	R Core Team (2012)
spec.mtm	multitaper	YES	YES	double	Rahim and Burr (2012)
SDF	sapa	YES	NO	single*	Percival and Walden (1993)

```
sessionInfo()

## R version 2.15.2 (2012-10-26)
## Platform: x86_64-apple-darwin9.8.0/x86_64 (64-bit)
##
## locale:
## [1] C
##
## attached base packages:
## [1] parallel datasets grDevices grid graphics tools stats
## [8] utils methods base
##
## other attached packages:
## [1] knitr_0.9
##
## loaded via a namespace (and not attached):
## [1] digest_0.6.0 evaluate_0.4.3 formatR_0.7 stringr_0.6.2
```

References

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- Lees, J. M. and Park, J. (1995). Multiple-taper spectral analysis: A stand-alone C-subroutine. *Computers & Geosciences*, 21(2):199–236.
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Index

amplitude spectrum, 2, 4

autocovariance, 2, 4

double-sided, 2, 4, 10

energy spectral density, 2, 4, 5

phase spectrum, 2, 4

power spectral density, 1, 2, 4, 5

single-sided, 1–4, 10