# Principles of Neurobiological Signal Analysis

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## **PREFACE**

More years ago than we care to think of or mention, we convinced ourselves of the need for a monograph on the principles of signal analysis as applied to the electrical activity of the nervous system. This book is the result. Our premise in organizing it has been simple: that neurobiologists are generally uneasy in their use of signal analysis simply because they have had little formal training in the mathematics underlying its framework and that therefore they have little intuitive feel for what signal analysis procedures mean. Our goal, consequently, is to provide neurobiologists with a reasonably detailed discussion of signal analysis as it has been variously applied to neuronal signals. We wish to make them more aware of what these analyses can and cannot do, their implications, and limitations. We have used mathematics where it is essential, but in doing so we have tried to avoid unnecessary rigor. We have assumed that mathematically the reader is equipped with a hazy recollection of calculus. Our hope is that we can dispel most of this haze in the early going. On another front, we have consciously refrained from treating the cuisine of signal analysis. Recipes or programs for signal analysis are readily available for a variety of computers. We do not feel they provide much elucidation of the basic issues.

The first three chapters establish the theoretical groundwork of signal analysis. Chapter 1 presents an introductory discussion of the properties of signal and noise, especially as they apply to the nervous system. It reflects our judgment that the essential ingredients of neurobiological signal analysis are the related concepts of signal spectra and covariance functions. They are likely to remain so even as the present, predominantly linear methods of signal analysis are broadened to encompass nonlinear techniques. Chapter 2 discusses the methods of sampling and converting biological signals into sequences of digital numbers readily digestible by a computer. Chapter 3 then develops more thoroughly the concepts of spectrum and covariance analysis. This chapter is mathematically somewhat more demanding than the first two. Those who find it too trying should not feel distressed since much of what appears subsequently will still be comprehensible. The loss is in the appreciation of some of the analytic details.

Chapters 4 and 5 deal with techniques for extracting evoked responses from background noise and with multivariate statistical procedures for treating evoked response waveshapes as variables dependent upon the experimental manipulations performed upon a subject. Chapters 6 and 7 deal with the analysis of spike (action potential) activity generated by individual neurons and small groups of neurons. Chapter 8 presents methods for studying how such spike activity may be related to

the concurrently observed slow wave (EEG-like) activity of the nervous system.

A number of individuals have contributed to the completion of this work. It was Dr. José del Castillo who provided us with facilities at the Laboratory of Neurobiology of the University of Puerto Rico. It was there that this book had its inception. Drs. Donald Childers, Emanuel Donchin, and George Gerstein reviewed various chapters and provided much helpful criticism. A special note of thanks goes to Drs. José Negrete and Guillermina Yankelevich de Negrete who lent much encouragement during the initial tribulations of writing. Finally, we would like to express our special appreciation to Mrs. Frances Pridgen who, equipped with an extensive background as a legal secretary, typed the manuscript and suffered with us in guiding it to completion. In a moment of relaxation, when all was done, we asked her opinion of the work. She flipped slowly through its pages, smiled and said, "Naturally, this is taxable." We wonder.

eface	•	X
	CHAPTER 1 SOME PROPERTIES OF BIOLOGICAL SIGNA	LS
1.1	Introduction	1
1.2	Continuous Signals and Their Discrete Counterparts	2
1.3	Repetitive and Periodic Signals	8
1.4	Sampled Representation of a Signal	9
1.5	Fourier Series Representation of a Signal	11
1.6	Bandwidth Limited Signals	15
1.7	Autocovariance Functions and Power Spectra	
	of Periodic Signals	17
1.8	Aperiodic Signals	20
1.9	Autocovariance Functions and Power Spectra	
	of Aperiodic Signals	21
1.10	Cross Covariance Functions and Cross Spectra for	
	a Pair of Periodic Signals	23
1.11	A Summary of Properties of Covariance Functions	
	and Spectra	25
1.12	Random or Probabilistic Signals	26
1.13	Some Important Probability Distributions	
	A. Probabilistic Descriptions of Dynamic Processes	32
	B. The Gaussian Distribution	33
	C. The Chi-Squared Distribution	34
1.14	Ensemble Autocovariance Functions	37
	Ensemble Autocovariance and Cross	
	Covariance Functions, and Stationarity	39
1.16	The Relationship between Ensemble	
.,,,	and Time Statistics	43
1.17	Mixtures of Signal and Noise	45
	Response Detection and Classification—	
	Hypothesis Testing	47
	CHAPTER 2 BASICS OF SIGNAL PROCESSI	٧G
2.1	Introduction	55
2.2	Analog-to Digital Conversion	55
2.3	Quantization Noise	58
2.4	Multiplexing: Monitoring Data	
	Sources Simultaneously	62

2.5	Data Filtering												64
2.6	The Digital Filter												65
	A. Filtering of the Constant Component .												67
	B. Filtering the mth Frequency Component												68
2.7	Impulse Response of a Digital Filter												70
2.8	Spectral Relations between Filter Input												
	and Output-the Discrete Fourier Transform												72
2.9	Filtering Aperiodic Signals									-			
	A. Short Duration Signals							_					74
	B. Maintained Signals		-				Ī	Ī	Ī	•	•	•	76
2.10	Data Smoothing by Digital Filtering					•	•	•	•	•	•	•	79
2.11	Digital Filters with Feedback—	-	-	•	•	•	•	•	•	•	•	•	
	Recursive Filters	_	_			•							84
2.12	The Linear Analog Filter			•		•	•	•	•	•	•	•	86
	The Laplace Transform, the Filter	•	•	•		•	•	•	•	•	•	•	•
	Transfer Function, and Impulse Response												87
2.14	The Operational Amplifier	•	•	•	• •	•	•	•	•	•	٠	•	94
2 15	The Amplitude Comparator	•	•	•		•	•	•	•	•	•	•	97
2 16	Time-Varying and Nonlinear Filters	•	•	•		•	•	•	•	•	•	•	100
	Time varying and recommend raters	•	•	•		•	•	•	•	•	•	•	100
	CHAPTER 3 POWER SPECTRA	A	NE	C	OV	ΑF	RI/	N	CE	F	UN	СТ	IONS
3.1	Introduction												103
3.2	Discrete Fourier Representations						-	-	-			-	
	of Continuous Processes								_	_			104
3.3	Aliasing		_			Ī	Ī		•	Ċ	·	•	110
3.4	Leakage	•	-	•	•	•	٠	٠	•	•	•	•	
	A. Fourier Series												116
	B. Discrete Fourier Transforms	•	•	•	• •	•	•	•	•	٠.	•	•	120
3.5	Trend	•	•	•		•	•	•	•	•	•	•	123
3.6	The Power Spectrum, General Considerations	•	•	•		٠	•	•	•	•	•	•	124
3.7	Power Spectrum of Continuous Random Signal		•	•	• •	•	•	•	•	•	•	•	130
3.8	The Power Spectrum of	•	•	•	• •	•	•	•	•	•	•	•	130
	T-Discrete Signals												135
3.9	The Fourier Transform for	•	•	•	•	•	•	•	•	•	•	•	135
	7-Discrete Signals												100
3 10	The Periodogram	•	•	•	•	•	•	٠	•	•	•	•	136
3 11	Statistical Errors of the	•	•	•	•	•	•	•	٠	•	•	•	138
· · · ·													
3 12	Periodogram—Bias	•	•		•	•	•	٠	•	٠	•	•	141
5.12													
2 12	Periodogram—Variance		•		٠	٠	•	•	•	٠		•	144
5.15	the Pertine Feriodogram—												
2 14	the Bartlett Estimator				•		•						147
3.14 3.1E	Variance of the Bartlett Estimator				•		٠						149
J. 15	the Fast Fourier Transform												
2 4 6	and Power Spectrum Estimation					•							150
3.16	Smoothing of Spectral Estimates												
~ <i>-</i> -	by Windowing												151
3.17	The Cross Spectrum												155
3.18	Covariance Functions												157

	A. Some Statistical Properties													158
	of the ACVF Estimators	٠	•		•	•	٠	•		•			•	162
	B. Estimation of the ACVF		•		•	•	•			•	•		•	
	C. Cross Covariance Function Estimation		•			•	•	•	•	•		•	•	166
3.19	Coherence Functions				•	•		•	•	•			•	168
3.20	Phase Estimation							•		•			•	174
														A A I D
	CHAPTER 4 EVOKED	PO	TE	N1										AND
					D	ISC	RI	MII	NA	/M	I A	N.	AL	YSIS
	Introduction									_		_		177
4.1	Estimation of Variability													180
4.2	Confidence Intervals													182
4.3	Comments on Assumptions													184
4.4	Alternative Measures of Variability								-					185
4.5	A. Split Sweep Assessment of Significance	•	•	•										185
	B. Plus-Minus Reference Method													188
		•	•	•	•	• •	•	•	•	•	•	•		
4.6	Correlated Noise, Overlap and Stimulus Spacing													189
									•	•	•	•	•	192
	A. Aperiodic Stimuli	• ••••	•	•	•	• •	•			•	•	•	•	194
	B. Narrow Band Noise and Aperiodic Stin								•	•	•	•	•	196
4.7	The Median Evoked Response Nonhomogeneous Sets of Evoked Potential	•	•	•	•	• •	•	•	•	•	•	•	•	201
4.8		5	•	•	•		•	•	•	•	•	•	•	
4.9	Correlation Estimation of a Constant													201
	Waveform with Varying Latency										:	•	•	205
	Homogeneous Subsets												•	206
4.11	The Cumsum Procedure												•	207
	A. The Precum Method	•	•	•	•	• •	•	•	•	•	•	•	•	207
	B. Distribution of Precums													210
	for Homogeneous Responses							•	٠	•	•	•	•	210
	C. An Algorithm for Computing Precum							•	•	٠	٠	•	•	
	The Sort Method						٠	٠	•	•	٠	•	٠	212
	Discriminant Analysis						•	٠	•	٠	•	•	•	220
	Stepwise Discriminant Analysis									•	•	•	•	225
4.15	Appendix: Histograms, Smoothing, N Mo	de		•	•		•	•	•	•	•	•	•	229
										_	~-	<b>.</b>	~~!!	CNITO
	CHAPTER 5 EVOKED POTEN	TIF	\L	<b>S</b> :										ENTS
					•	ANI	ν	Ar	( IIV	1A	X A	٨N	IAL	YSIS.
5.1	Introduction						٠.							233
5.2	Linear Representation of Waveforms													235
5.3	The Cross Correlation Coefficient													236
5.4	Signal Space													236
5.5	Linear Expansion Methods, Factor Analysi													
0.0	and Other Techniques													241
5.6	Factor Analysis and Principal Factors													244
5.7	Matrices and Signal Analysis													245
···	A. Matrix Properties, Definitions					. •	•	-	•	•				245
	B. Matrix Addition			-	•				•					246
	C. Scalar Multiplication							•	•	•		-		247
	D. Matrix Multiplication							•	•	•	•	•	•	247
5.8	Matrices and Linear Expansions of Wavefo							•	•	•	•		•	247
J.J	A. Cross Correlation							:	•	•	•	•	•	249
	A. 0109 COHEIGHOH	•	•	•	•		•	•	•	•	•	•	•	270

5.9	The state of the s	249
5.	10 Some Special Matrices	250
5.	11 Matrix Inverse	251
5.1	12 Orthogonal Matrices	252
5.1	13 Properties of Linear Expansions Based upon	202
	Orthonormal Basic Waveforms	252
5.1	4 Principal Components	255
5.1	5 Computation of Principal Components	255
5.1	6 Covariances and Correlation Coefficients	263
5.1	7 Dimensionality and Eigenvalues	264
5.1	8 Varimax Rotation of the Weighting Coefficients	
5.1	9 Varimax Rotation of the Basic Waveforms	265
5.2	O Principal Component Analysis and the	268
	Karhunen-Loeve Expansion	000
5.2	1 Principal Components—Varimax Analysis of	268
	Deviation Waveforms	
5.2	2 Covariances, Correlation Coefficients, and	270
	Implied Baselines	
5.2	3 Principal Component—Varimax Analysis Based	273
	upon Orthonormal Weighting Coefficients	
5.24	4 Examples	274
	A. Auditory Evoked Responses and Masking Effects	
	B. Effects of Drugs upon Evoked Responses	277
	C. Independence of Components Recorded from	280
	Scalp of Humans	
5.25	5 Some General Remarks on Linear Expansions	282
	Expansions	285
	CHAPTER 6 SPONTANEOUS AND DRIVEN SINGLE UNIT ACTIV	<del></del> .
	AND THE STATE OF T	/IIY
6.1	Introduction	291
6.2	For Frocess—an Idealization of Neuronal Spike Activity	292
	A. Spontaneously Active Processes	293
~ ~	B. Driven Processes	294
6.3	Classification of Spontaneously Active Processes	296
	A. Renewal Processes	301
	B. Poisson Processes	302
6.4	Spike Data Acquisition	303
6.5	Interval Distribution, Mean and Variance	306
6.6	rests for Mean Interval and Rate of a Poisson Process	309
6.7	lests for a Poisson Process	313
6.8	lests for the Parameters of a Gamma Renewal Process	316
6.9	Serial Statistics and Nonrenewal Processes	317
6.10	interval Southing as a Test for Renewal Processes	324
6.11	The Expectation Density and Covariance Function	J24
	of Point Processes	325
6.12	abase at August 201 abive Seddelices	320
	A. Relationship to the Expectation Density	335
	D. Shoothed Estimates of Point Process Spectra	
6.13	Solicial Considerations in Spectral Smoothing	340
6.14	The opecition of intervals and its Relationship to	343
	the Serial Correlogram	247
R 1E		347
0.15		351

6.16	Peristimulus Time Histogram Analysis	
	of Driven Activity	357
6.17	Tests for Response Dependency on the Stimulus	358
6.18	Response Trends	362
6.19	Data Displays	367
	CHAPTER 7 MULTIPLE UNIT ACTIV	ITY
7.1	Introduction	373
7.2	Cross Covariance Methods	376
	A. Cross-Expectation Density Analysis	378
	B. The Cross-Expectation Density during Spontaneous	
	Activity or Continuous Stimulation	379
	C. The Cross-Expectation Density during Stimulation	382
7.3	Interspike Interval Tests for Unit Dependency	385
7.4	Data Displays for Two Stimulated Units—	-
	the Peristimulus Time Scatter Diagram	394
7.5	Data Displays for Three Units—the Snowflake Diagram	400
7.6	Separation of Activity of Concurrently Discharging Units	405
	A. Use of Salient Waveform Parameters:	703
	Amplitude, Width, Slope	407
	P. Has of the Cutine Cuite Message	
	B. Use of the Entire Spike Waveform	412
	C. Identification Errors and the Analysis	404
	of Unit Interactions	421
7.7	Multiunit Activity as an Entity	423
СНА	PTER 8 RELATIONS BETWEEN SLOW WAVE AND UNIT ACTIV	ITY
8.1	Introduction	427
8.2	Some General Considerations on Covariance	
	and Spectral Analysis	428
8.3		432
8.4	Estimation of Cross Covariance between a	
	Continuous and a Point Process	435
8.5	Some Dangers in Cross Covariance Estimation	
		441
8.6	Cross-Expectation Relations between a Point Process and	• • • •
		443
8.7		447
8.8		450
8.9		454
2.10	Continuous Process Probabilities Conditioned	454
J. 1 U		455
R 11	by Point Process Events	455
J. 1 I		4
Ω 12		458
U. I Z	Changes of State in Point and Continuous Processes	461
ject	Index	465

### Chapter 1

# SOME PROPERTIES OF BIOLOGICAL SIGNALS

### 1.1. INTRODUCTION

Speaking in a somewhat general way, we say that all biological data can be considered to be signals. Obviously, however, some data are more signallike than others. The dividing line between data that can be profitably considered to be signallike and data that cannot depends upon both the origin of the data and how we propose to process it and analyze it conceptually. A discussion of the many facets of this idea in the light of modern computer data processing methods is one of the major purposes of this book. Embarking in this direction requires that we first establish some of the major concepts and properties of signals insofar as they relate to biological processes. The properties of these signals influence, guide, and sometimes determine the ways in which computer programs are developed to perform signal analysis.

Signal: A variation in the amplitude and polarity of an observed physical quantity produced by a process whose mechanisms we desire to understand by experimental investigation.

The requirement that the variation be produced by a mechanism we are interested in is of basic importance and brings us to consider at once, noise, the inseparable companion of signal.

Noise: A variation in the size of an observed physical quantity we are investigating produced by a process or an aspect of a process that we have no present interest in.

Data: Some combination, often additive, of signal and noise. The additive situations are easiest to deal with in terms of analysis and interpretation of results. In much of what follows we will assume it applies. In general, however, additivity should not be taken for granted.

### PRINCIPLES OF NEUROBIOLOGICAL SIGNAL ANALYSIS

The errant course of scientific progress is such that often what is considered to be a signal in one investigation turns out to be noise in another. Or more colloquially, one man's signal is another man's noise.

The variations in the size of a physical quantity are often time-dependent. When they are, the data is said to be a function of time and written x(t). Temporal data variation is most convenient for us to consider and also most appropriate since a realtime computer generally accepts data in time sequential form. However, we may also profitably consider data which are functions of such variables as distances or angle, for it is usually a simple matter to convert them into functions of time by a signal transducer. As an example, a scanning densitometer converts the spatially varying density of a translucent object into a function of time as the densitometer is moved over the scanned object. oscilloscope screen is an example of the process in reverse for there the time-varying data is converted into a function of distance along the horizontal axis of the oscilloscope screen. after, when we mention data signals and noise, we will consider them to be temporally varying.

We are interested in establishing the basic principles of a wide assortment of procedures by which we analyze the signallike data of neurobiological investigations. Temporally generated signals and noises exhibit a wide variety of waveform features or parameters, and it is essential to classify them according to such features, for the validity of much of the subsequent data processing depends upon the presence or magnitude of these features. The following pages contain a discussion of some of the properties of signals to serve as the basis of understanding the signal analysis procedures and techniques to be described in later chapters.

# 1.2. CONTINUOUS SIGNALS AND THEIR DISCRETE COUNTERPARTS

Let us begin with data which consist only of signals. A

### SOME PROPERTIES OF BIOLOGICAL SIGNALS

signal is said to be continuous if it is defined at all instants of time during which it occurs. A continuous signal may, however, possess discontinuities or sudden changes in amplitude at certain instants of time. At these instants the slope of the signal is infinite. At other times the signal amplitude changes gradually so that by choosing an interval short enough, the corresponding change in amplitude can be made as small as we like. While continuous signals without discontinuities are the rule in such biological phenomena as the EEG, deliberately generated discontinuous signals may be generated by the instrumentation associated with neurobiological investigations. As an example, the signal produced by a rat when it pushes a switch to obtain food is discontinuous. This type of signal is referred to as a step function. Illustrations of continuous and discontinuous signals are shown in Fig. 1.1. It is also to be noted that whether continuous or not, the signals are always single valued: they have only one value at any particular instant in time. A particularly interesting and important discontinuous signal is the unit step signal of Fig. 1.1(c):

$$u(t) = \begin{cases} 0 & \text{when } t \leq t_d, \\ 1 & \text{when } t > t_d \end{cases}$$
 (1.1)

 $t_d$  is the instant of discontinuity. The equation indicates that the signal jumps to 1 as soon as t becomes greater than  $t_d$ . The unit step is used, among other purposes, to describe a stimulus that has a sudden onset.

Besides speaking of a continuous signal, x(t), we will also have occasion to speak of its time derivatives, the first derivative being written dx(t)/dt or, alternatively, x'(t). The first derivative is, of course, the time rate of change of the variable. When it is zero, the variable itself is at a local maximum or minimum value or, less frequently, at an inflection point. (The derivative of a constant signal is always zero.) This property is often used in determining when a spikelike waveform reaches a maximum or minimum. A peak detection device which essentially

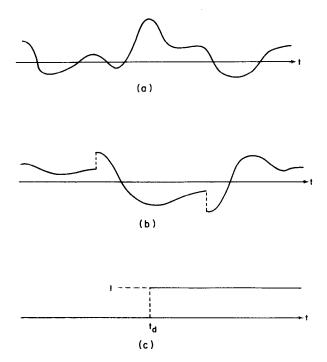


Fig. 1.1. (a) A continuous signal; (b) a discontinuous signal; (c) the unit step  $\mu(t)$ , showing step onset at  $t=t_d$ .

takes the time derivative of the waveform is commonly employed for this. When its output, the waveform time derivative, goes through zero in a negative direction, a positive maximum has occurred; when it goes through zero in a positive direction, a negative maximum has occurred. Figure 1.2(a) illustrates the situation for the former case. The first derivative is also important in indicating when the signal is changing most rapidly because it has its greatest value at that time. A positive maximum in the first derivative indicates the time when the signal is increasing most rapidly; a negative maximum, when it is decreasing most rapidly. Just as a continuous signal may exhibit discontinuities, so may its derivatives. A discontinuity in the first derivative occurs when there is a cusp in the original signal. An example is the sawtooth signal of Fig. 1.2(b). When it is at its maximum and

### SOME PROPERTIES OF BIOLOGICAL SIGNALS

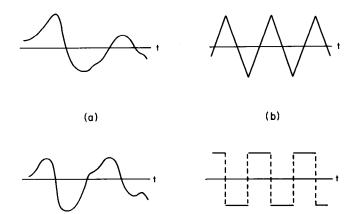


Fig. 1.2. (a) Above, a continuous signal; below, its time derivative. The negative and positive going zero crossings of the derivative correspond to positive and negative peaks in the signal. (b) Above, a periodic sawtooth signal; below, its time derivative which is a periodic discontinuous square wave.

minimum values, discontinuities occur in its first derivative, a square wave.

The derivative operation is not without practical difficulties since noise contributions tend to corrupt the derivative measurement. In computer analysis of data, the derivative operation is approximated by comparing successive sampled values of the signal with one another to see when maximum and minimum rates of change occur. Although this is an approximation, the results are often more than adequate. It is worth noting here that approximation is different from estimation, the latter being a statistical procedure whose meaning will be made clear in the subsequent pages.

In contrast to temporally continuous (T-continuous) signals are the temporally discrete (T-discrete) signals. These are signals which exist only at discrete instants in time. For our purposes the most important discrete signals are those which occur when a continuous signal has its amplitude measured or sampled at discrete instants of time that are usually equally spaced. A T-discrete signal is thus a sequence of measurements  $x_1, x_2, \ldots$ 

### PRINCIPLES OF NEUROBIOLOGICAL SIGNAL ANALYSIS

 $\mathbf{x}_T$  lasting for the duration of the time the signal is observed. In digital data processing it is furthermore usually quantized in amplitude by an analog-to-digital converter. This gives it the property of being amplitude discrete (A-discrete). The result is a signal, T- and A-discrete, which provides the basic data thereafter for all subsequent computer analyses of the original signal.

Having introduced the continuous signal and its sampled T-discrete representation, it is useful to establish here a form of notation which permits us to distinguish between them with a minimum amount of confusion. We will use the symbol ° to distinguish a sampled T-discrete signal from its continuous source signal. We will drop the ° when no confusion seems possible. Similarly, we will use t to represent continuous time and t° $\Delta$  to represent those instants that a signal is sampled at a uniform rate.  $\Delta$  is the interval between neighboring samples, and t° is an integer-valued index: 1, 2, 3, ..., etc. Signal analyses are often most easy to describe when  $\Delta$  = 1. This results in no loss of generality. When there is no possibility of confusion, the  $\Delta$  will be dropped.

The signals or data handled by a digital computer are discrete not only in time but also in amplitude. This arises from the fact that the amplitude of a signal at a particular sampling instant is represented as a number within the computer, a number containing a limited number of digits or bits depending upon the computer's structure. To arrive at this numeric representation a continuous signal is first transformed into its A-discrete amplitude version by quantization in an analog-to-digital (A-D) converter. At each sampling time the quantization procedure assigns to the signal amplitude one of a finite number of levels. This level has a numeric value which represents the sample in subsequent data analysis computations. The subject of A-D conversion, or quantization, is discussed more thoroughly in Chapter 2.

Perhaps the simplest way of reconstructing a continuous signal from a set of its samples is shown in Fig. 1.3. Here the signal is assumed to remain constant at its sampled value for the

### SOME PROPERTIES OF BIOLOGICAL SIGNALS

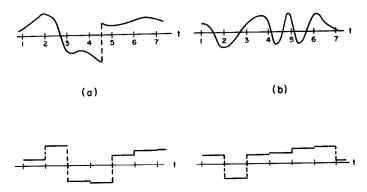


Fig. 1.3. (a) Above, a signal with a discontinuity between t = 4 and 5; below, a reconstruction of that signal by interpolation with a constant value between sampling instants. (b) Above, another continuous signal fluctuating rapidly between the 4th and 5th sampling instants. Note how the same type of sampling reconstruction totally lacks evidence of the rapid fluctuation of the original.

time interval between the present and the next sample time. It is important to recognize that the sampling and interpolation process can produce severe alterations of the signal depending upon the interrelationships between signal and sampling parameters. Two of the simplest errors are seen in Fig. 1.3 where in (a) a discontinuity is lost and in (b) a rapidly fluctuating component is suppressed because the sampling rate is too low. This type of error occurs regardless of how the interpolation between sampling instants is performed. A more thorough discussion of sampling problems is also presented in Chapter 3.

In some cases a signal is intrinsically T-discrete as for example is the count of the number of events occurring within an interval of time, such as the number of times an EEG waveform has a zero-crossing (a transition through zero amplitude) in one second. A second example is a list of measurements characterizing the structure of an object. It is important to note, however, that in the latter example the order in which the measurements are placed into a sequence may be of little or no importance. In temporal measurements or in measurements that are functions of a scanning