

A Time-Domain Analysis of Intracardiac Electrograms for Arrhythmia Detection

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DICARLO, L.A., ET AL.: A Time-Domain Analysis of Intracardiac Electrograms for Arrhythmia Detection. The analysis of intracardiac electrogram morphology has been proposed as a complementary method for accurate discrimination between sinus rhythm (SR), supraventricular dysrhythmias, and ventricular dysrhythmias by automatic antitachycardia and cardioverter defibrillator devices. In this study, the performance of a traditional time-domain method for surface electrocardiogram interpretation—Correlation Waveform Analysis (CWA) and a newly developed technique—Bin Area Method (BAM) were used to analyze unfiltered intraatrial and intraventricular electrograms obtained from 47 patients during routine cardiac electrophysiology studies. Nineteen patients had 31 distinct, sustained, monomorphic ventricular tachycardias (VTs) induced; 13 patients had paroxysmal bundle branch block of supraventricular origin (BBB) induced; 19 patients had retrograde atrial activation during ventricular overdrive pacing. Three patients were common to two or more groups. Using a best fit electrogram alignment, both CWA and BAM distinguished VT from SR in 28/31 cases (90%), BBB from SR in 15/15 patients (100%), and anterograde from retrograde atrial activation in 19/19 patients (100%). We conclude that the use of time-domain techniques that are independent of amplitude and baseline fluctuations appear to be reliable for discrimination of retrograde atrial activation, paroxysmal BBB, and VT from SR using intracardiac electrograms. Reduction of computational time and power constraints, without sacrificing reliable dysrhythmia discrimination, is possible. These features may make real-time morphology analysis of intracardiac electrograms feasible for automatic antitachycardia and cardioverter-defibrillator devices. (*PACE*, Vol. 14, February, Part II 1991)

arrhythmia analysis, computer recognition of arrhythmia

Introduction

The use of rate and rate variation alone by currently available automatic antitachycardia and cardioverter-defibrillator devices to discriminate between sinus rhythm (SR), supraventricular dysrhythmias, and ventricular dysrhythmias has not been satisfactory.¹⁻⁴ The utilization of comple-

mentary algorithms which analyze the morphology of intracardiac electrograms has been proposed as one means of achieving more accurate discrimination.⁶⁻¹⁰

In this study, we assessed the performance of a time-domain method traditionally utilized for surface electrocardiogram interpretation—correlation waveform analysis (CWA) and a new technique—Bin Area Method (BAM) for discrimination of retrograde atrial activation, paroxysmal bundle branch block (BBB), and ventricular tachycardia (VT) from SR with and without chronic BBB, an intraventricular conduction delay, and/or antiarrhythmic therapy.

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Methods and Materials

Electrophysiology Study

Bipolar (1 cm) atrial and ventricular endocardial electrograms were recorded during elective clinical cardiac electrophysiology studies as previously reported.⁹ Nineteen consecutive patients had 31 distinct, sustained, monomorphic VTs induced (group 1)(Table I); 13 patients had paroxysmal BBB of supraventricular origin induced (group 2)(Table II); 19 consecutive patients had 1:1 retrograde atrial activation during ventricular overdrive pacing (group 3)(Table III). None of the patients had dual atrioventricular nodal pathways or accessory atrioventricular connections. One patient was common to all three groups, and two patients were common to two groups.

Methods of Analysis

Recorded endocardial electrograms were subsequently replayed and digitized on a personal computer with a Tecmar Lab Master (Scientific Solutions, Inc., Solon, OH, USA) analog-to-digital system at a sampling rate of 1000 Hz. Programs for digitization and subsequent waveform analysis were written in the C programming language and 8086 assembly language. Data sets consisted of three passages from each patient.

An initial passage of SR/AF, NSR, or antero-gradual atrial activation was used to construct a ventricular or atrial electrogram template by signal averaging. The template was used for subsequent comparison with a second, separate passage of SR/AF, NSR, or antero-gradual control passage and a third passage of either VT (group 1), BBB (group 2), or retrograde atrial activation (group 3). A careful selection of window size effectively excluded any local repolarization in order to avoid the inclusion of injury current caused by temporary endocardial damage adjacent to the catheter. The template and the electrogram under analysis were compared using a best fit alignment.

In presenting both CWA and BAM, the following notation will be used: N = the number of points in the template; t_i = the template points; s_i = the signal points to be processed; \bar{t} = the template average; and \bar{s} = the signal average.

CWA

The correlation coefficient, ρ ,⁹ is independent of amplitude fluctuations, baseline changes, and produces an output between -1 and 1 . Mathematically, the correlation coefficient is defined as,

$$\rho = \frac{\sum_{i=1}^{i=N} (t_i - \bar{t})(s_i - \bar{s})}{\sqrt{\sum_{k=1}^{k=N} (t_k - \bar{t})^2} \sqrt{\sum_{k=1}^{k=N} (s_k - \bar{s})^2}}$$

BAM

BAM compares corresponding areas or bins constructed from the template with bins constructed from subsequent depolarizations using a simple error measure. Consecutive sample points are summed to estimate the areas using a rectangular area rule in equal sized bins. The average of these bin values is then removed resulting in a correction of baseline shift, and then these corrected bin values are normalized by the absolute sum of all corrected bin values. As a final step, the sum of the absolute difference of these normalized and corrected bins with an identically processed template is computed.

To form three-point bins, $S_1 = s_1 + s_2 + s_3$, $S_2 = s_4 + s_5 + s_6 \dots$ and $S_M = s_{N-2} + s_{N-1} + s_N$.

Template points, t_i , are processed similarly to form the T_i . For M equally sized bins in the template, the index of merit for BAM is given as:

$$\rho = 1 - \sum_{i=1}^{i=M} \left| \frac{T_i - \bar{T}}{\sum_{k=1}^{k=M} |T_k - \bar{T}|} - \frac{S_i - \bar{S}}{\sum_{k=1}^{k=M} |S_k - \bar{S}|} \right|$$

where

$$\bar{S} = \frac{1}{M} \sum_{k=1}^{k=M} S_k, \quad \text{and} \quad \bar{T} = \frac{1}{M} \sum_{k=1}^{k=M} T_k.$$

Because of the design of BAM, all template processing is performed in advance, i.e., prior to comparison of the template with subsequent electrograms under analysis. BAM is designed such

COMPUTER ANALYSIS OF ELECTROGRAMS

Table I
Patient Data for Discriminating VT from SR/AF

Patient	Sex	Heart Disease	Drugs	Sinus Rhythm/Atrial Fibrillation QRS Morphology	Ventricular Tachycardia QRS Morphology
1	M	CAD	None	SR-Normal	RBB-S/R
2	M	CAD	None	SR-Normal	LBB-S/L
3a	M	CAD	None	SR-LBBB	LBB-S/R
3b					LBB-S/L
4a	M	CAD	None	AF-LBBB	LBB-S/L
4b					LBB-S/L
5a	M	CAD	Proc	SR-Normal	RBB-I/R
5b					LBB-I/R
6a	M	CAD	Proc	SR-Normal	RBB-I/L
7a	F	CAD	Am	SR-Normal	RBB-S/L
7b					RBB-S/L
8	F	None	Am	SR-Normal	RBB-S/R
9	M	CAD	Qu Me	SR-Normal	LBB-S/L
10a	M	CAD	Am	SR-LBBB	LBB-S/L
11	M	VHD	Am	SR-LBBB	LBB-S/R
12	M	CAD	Am Me	SR-LBBB	LBB-I/R
13	M	CAD	En	SR-LBBB	LBB-I/L
14	M	CAD	Qu Me	SR-RBBB	RBB-S/R
15	M	CAD	Proc	SR-RBBB	LBB-S/R
16a	M	CAD	Am	SR-IVCD	RBB-I/R
16b					RBB-S/L
17	M	CAD	Am	SR-IVCD	LBB-S/L
18a	M	CAD	Am	SR-IVCD	†RBB-S/R
18b					†LBB-S/L
18c					†LBB-S/L
18d					†RBB-S/R
18e					†RBB-S/R
10b*	M	CAD	Am En	SR-IVCD	RBB-S/R
10c					LBB-I/L
6b*	M	CAD	En Proc	AF-IVCD	RBB-S/R
19	M	CAD	Qu Di	AF-IVCD	RBB-S/L

* These patients were studied on two different days with two different VT morphologies. † This patient had five distinct VT morphologies. CAD = coronary artery disease, VHD = valvular heart disease; Am = Amiodarone, Di = digoxin, En = encainide, Me = mexiletine, Proc = procainamide, Qu = quinidine; SR = sinus rhythm, AF = atrial fibrillation; LBBB = left bundle branch block, RBBB = right bundle branch block, IVCD = non-specific intraventricular conduction delay; LBB = left bundle branch morphology, RBB = right bundle branch morphology; I = inferior axis, S = superior axis; L = leftward, R = rightward.

Table II.

Patient Data for Discriminating Paroxysmal Bundle Branch Block of Supraventricular Origin From Normal Sinus Rhythm

Patient	Sex	Heart Disease	Drugs	Method of BBB Induction	Aberration Morphology
20	F	None	None	Spontaneous	RBBB
21	F	None	None	AOP	RBBB
22	F	None	None	AOP	RBBB
8	F	None	None	Spontaneous	RBBB
23	F	None	Iso	AOP	RBBB
24	F	None	Iso	Atrial Fibrillation	RBBB
25	F	None	Iso	Spontaneous	RBBB
26	M	COPD	None	AOP	RBBB
27	M	CAD	None	Spontaneous	RBBB
28	M	CAD	None	AOP	RBBB
29	M	CAD	Proc	AOP	RBBB
30	M	None	Iso	AOP	LBBB
31	M	CAD	None	AOP	LBBB

CAD = coronary artery disease, COPD = chronic obstructive pulmonary disease; Iso = isopril, Proc = procainamide; LBBB = left bundle branch block, RBBB = right bundle branch block; AOP = atrial overdrive pacing.

Table III

Patient Data for Discriminating Retrograde from Anterograde Atrial Activation

Patient	Sex	Heart Disease	Drugs	Atrial Electrode Location
23	F	None	Iso	RAA
32	M	None	None	RAA
33	M	None	None	RAA
34	F	None	None	RAA
35	M	CAD	None	RAA
36	M	None	None	RAA
37	M	None	None	RAA
38	F	None	None	RAA
26	M	COPD	None	RAA
39	F	None	None	RAA
40	M	CAD	None	RAA
41	M	CAD	Ve	RAA
8	F	None	None	RAA
42	M	None	None	RAA
43	F	None	None	HRA
44	M	None	Qu Ve	HRA
45	M	None	None	HRA
46	F	None	Ep	HRA
47	M	None	En	HRA

En = encainide, Ep = epinephrine, Iso = isopril, Qu = quinidine, Ve = verapamil, CAD = coronary artery disease, COPD = chronic obstructive pulmonary disease, RAA = right atrial appendage, HRA = high right atrium.

that it requires only one-sixth the computation of CWA for three-point bins. BAM is similar to another time-domain method of analysis-area of difference (AD).⁷ However, like CWA, BAM differs from AD in that it is independent of amplitude and baseline fluctuations.

Results

For either method, BAM or CWA, there was no universal threshold which separated VT from SR/AF, BBB from NSR, or retrograde atrial activation from anterograde atrial activation, in the patient population studied. Instead, patient specific thresholds were required for each method. Figure 1 summarizes the results of using CWA and BAM to distinguish VT from SR/AF. CWA and BAM both discriminated VT from SR/AF in 28/31 (90%) cases. Figure 2 summarizes the results of using CWA and BAM to distinguish BBB from NSR. CWA and BAM both discriminated BBB from NSR in 13/13 (100%) patients. Figure 3 summarizes the results of using CWA and BAM to distinguish retrograde atrial activation from anterograde atrial activation. CWA and BAM both discriminated retrograde from anterograde atrial activation in 19/19 (100%) patients.

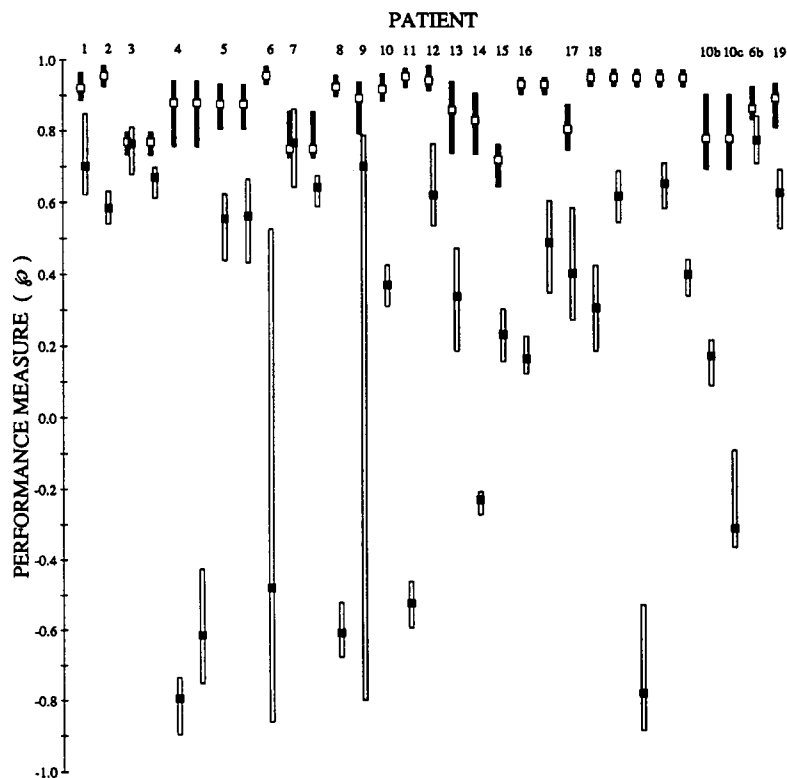
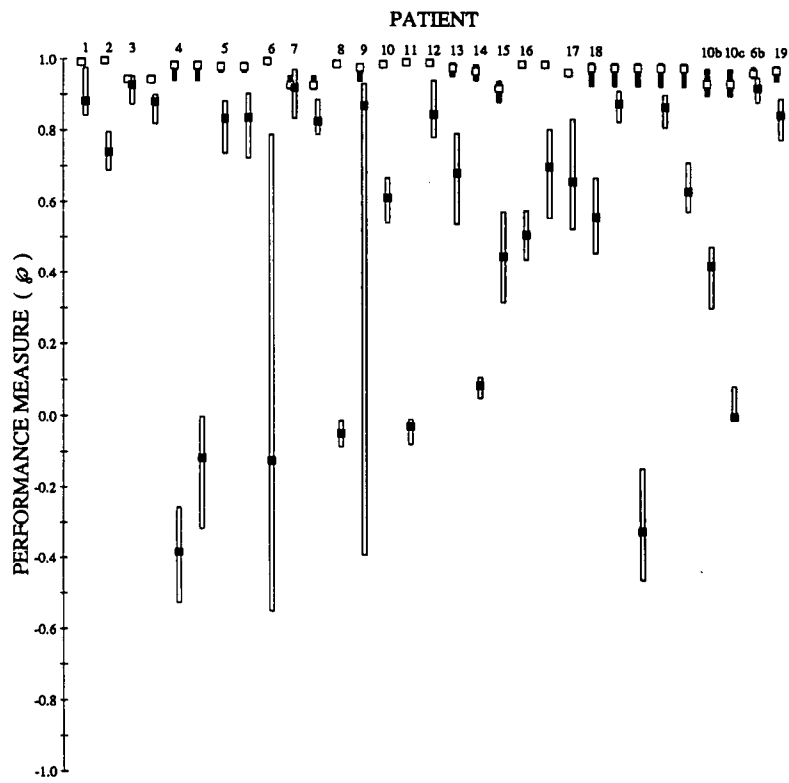


Figure 1. Results of CWA and BAM with three-point bins (A) (B) using the best fit alignment for distinguishing VT from SR (group 1). The ranges of ρ during SR/AF is shown in white, with a black box at the mean, while the ranges of ρ for VT is shown in black, with a white box at the mean.

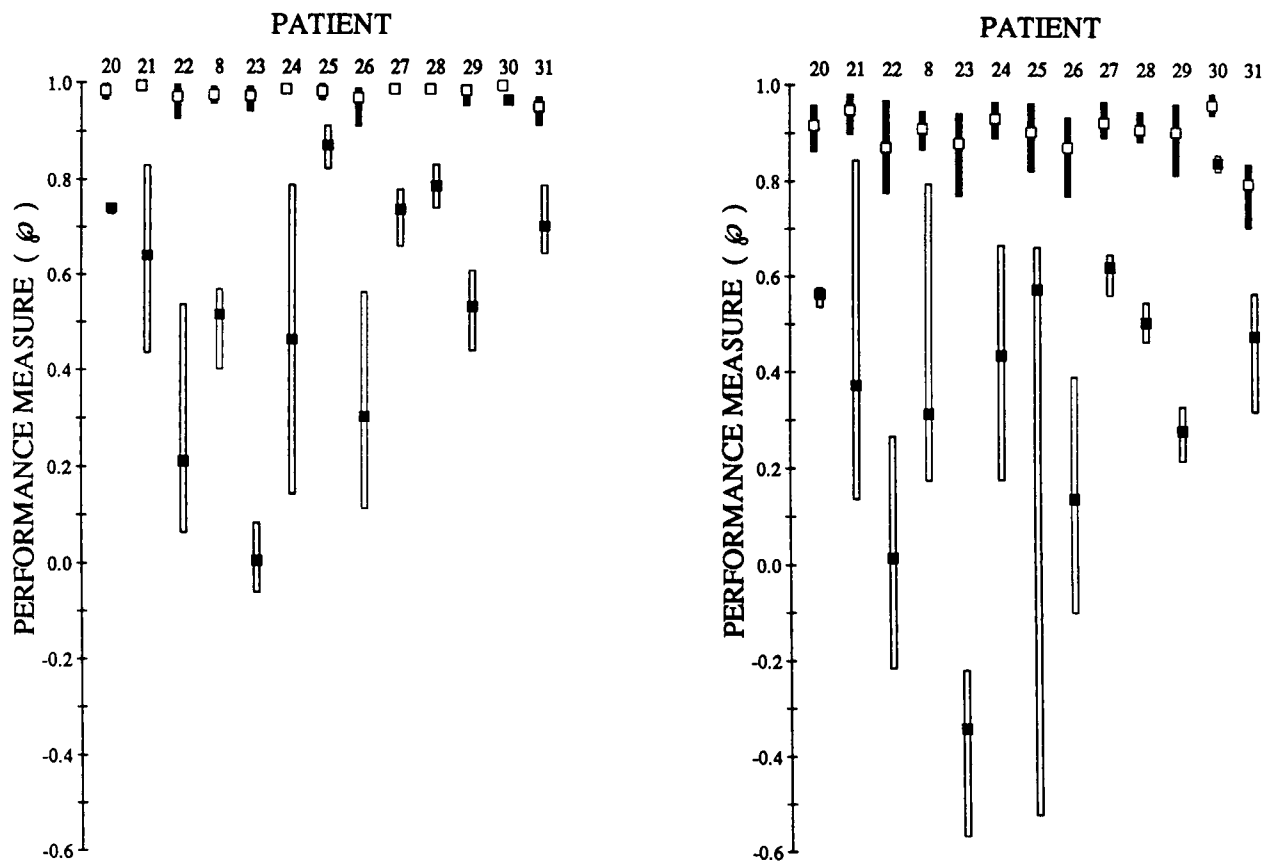


Figure 2. Results of CWA and BAM with three-point bins (A) (B) using the best fit alignment for distinguishing BBB from NSR (group 2). The ranges of ρ during NSR is shown in white, with a black box at the mean, while the ranges of ρ for BBB is shown in black, with a white box at the mean.

Discussion

Ideally, a universal threshold would be preferable to a patient-specific threshold for separating diverse cardiac dysrhythmias. However, the critical features of intracardiac electrogram morphology that determine accurate discrimination between diverse cardiac dysrhythmias remain elusive.

The present study was limited to an analysis of bipolar intracardiac electrograms. Whether bipolar electrograms are preferable for discriminating between dysrhythmias remains to be determined. Preliminary work from our laboratories would suggest that similar rates of success in discriminating VT from SR are achievable using time-domain analysis of either bipolar or unipolar elec-

trograms in a population of patients with inducible, sustained monomorphic VT. In individual patients within that population, however, either bipolar or unipolar analysis may be preferable for maximizing the difference between SR and VT electrograms.¹¹

The results of the present study demonstrate that time-domain techniques such as CWA are reliable for discrimination of retrograde atrial activation, paroxysmal BBB, and VT from SR. The similarity of performance of the BAM in this study supports the feasibility of developing alternative techniques which are also independent of electrogram amplitude and baseline fluctuations and have the added advantage of requiring less computational time without sacrificing diagnostic ac-

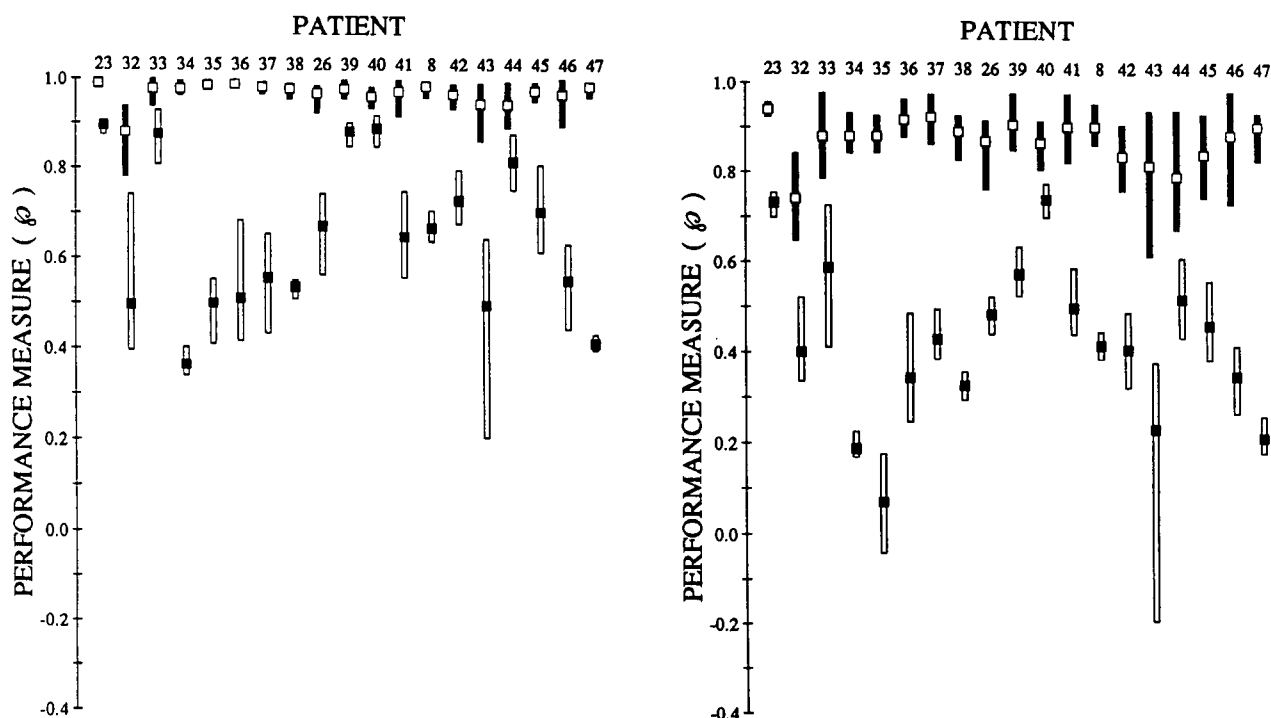


Figure 3. Results of CWA and BAM with three-point bins (A) (B) using the best fit alignment for distinguishing RAA from AAA (group 3). The ranges of ρ during AAA is shown in white, with a black box at the mean, while the ranges of ρ for RAA is shown in black, with a white box at the mean.

curacy.¹² Further reduction of time and power constraints may make real-time morphology analysis of intracardiac electrograms feasible for automatic antitachycardia and cardioverter-defibrillator devices.

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