

Two Types of Distribution Patterns of Bigeminy and Trigeminy in Long-Term ECG: a Model-Based Interpretation

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Abstract

Two types of distribution patterns of bigeminy and trigeminy are found in analysis of long-term ECG. To investigate the mechanism underlying this finding, a simplified equation of a modulated parasystole model was used and symbolic solutions for cyclic VPC patterns were obtained. The map of these solutions in a model parameter plane showed two different solutions for bigeminy, four kinds of trigeminy, two kinds of quadrigeminy, combinations of these patterns, and apparently normal ECG. These results are used to explain the features of the distribution patterns.

1. Introduction

1.1. Background and purpose

The rate of appearance of ventricular premature contraction (VPC) depends on heart rate (HR) in analyses of a Holter ECG and in exercise stress tests. Different patterns of HR-dependency in VPC have been reported: an increase in VPC frequency as HR increases, a decrease in VPC frequency, non-monotonous changes, and disappearance of VPC at a specific HR. Nakata and Takayanagi reported two types of distribution patterns of bigeminy and trigeminy in plots of VPC frequency in a Holter ECG using the coupling interval between each VPC and the next normal sinus beat (XN) as the abscissa, where “X” designates a VPC and “N” is a normal sinus beat [1-3]. On this histogram, a trigeminy-bigeminy order of appearance (with increasing XN) was defined as the standard type and a bigeminy-trigeminy order as the reverse type. The purpose of the current study is to provide an interpretation of this phenomenon using a mathematical model of parasystole.

1.2. Two types of distribution patterns of bigeminy and trigeminy

ECG recordings are shown in Figure 1. In the X-N sequence, a pattern XNX is defined as bigeminy and XNNX as trigeminy. Figure 1 shows two cases of Holter ECGs. In Figure 1A, the XN interval for bigeminy is larger than that for trigeminy, and in Figure 1B, the XN interval for trigeminy is larger than that for bigeminy. Histograms of bigeminy (blue) and trigeminy (yellow) over long ECG recordings are shown in Figures 2 and 3.

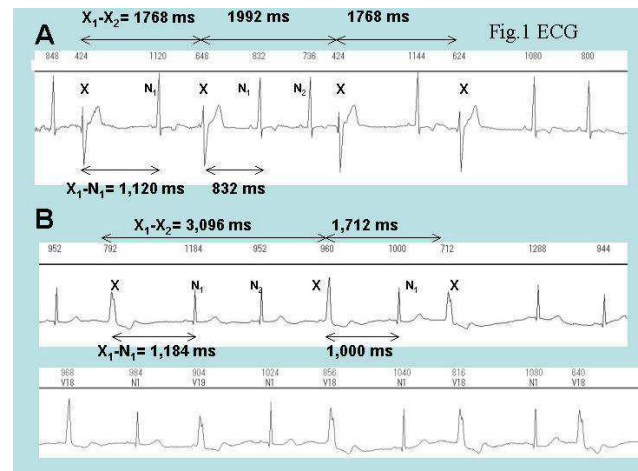


Figure 1. Two types of bigeminy-trigeminy patterns. A: standard type in which the bigeminy distribution is centered at a larger XN, and B: reverse type in which the bigeminy distribution occurs at a smaller XN.

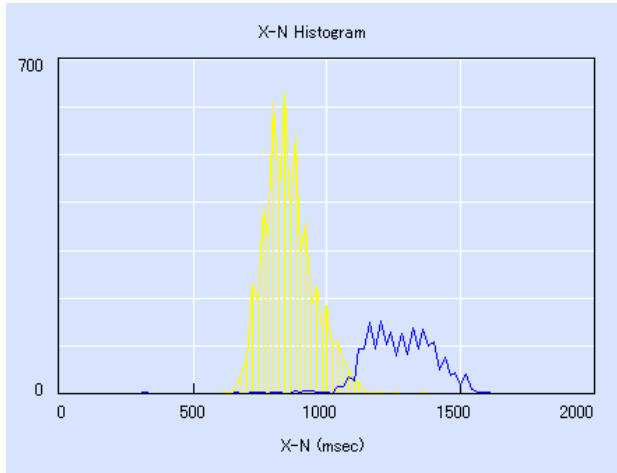


Figure 2. Histogram of bigeminy (blue) and trigeminy (yellow) for XN values (standard type).

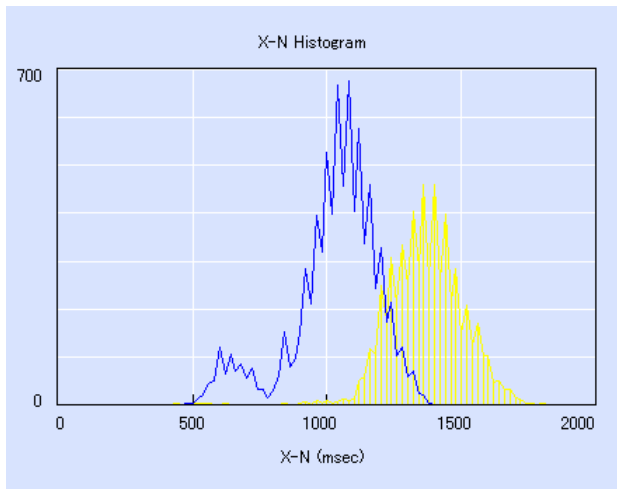


Figure 3. Histogram of bigeminy (blue) and trigeminy (yellow) for XN values (reverse type).

Two types of patterns were distinguished: the standard type (Figure 2) in which the bigeminy distribution is centered at a larger XN, and the reverse type (Figure 3) in which the bigeminy distribution occurs at a smaller XN.

2. Methods

We assumed a *modulated parasystole model* and formulated the model as a system of difference equations in a phase of the sinus pacemaker's

stimulus on an ectopic pacemaker. Assuming that the variable t_n is the n -th phase of the ventricular excitation induced by the sinus pacemaker in the oscillation of the ectopic pacemaker, the following equations are obtained between the successive phases, t_n and t_{n+1} [4, 5].

$$t_n + Ts - PRC(t_n) \quad (0 \leq t_n < Te) \quad (1N)$$

$$t_{n+1} = t_n - Te + Ts \quad (Te \leq t_n < Te + (1-\gamma)Ts) \quad (1X)$$

$$t_n - Te \quad (Te + (1-\gamma)Ts \leq t_n < Te + Ts) \quad (1.)$$

where Ts is the intrinsic period of the sinus pacemaker, Te is the intrinsic period of the ectopic pacemaker, and γTs is the refractory period due to the sinus R wave. The function $PRC(t)$ represents the phase response curve (PRC) of an ectopic pacemaker.

Approximating the PRC function by

$$PRC(t) = \begin{cases} at & (t \leq wTe) \\ b(t - Te) & (t > wTe), \end{cases}$$

where w is a break point of the PRC, equation (1N) becomes

$$t_{n+1} = \begin{cases} (1-a)t_n + Ts & (0 \leq t_n < wTe) \quad (1a) \\ (1-b)t_n + bTe + Ts & (wTe \leq t_n < Te) \quad (1b) \end{cases}$$

and equation (1) becomes

$$(1-a)t_n + Ts \quad (0 \leq t_n < wTe) \quad (1a)$$

$$(1-b)t_n + bTe + Ts \quad (wTe \leq t_n < Te) \quad (1b)$$

$$t_{n+1} = t_n - Te + Ts \quad (Te \leq t_n < Te + (1-\gamma)Ts) \quad (1X)$$

$$t_n - Te \quad (Te + (1-\gamma)Ts \leq t_n < Te + Ts) \quad (1.)$$

Equations (1a) and (1b) correspond to a normal beat, equation (1X), an ectopic beat, and equation (1.), a silent ectopic beat that is concealed under the refractory period of the ventricle. We define a cyclic solution with period n of (1) to be a sequence $\{t_0, t_1, \dots, t_n\}$ with $t_n = t_0$, and the solution is represented as $\{bX\}$, $\{abX\}$, $\{b.aX\}$, etc., using symbols that correspond to the branches of equations (1a), (1b), (1X) and (1.).

A symbolic formula processor (MathCad8, MathWorks Inc.) was used to obtain stable solutions of ECG patterns.

3. Results

3.1. Regions of stable solutions

Theoretically, we obtained stable solutions for bigeminy ($\{bX\}$ and $\{aX\}$), trigeminy ($\{bbX\}$, $\{abX\}$, $\{b.bX\}$ and $\{b.aX\}$), quadrigeminy, and apparently normal ECG. Patterns of bigeminy and trigeminy and the corresponding regions with respect to T_s were obtained as follows:

Bigeminy $\{bX\}$:

$$\max[(1-b+w b)Te/2, (1-b)Te/(2-b)] \leq T_s < (1-b)Te/(2-2b+\gamma) \quad (2)$$

Bigeminy $\{aX\}$:

$$Te/(2-a) \leq T_s < \min[(1+wa)Te/2, Te/(2-2a+\gamma)] \quad (3)$$

Trigeminy $\{bbX\}$:

$$\max[(1-b+w b)Te/2, (1-b)Te/(2-b)] \leq T_s < \min[(1+wa)Te/2, Te/(2-2a+\gamma)] (1-b)Te/(2-2b+\gamma) \quad (4)$$

Trigeminy $\{abX\}$:

$$\max[(1-b)Te/(3-b), (1-a-b+ab+w(a+b-ab))Te/(3-2b), (1-b)Te/(3-a-2b+ab)] \leq T_s < \min[(1-b+w(a+b-ab))Te/(3-b), Te/(3-2a), (1-b)Te/(3-2a+2ab+\gamma(a+b-ab))] \quad (5)$$

Trigeminy $\{b.bX\}$:

$$\max[(2-b)(1-b+w b)Te/(3-b), (1-b)(2-b)Te/(3-4b+b^2+\gamma(2-b)), (2-b)(1-b+wb)Te/(3-2b), (1-b)(2-b)Te/(3-4b+b^2+\gamma(2-b))] \leq T_s < \min[(2-b)Te/(3-b), (1-b)(2-b)Te/(3-5b+2b^2+\gamma(2-b))] \quad (6)$$

Trigeminy $\{b.aX\}$:

$$\max[(2-b)(1-b+w b)Te/(3-b), (1-b)Te/(3-2b-(1-\gamma)(a+b-ab)), (1-b)Te/(3-2b), (2-a-b+ab)Te/(3-2a-b+ab)] \leq T_s < \min[2Te/(3-a), (2-2b+w(a-b+ab))Te/(3-2a), (2-a-b+ab)Te/(3-a+(2-\gamma)(a+b-ab))] \quad (7)$$

etc.

A map of these regions was drawn as two-dimensional areas in the $T_s/Te - w$ plane (Figure 4). We also derived a linear relationship between XN and T_s as follows: for the solution $\{bC\}$ to exist, $t_0 = XN$ satisfies $wTe \leq t_0 < Te$, $t_1 = (1-b)t_0 + bTe + Ts$ obtained from equation (1b) satisfies $Te \leq t_1 < Te + (1-\gamma)Ts$, and $t_2 = t_1 - Te + Ts$ obtained from equation (1X) must satisfy $t_2 = t_0$. Solving these equations gives region (2) for T_s , and the relationship between $XN (= t_0)$ and T_s as follows:

$$XN = (2Ts - (1-b)Te)/b \quad (6).$$

Similar relationships can be obtained for other solutions.

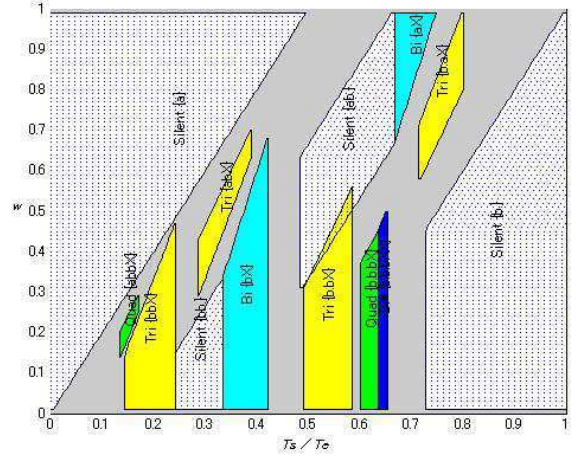


Figure 4. Regions of stable solutions in the $T_s/Te - w$ plane, including two of bigeminy (cyan), four of trigeminy (yellow), and two of quadrigeminy (green). Between these regions, combinations of the patterns (gray) and silent patterns (dots) are shown. Parameters: $a = b = 0.5$ and $\gamma = 0.375$.

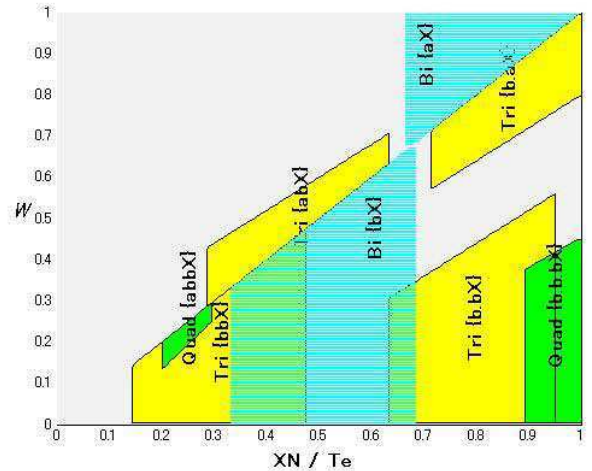


Figure 5. Regions of bigeminy (cyan), trigeminy (yellow) and quadrigeminy (green) in the $XN/Te - w$ plane.

Thus we have a solution map in the $XN/Te-w$ plane (Figure 5). Clinically observed VPC distributions were explained from this map: when XN/Te is small, bigeminy {bX} is preceded by trigeminy {bbX} (for small w) or by {abX} (for large w) as XN increases, the standard type pattern (Figure 2); and when XN/Te is large, bigeminy {bX} is followed by trigeminy {b.bX}, the reverse type pattern (Figure 3).

3.2. Computer simulation

The results shown in Figures 4 and 5 indicate regions of steady-state solutions of equation (1), i.e., under conditions in which the model parameters were held constant. Clinically observed distributions (Figures 2 and 3) arose from conditions of long-term variation of biological parameters. To simulate these results, computer simulations were performed using equation (1) with random fluctuations of the parameters Ts and w . The results are shown in Figure 6 as a histogram of frequency of bigeminy and trigeminy.

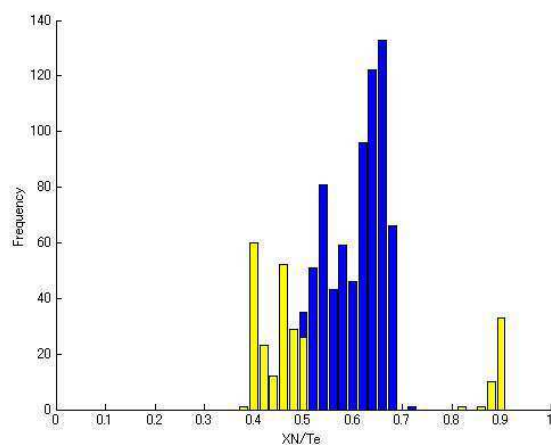


Figure 6. Histogram of bigeminy (blue) and trigeminy (yellow) for XN values obtained from the computer simulation.

The values of Ts and w were changed randomly at a certain period of time according to a normal distribution. The two types of distributions (standard and reverse) appear in the same graph because of the wide variation in parameter values.

4. Discussion and conclusions

We have shown that the two clinically observed types of distribution patterns of bigeminy and trigeminy and their related properties can be explained by the modulated parasystole model, although other mechanisms such as reentry and delayed afterdepolarization cannot be excluded.

Acknowledgements

This study was supported by a grant from Kitasato University School of Allied Health Sciences (Grant-in-Aid for Research Project, No 2007-1007).

References

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