

Clinical tissue characterization: online determination of magnitude and time delay of myocardial backscatter

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Implementation of an algorithm for the automated determination of the magnitude and time delay of the cardiac cycle-dependent variation of integrated backscatter is reported. This algorithm has been implemented in the experimental firmware of a commercially-available echocardiographic imaging system. Integrated backscatter and the cardiac cycle-dependent variation (cyclic variation) of integrated backscatter are described and their roles in myocardial tissue characterization are discussed. A brief description of the algorithm used for the determination of the magnitude and time delay is given, followed by accounts of the process of collecting cyclic variation data and the algorithm implementation on the cardiac system. This implementation demonstrates how tissue characterization techniques could be used to augment diagnostic ultrasound and may facilitate the further investigation of the diagnostic potential of the cyclic variation of myocardial backscatter.

INTRODUCTION

An evolving application of ultrasonic examination of the heart is tissue characterization, the goal of which is to classify the health or pathology of myocardial segments based on the intrinsic interaction of ultrasound with the tissue.¹ The specific approach to tissue characterization followed in this work involves the derivation of quantitative parameters from the radio frequency signals received due to the scattering of ultrasound from tissue. These derived parameters should be reproducible and independent of instrumentation. The utilization of tissue characterization methods in a clinical setting requires the rapid determination of these quantitative indices from which the physical state of the tissue in question can be ascertained. In this article, we describe and demonstrate the implementation of an algorithm for the automated determination of the magnitude and time

delay of the cardiac cycle-dependent variation of integrated backscatter in experimental firmware on a commercially-available cardiac imaging system. This pilot implementation can facilitate the evaluation of the magnitude and time delay indices as clinically-relevant parameters for tissue characterization and demonstrate the role that tissue characterization can play in augmenting the diagnostic capabilities of clinical ultrasound.

INTEGRATED BACKSCATTER

Integrated backscatter is a measure of the acoustic energy returned to the transmitting transducer due to the scattering of the ultrasound by a tissue segment. For the purposes of myocardial tissue characterization, the acoustic backscatter signals of interest are those emanating from the intramural myocardium, avoiding signals due to tissue boundaries (i.e., blood-tissue or tissue-pericardial interfaces). This scattering of ultrasound by intramural myocardium is due to the small-scale differences in the mechanical properties among the material constituents of the tissue. The determination of integrated backscatter from the raw signal backscattered by tissue is a multistep process. First, a gate is used to isolate the portion of the signal that is due to

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the backscatter from the tissue of interest. This gated signal represents the scattering from a volume of tissue that is defined by the length of the gate and the effective cross-sectional area of the insonifying ultrasonic beam. In one approach, a Fourier decomposition of this gated signal is performed. This allows the determination of the signal's power spectrum, i.e., the acoustic power backscattered to the transducer as a function of the respective temporal frequencies that comprise the signal. (An alternate approach to determining integrated backscatter using only time-domain techniques is described in the accompanying video).

In order to determine the absolute level of energy contained in the signal, its power spectrum is normalized by the spectrum from a reference of known backscatter strength, usually an ultrasonic mirror such as a steel plate in water. The resulting power curve is then averaged (i.e., integrated) over the usable frequency range of the signal to obtain the integrated backscatter value. This averaging over frequency serves to minimize the influences of interference and phase cancellation effects that arise due to the presence of many scattering sites within the interrogated tissue volume.

Typically, the integrated backscatter values from nearby insonified tissue volumes are averaged to obtain a single value over some localized region. The spatial extent of the region will depend on the center frequency of the transmitted ultrasound. For clinical scanners, where the center frequency is in the 2 to 5 MHz range, the extent of spatial averaging used is on the order of several millimeters in each direction. This technique of spatial averaging, while reducing localization somewhat, aids significantly in making the measurements stable and reproducible.

In vitro and *in vivo* studies of both animal and human hearts have shown that integrated backscatter and other related acoustic indices differentiate among acutely ischemic tissue, old infarct and normal tissue, and correlate with collagen content and other measures of tissue state.²⁻²⁵ Studies of humans have also shown that integrated backscatter can be used to delineate blood-tissue boundaries in real-time, facilitating online analysis of cavity areas and volumes during echocardiographic examination.²⁶⁻²⁸

CYCLIC VARIATION OF INTEGRATED BACKSCATTER

In addition to the correlation of absolute integrated backscatter (i.e., normalized by some reference scatterer or reflector) with tissue properties, the integrated backscatter from myocardium is observed to undergo a characteristic variation over the cardiac cycle.²⁹⁻³² This pattern of cyclic variation is characterized by two distinct regimes over the heart cycle. In general, the integrated backscatter level approximates a relative maximum value for the majority of the cycle and falls to a minimum level, or trough, for a time duration roughly equivalent to the systolic interval (Fig. 1). The position of this trough in the variation relative to the heart cycle can vary depending on the details of the acquisition and the state of the insonified tissue. Changes in the pattern of cyclic variation have been correlated with changes in cardiac function and specific pathologic myocardial states.³³⁻³⁹ In order to yield quantitative comparisons, cyclic variation data are characterized by two parameters, the magnitude and time delay. The magnitude is a measure of the difference in the respective average integrated backscatter values between the minimum and maximum levels of the variation (Fig. 2). Because integrated backscatter values are expressed on a decibel (dB) scale, the magnitude is similarly expressed in decibels. The time delay begins as a measure of the time

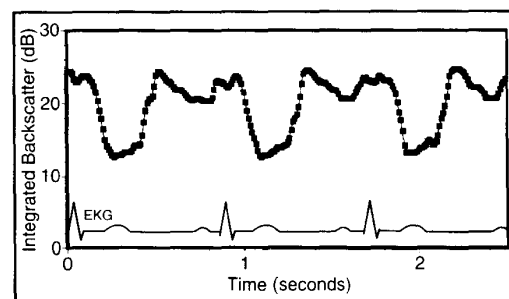


Fig. 1 Cyclic variation of integrated backscatter over three heart cycles. This data set illustrates the two-level pattern that is characteristic of cyclic variation. For the majority of each heart cycle, the backscatter fluctuates about a baseline value and then dips into a "trough" or minimum level for a period of time on the order of the systolic interval. This data was collected from the left-ventricular posterior wall of a normal human volunteer using an echocardiographic imaging system with real-time integrated backscatter processing.

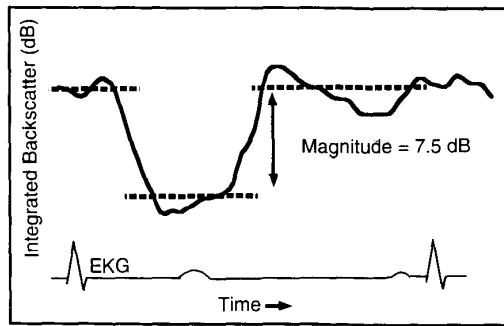


Fig. 2 Magnitude of cyclic variation. The magnitude is a measure of the size of the primary variation in the integrated backscatter data over the heart cycle. The magnitude is determined by taking the difference between the average integrated backscatter levels of the baseline region and the trough, respectively.

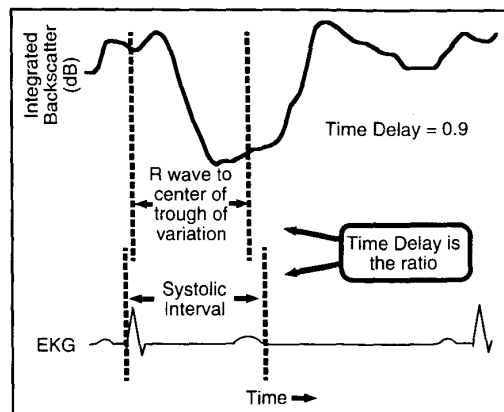


Fig. 3 Time delay of cyclic variation. The time delay measures the offset of the center of the trough of variation from the R wave of the EKG. This time interval is then normalized by the systolic interval to arrive at the dimensionless time delay.

interval from the R-wave of a simultaneously-recorded EKG to the center of the minimum-value phase of the variation curve. This time interval is then normalized (i.e., divided) by the systolic interval to yield a dimensionless parameter (Fig. 3). Many studies in both humans and experimental animals have shown correlation between the cyclic variation parameters and physical state of the tissue.^{8,21,25,33-37,40-55} One such study in dogs showed that the magnitude and time delay values return to normal faster than the recovery of wall thickening during the reperfusion of previously ischemic but viable tissue.⁵⁶ Transthoracic studies in humans have also shown that magnitude and time delay of

cyclic variation can distinguish between normal and infarcted tissue as well as normal and acutely ischemic tissue.^{42,45,47,57} A further study has also suggested that the magnitude and delay of the cyclic variation may be a more sensitive indicator than conventional echocardiographic measurements for the detection of the onset of diabetic heart disease.⁴⁹ Because the quantitative analysis of cyclic variation data only requires the consideration of relative changes in integrated backscatter rather than its absolute determination, it is particularly attractive for use in situations where ascertaining absolute levels of backscatter may be difficult.⁵⁸⁻⁶⁰

Initial investigations of the cyclic variation showed a maximum in integrated backscatter to occur near end-diastole and a minimum to occur at end-systole. This observation led to the use of a two-point analysis of cyclic variation in which integrated backscatter values near end-diastole and end-systole were taken as maximum and minimum values of the backscatter curve. This method of analysis resulted in a magnitude-only parameterization of cyclic variation, with no explicit recognition of the timing aspects of the variation. However, subsequent work has shown that this relationship strictly held only under specific conditions. Thus, the two-point end-systolic/end-diastolic method of analysis is inadequate for several reasons. Due to the organization of its muscle fibers, the heart has anisotropic acoustic properties, i.e., the interaction of ultrasound and myocardium depends on the angle of incidence of the sound beam relative to the orientation of the myofibers.^{9,43,61-68} Thus, the amount of backscatter produced by a tissue segment depends on the direction from which it is insonified. Recent work suggests that normal myocardium exhibits anisotropy in its pattern of cyclic variation as well.⁶⁹⁻⁷¹ This work shows that the minimum and maximum levels of the cyclic variation pattern from healthy segments are not always coincident with end-systole and end-diastole, respectively, for directions of insonification parallel or oblique to local myofibers. The state of health of the tissue under study also influences the position of the two levels in the variation curve relative to the cardiac cycle. Furthermore, the use of only two points from the backscatter curve for measurement purposes is

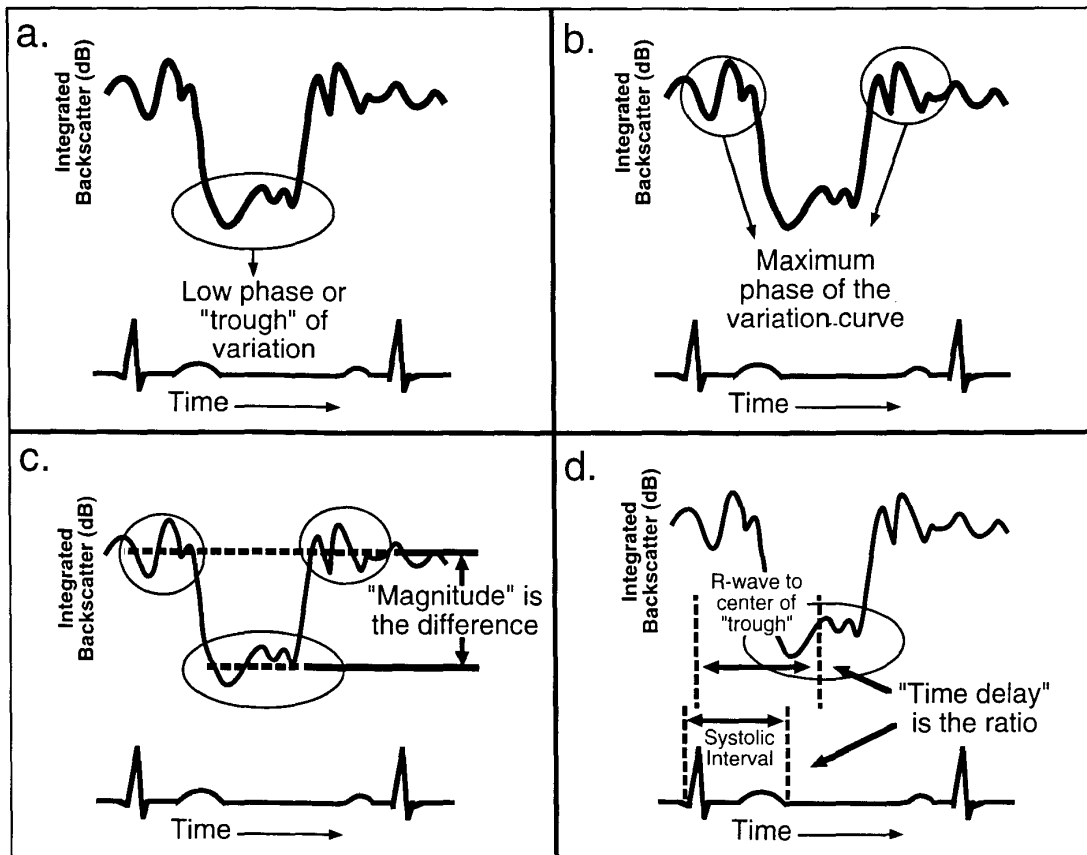


Fig. 4 Interpretation of cyclic variation data. (a) Identification of the low phase (or trough) of the cyclic variation curve. This is the portion of the curve that, on average, lies below the mean backscatter level over the heart cycle. (b) Identification of the maximal portion of the curve. This is the portion of the curve that, on average, lies above the mean backscatter level over the heart cycle. (c) Average backscatter values over the two regions are determined. The magnitude of variation is the difference in the two average integrated backscatter values. (d) Time delay determination begins by finding the time interval from the R-wave of the EKG to the center of the trough of the variation. This interval is then divided by the time duration of systole to complete the time delay calculation.

more susceptible to measurement noise and small scale variations than a quantitative analysis that employs the entire data curve to extract the size of the primary variation. Also, the explicit inclusion of the timing information, which we quantify as the normalized time delay, complements the magnitude parameterization of the cyclic variation data augmenting its diagnostic potential.⁷² Thus, we advocate the interpretation of cyclic variation data by identifying the following features of the integrated backscatter versus time curve: (1) a region of the curve with a time duration on the order of the systolic interval for which the backscatter values fall below the mean value of the entire data set; and (2) the region of the data set where the data lie

above the mean data set value. After identification, the average integrated backscatter value for each region is determined and the absolute difference between the two levels is then the magnitude of variation for the data. Additionally, the position in time of the center of the region of reduced backscatter relative to end-diastole is determined. This time interval is divided by the systolic interval duration to determine the time delay (Fig. 4).

In general, abnormal myocardium exhibits a decrease in magnitude and an increase in time delay in comparison with healthy tissue. The range of values for normal tissue varies depending on the segment insonified and the direction of interrogation. Typical values for the magni-

tude and time delay of normal myocardium imaged in standard echocardiographic views have been reported by Finch-Johnston et al.⁷⁰

ALGORITHM FOR DETERMINING MAGNITUDE AND TIME DELAY OF CYCLIC VARIATION

In order to permit an objective analysis of cyclically-varying integrated backscatter data, an algorithm was developed for the automated extraction of the magnitude and time delay.⁷³ In addition to the variation curve, the algorithm also requires some independent indicator of the timing of the cardiac cycle relative to the backscatter data, which can be provided by a simultaneously-recorded EKG. An idealized model of the variation is employed to determine the parameters. This model exhibits two integrated backscatter levels over the heart cycle, with the duration of the minimum level equivalent to the systolic interval (as approximated by the QT interval from the EKG). The initial values for the two levels of the model function are chosen such that the model has a zero average over the heart cycle and a magnitude (difference in integrated backscatter value between the two levels) of unity. These specific choices are made to avoid complications in the time delay determination and to simplify the magnitude calculations. Using Fourier analysis, the amplitude of variation of the model is compared with that of the backscatter data, resulting in the extraction of the magnitude. By performing a cross-correlation, the position of the minimum phase of the model is adjusted to best fit the data. This operation in concert with timing information from the EKG (position of R-wave and length of QT interval) yields the necessary input to determine the time delay parameter. Earlier versions of the algorithm have undergone validation tests, pitting the best manual estimates versus the automated determinations of the magnitude and time delay.^{73,74} These studies have confirmed the utility and accuracy of the algorithm. However, in a fraction of the cases studied, the algorithm produced clear overestimates of the magnitude.⁷⁵ The latest version of the algorithm, demonstrated in the accompanying video, includes an additional step in the magnitude determination that

extends its applicability to a broader range of cyclic variation data. This current version can detect the special conditions under which the older reported versions^{73,74} would produce overestimates of the magnitude. When these conditions are present, this updated algorithm implements a correction procedure that results in a significantly more accurate magnitude determination.

IMPLEMENTATION AND USE OF ALGORITHM ON A MEDICAL IMAGING SYSTEM

The algorithm has been implemented in experimental firmware on the Hewlett-Packard SONOS 1500™ cardiac imaging system. The real-time integrated backscatter processing capability of that system allows for the online acquisition of integrated backscatter data as a function of time. In contrast with the frequency domain (Fourier transform) method described above, the system employs a time-domain sum-of-squares approach to the estimation of integrated backscatter using an integration time of 3.2 μs .^{76,77} (For more detail on the time-domain approach and real-time integrated backscatter, see the accompanying video). The algorithm resides within the M-mode integrated backscatter operating environment of the system. In this operating environment, the system produces M-mode images in which each pixel has an assigned integrated backscatter value. To begin the data acquisition process, the operator selects a specific line from the two-dimensional sector image that intersects the tissue segment of interest. This line is then used in M-mode to form the desired real-time integrated backscatter images. The system can hold approximately 30 seconds of M-mode integrated backscatter image data. An EKG is displayed and recorded along with the image data. Once the operator has acquired satisfactory images, several heart cycles are selected out of the many obtained for the generation of the cyclic variation data. After the cycles of interest are chosen and displayed on-screen, the operator moves a region-of-interest cursor (ROI) through the M-mode representation of the myocardial tissue segment under investigation. All of the integrated backscatter

values assigned to the image pixels that fall within the ROI are averaged, producing a single integrated backscatter value. The ROI is square in shape and its size can be varied. The size is chosen to be as large as possible, to achieve good spatial averaging, while staying entirely within the myocardium to avoid corruption from specular reflections at the tissue boundaries. An integrated backscatter trace is then simultaneously acquired and plotted on-screen by driving the ROI, via a trackball, through the image of the segment of interest.

Once the integrated backscatter data is obtained, the algorithm process can be activated. The operator confirms that the algorithm processing is active and selects the number of consecutive heart cycles of the data that they wish to analyze. Once the preliminary settings are approved, the operator begins the process of entering the timing information. Using the cursor under trackball control, the operator enters the timing information by marking the following locations on the EKG for the consecutive heart cycles to be analyzed: onset of QRS, end of T-wave, and the R-wave. In addition, the QRS following the last cycle of interest is marked to frame the data of interest. After the timing markers have been set and approved, the calculations of the cyclic variation parameters begin. At the conclusion of the calculations, the results for the magnitude and time delay are displayed automatically in a results window and the computed model fit is superimposed on the displayed integrated backscatter data plot.

The implementation reported here is expected to facilitate the evaluation of the magnitude and time delay indices as clinically relevant parameters for tissue characterization. It is hoped that these efforts will help to establish myocardial tissue characterization as a complement to diagnostic echocardiography that can be readily performed at the ambulatory patient lab or at the bedside of critically-ill patients with myocardial dysfunction.

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Clinical tissue characterization: online determination of magnitude and time delay of myocardial backscatter

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Implementation of an algorithm for the automated determination of the magnitude and time delay of the cardiac cycle-dependent variation of integrated backscatter is reported. This algorithm has been implemented in the experimental firmware of a commercially-available echocardiographic imaging system. Integrated backscatter and the cardiac cycle-dependent variation (cyclic variation) of integrated backscatter are described and their roles in myocardial tissue characterization are discussed. A brief description of the algorithm used for the determination of the magnitude and time delay is given, followed by accounts of the process of collecting cyclic variation data and the algorithm implementation on the cardiac system. This implementation demonstrates how tissue characterization techniques could be used to augment diagnostic ultrasound and may facilitate the further investigation of the diagnostic potential of the cyclic variation of myocardial backscatter.