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A Survey of Different Compression Technique for Biomedical Signal (ECG)

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ABSTRACT: Capacity and transmission impediments have made biomedical flag information pressure an imperative element for most biomedical automated frameworks. Electrocardiogram (ECG) flag is a critical measure to know the Heart genuine conditions so that effectively discovered expires. Different methods have been proposed throughout the years for tending to the issue. In our paper, we are breaking down various methodologies of biomedical flag pressure, for example, ECG motion in secure correspondence network.ECG signs are gathered both over drawn out stretches of time and at high determination from patients. This makes considerable volumes of information for capacity and transmission. Information pressure tries to lessen the quantity of bits of data required to store or transmit digitized ECG signals without critical loss of flag quality. An extensive variety of pressure procedures in view of various change methods like MSVQ, hereditary enhancement, SPARSE 2D SEPARABLE TRANSFORM. DST and DCT were assessed to locate an ideal pressure system for ECG information pressure. All Testing was performed on falsely motions from the standard CSE and MIT-BIH database.

KEYWORDS: ECG, MSVQ, Sparse 2D, DST, DCT

I. INTRODUCTION

An electrocardiogram is utilized to screen your heart. Each beat of your heart is activated by an electrical motivation typically created from exceptional cells in the upper right council of your heart. An electrocardiogram — likewise called an ECG or EKG — records these electrical flags as they go through your heart.



Biomedical signs can be packed in time area, recurrence space, or time-recurrence space. ECG information pressure calculations have been for the most part arranged into three noteworthy classes [3]: 1) Direct time-space strategies, e.g., defining moment (TP), plentifulness zone-time age coding (AZTEC) [4], facilitate lessening time encoding framework (CORTES) and Fan calculation. 2) Transformational approaches [3], e.g., discrete cosines change (DCT), quick Fourier change (FFT), discrete sine change (DST), wavelet change (WT) and so forth 3) Parameter extraction strategies, e.g., Prediction and Vector Quantization (VQ) techniques [2]. The time space systems which depend on direct techniques were the prior ways to deal with biomedical flag pressure. Change Coding (TC) is the most critical recurrence area advanced waveform pressure technique. When we look at these strategies we locate that immediate information pressure is a period space pressure calculation which straightforwardly examinations tests where between beat and, intra-beat relationship is abused. In this paper distinctive changes like MSVQ, Sparse 2D, DST, DCT and are examined for ECG flag pressure.





Fig.2: ECG signal before compression

II. EVALUATION CRITERIA & COMPRESSION RATIO

The evaluation of performance for testing ECG compression algorithms includes followings three components: First one is compression efficiency second is reconstruction error and third is computational complexity. The compression efficiency is given by compression ratio (CR).

 $CR = \frac{Number of samples before compression}{Number of samples after compression}$

III. DIFFERENT TECHNIQUE

A. MSVQ

The motivation for introducing MSVQ for ECG signal compression is determined from the observation of the waveform itself. If the time sequences of ECG signals are analyzed, it is possible to conclude that many short-length segments differ mainly in their mean values, while their wave shapes have less variability. Following this reasoning, the compression system described here defines vectors as segments from the sequence of samples in a digitized ECG signal, and then calculates and subtracts the mean value from each vector. The subsequent signal processing is as outlined above, and is summarized below in four steps.

1) An ECG signal sampled at an adequate rate is appropriately segmented into vectors.

2) The mean is subtracted from the vectors created in 1) and a training sequence is derived from a very large number of segments.

B. SPARSE 2D

The basic idea of sparse 2D is achieving an enhanced sparse representation by grouping similar segments of the input ECG signal into 2D arrays, and then using a 2D transformation (which can be complete or over complete) to transform 2D arrays. A simple justification for the effectiveness of the idea is as follow:

• Assume that the grouping is done, i.e. similar segments are placed in groups and a 1D transformation (A) is used for each group.

• In each group we have similar segments and hence after transformation we will have the same number of high magnitude coefficients for each segment in a group, say high-magnitude coefficients for each segment.

• Assuming n segments in each group, we will have n_high-magnitude coefficients in that group. In other words this group can be represented by n_ coefficients.

• Now we should perform another 1D transform (B) on the second dimension (along each row) of each group.

• Components of this row are similar (because only similar segments are in this group), i.e. there is a kind of similarity for all members of the row.

C. GENETIC SEGMENTATION

The genetic segmentation of ECG signals discussed here can be viewed as a fundamental introductory phase for a number of detailed compression schemes. It is worth stressing that the genetic segmentation of the signal does not confine itself to a specific form of compression such as a piecewise linear approximation or any other technique by that matter. We have used the model of linear approximation realized within the individual segments. Obviously, the segmentation can serve as a starting point when proceeding with more sophisticated compression schemes, say via quadratic functions or higher order polynomials. One should emphasize that the genetic segmentation is concerned with a global optimization and this stands in sharp contrast with the most compression techniques. Simply, in genetic



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segmentation once we decide upon their number of segments, the optimization of the segments looks at all data *globally*. There are several possible enhancements and extensions of the approach introduced in this study. First, which is usually a crucial point in genetic optimization; one can augment the form of the fitness function. Here the GA mechanisms were guided by the absolute values of the differences between the external values (minimal and maximal) of the derivatives or their estimates. One can envision situations (especially in case of highly noisy data) that these extreme values become too sensitive and can be highly corrupted by noise. To alleviate such shortcomings, we may use more robust estimates of the derivatives. Similarly, we can base the computations of the variability not on the extreme values of the derivatives but employ some statistical measures, say quartiles of the differences. Any combination of these measures can easily remedy the potential drawbacks of noisy data.

D. DCT

A discrete cosine transform (DCT) is finite sequence of data points in terms of a sum of cosine functions oscillating at different frequencies. DCTs are important to numerous applications in science and engineering, from lossy compression .Where small high-frequency components can be discarded, to spectral methods for the numerical solution of partial differential equations. The use of cosine rather than sine functions is critical for compression, since it turns out that fewer cosine functions are needed to approximate a typical signal.



Fig.3: ECG signal after DCT compression

DCT-II technique, which is very advance. When there is high correlation among the input samples, which is the case in many digital waveforms including speech, music, and biomedical signals. This transform is exactly equivalent to a DFT of 4n real inputs of even symmetry where the even-indexed elements are zero.



E. DST

Fig. 4:ECG signal after DCT-II compression

The discrete sine transform (DST) is a Fourier-related transform similar to the discrete Fourier transform (DFT), but using a purely real matrix. It is equivalent to the imaginary parts of a DFT of roughly twice the length, operating on real data with odd symmetry (since the Fourier transform of a real and odd function is imaginary and odd), where in some variants the input and/or output data are shifted by half a sample. A related transform is the discrete cosine transform (DCT), which is equivalent to a DFT of real and even functions. See the DCT article for a general discussion of how the boundary conditions relate the various DCT and DST types. The DST-I matrix is orthogonal (up to a scale factor). A DST-I is exactly equivalent to a DFT of a real sequence that is odd around the zero-th and middle points, scaled by 1/2. For example, a DST-I of N=3 real numbers (a,b,c) is exactly equivalent to a DFT of real numbers (0,a,b,c,0,-c,-b,-a) (odd symmetry), scaled by 1/2. (In contrast, DST types II-IV involve a half-sample shift in the equivalent DFT.) This is the reason for the N+1 in the denominator of the sine function: the equivalent DFT has 2(N+1) points and has $2\pi/2(N+1)$ in its sinusoid frequency, so the DST-I has $\pi/(N+1)$ in its frequency.



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Fig.5: ECG signal after DST compression

TABLE- I

COMPARISON OF DCT	& DST COMPR	ESSION TECHNIOU	ES
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Method	Compression Ratio	PRD
DCT	91.6800	0.8392
DST	70.4073	1.1967
DCT-II	94.28	1.5729

Considering that the number of electrocardiogram records annually numbers in the millions and the use of sending electrocardiogram records over telephone lines for remote analysis is increasing, the need for effective electrocardiogram compression techniques is great. Many existing compression algorithms have shown some success in electrocardiogram compression; however, algorithms that produce better compression ratios and less loss of data

IV. CONCLUSION

We have studied maximum approaches for Biomedical signal (ECG) compression in secure communication network, and we found the idea is based on 2D transform (complete or over complete) to enhance the scarcity of the coefficients. By using Kalman smoother filtering about 2dB for higher input SNRs, but it does not give outstanding results for ECG signal compression. MSVQ has been introduced for ECG signal compression, with good results in comparison with other compression methods. It does not employ any QRS detection and the bit rate of the compressed signal does not depend on sampling conditions. Finally, through the use of more sophisticated VQ construction schemes and the use of entropy coding techniques, further improvements are to be expected, both in PRD-CR performance as well as in memory requirements. It is worth stressing that the genetic segmentation of the signal does not confine itself to a specific form of compression such as a piecewise linear approximation or any other technique by that matter. DST approach provides lowest CR and distortion is also high and technique of DCT improves CR with lowers PRD.

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