An improved method for R-peak detection by using Shannon energy envelope

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Abstract. R-peaks in electrocardiogram (ECG) play a vital role in diagnosis of heart rhythm irregularities and also estimating heart rate variability. However, almost all existing R-peak detectors suffer from the non-stationary of both QRS morphology and noise. To overcome these difficulties, we propose a four-stage improved method to detect R-peak using Shannon energy envelope. In the first stage, noise is suppressed and QRS complex is enhanced by using band pass filter, first order differentiation, and amplitude normalization. In the second stage, Shannon energy envelope is extracted. In the third stage, peak is estimated without considering any threshold amplitude. In the final stage, true R-peaks are detected. Our proposed R-peak detection method is validated using 48 first channel ECG records of the MIT-BIH arrhythmia database with the accuracy of 99.84%, sensitivity of 99.95% and positive predictability of 99.88%. Our proposed method outperforms other well-known methods in case of pathological ECG signals.

Keywords. Electrocardiogram; MIT-BIH database; QRS complex; Shannon energy envelope.

1. Introduction

Electrocardiogram (ECG) is an important biological signal that reflects the electrical activity of heart. It provides a lot of information on the normal and pathological physiology of heart. An ECG signal consists of different parts such as P wave, QRS complex and T wave [1]. As it is a non-stationary signal, it is very difficult to analyse it visually and therefore a computer based ECG analyzer is required. R-peak detection in ECG is a popular method in order to diagnose heart rhythm irregularities and also to estimate heart rate variability (HRV) [2–4].

Various signal processing algorithms have been developed for detecting R-peak in the ECG. Different methods for detecting R-peaks in ECG are Pan–Tompkins method [5], artificial neural network (ANN) [6], heuristic methods [7, 8], wavelet transform (WT) [9–14], Hilbert transform [15], empirical mode of decomposition [16], Shannon energy with Hilbert transform method [17], and Shannon energy envelope [18].

Pan–Tompkins method (PT) [5] based on moving average filter and peak detection is a low complexity algorithm but yields a lower accuracy. Moreover as the peak detection is based on amplitude threshold that degrades accuracy. ANN [6] requires complex rules for detecting R-peak. The

performance of heuristic methods [7, 8] depends on suitable choice of bandwidth of band pass filter (BPF) and also the duration of moving window used for integration. Wavelet based methods [9–11, 13, 14] require selection of suitable mother wavelet and scales to obtain QRS event. Hilbert transform and empirical mode of decomposition give high accuracy but on a limited number of records in MIT database. Shannon energy with Hilbert transform method (SEHT) [17] provides a good accuracy for detecting R-peak. Hilbert transform in SEHT requires high memory and delays making it unsuitable for real time implementation. For a long pause ECG data, SEHT detects a large number of noise peaks. Computation of Hilbert transform in frequency domain requires very long memory buffer for storing samples of ECG data. In 2013, Zhu and Dong [18] have developed an R-peak detection method called PSEE by using only Shannon energy envelope. Again for valid pair peak detection there is a use of amplitude threshold that affects the performance of the algorithm.

In this paper, we have proposed an algorithm based on the Shannon energy envelope which does not require any amplitude threshold for detecting R-peak. It has also overcome the drawbacks of the R-peaks detection techniques based on Shannon energy envelope [18] with a better accuracy. Our proposed method shows an accuracy of 99.84% evaluated with all the records of MIT-BIH arrhythmia database [19, 20].

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In section 2, the four-stage R-peak detection algorithm is described in detail. We introduce the R-peak detection methodology based on the Shannon energy envelope without requiring any amplitude threshold. The sensitivity, positive predictability and accuracy of our proposed algorithm evaluated from MIT-BIH arrhythmia database are explained in section 3. Finally, we discuss and conclude our study in section 4.

2. The proposed R-peak detection methodology

The block diagram of proposed R-peak detection methodology is shown in figure 1. It consists of four stages namely pre-processing, Shannon energy envelope extraction, peak estimation logic and true R-peak detection. In the pre-processing stage noise is suppressed and QRS complex is enhanced by using a BPF, first order differentiation and followed by amplitude normalization. In the Shannon energy envelope extraction stage, smooth Shannon energy envelope is produced by using Shannon energy calculation and a moving average filter. In peak estimation logic stage, the pre-processing of R-peak detection is performed by using first order differentiation, amplitude normalization, squaring, followed by a moving average filter for smoothing. In true R-peak detection stage, the actual location of

R-peak is detected. The detailed discussion of each stage in figure 1 is presented in the following sections.

2.1 Data pre-processing

In reality, the pathological ECG signals are corrupted by various noises such as power line interference noise (60 Hz), motion artefact due to electrode, skin interface, patient breathing, etc. Moreover, ECG signal contains P wave, T wave other than QRS complex which are not of interest in our work. Therefore, we need a filtering system in order to extract QRS complex. From the power spectral analysis of various signal components in ECG signal, we observe that the maximum power density of QRS complex lies between 5 and 20 Hz [21]. In our proposed method, the ECG is passed through a Chebyshev type I filter with bandwidth of [6 18] Hz for this purpose.

After bandpass filtering the signal, we perform the first order differentiation of the signal to get the slope information. The first order differentiation is nothing but a high pass filter which passes high frequency components (QRS complex) and attenuate lower frequency components (P and T wave). The mathematical implementation of the first order differentiation is shown below:

$$d[n] = f[n + 1] - f[n].$$

Figure 1. Block diagram of four-stage R-peak detection methodology.

After filtering and differentiating, the signal is normalized in order to scale its value to 1. This is done for preparing the signal for Shannon energy envelope computation.

$$\widehat{d[n]} = \frac{d[n]}{\max(|d[n]|)}.$$

2.2 Shannon energy envelope extraction

After differentiating the ECG signal it becomes a bipolar signal. Since the method is based on peak detection, we have to transform the differentiated signal into a unipolar signal. The unipolar signal can be obtained by absolute value method, squared value method, Shannon entropy method, and Shannon energy envelope method. The mathematical formulation of these methods is described below.

Absolute value: $y[n] = |a[n]|$

Squared value: $y[n] = a[n] \times a[n]$

Shannon entropy value: $y[n] = -|a[n]| \times \ln(|a[n]|)$

Shannon energy value: $y[n] = -(a[n])^2 \ln((a[n])^2)$

where $a[n]$, $y[n]$ are input and output for the above equations respectively.

In the proposed method, we have used Shannon energy envelope method as it has some advantages over the others. Figure 2 shows that absolute value method produces same weight value irrespective of input amplitude. Shannon entropy method attenuates high intensity components and gives high preference to low intensity components. The squared value method gives an exponential weightage response to the high intensity components which will make detection of low intensity components difficult. For Shannon energy method, middle intensity components are more emphasized and other intensity values are attenuated which gives a better detection of R-peaks.

For the sake of getting a unipolar signal, Shannon energy envelope calculation has been employed. When we are taking the logarithm of the normalized derivatives by using the Shannon energy envelope, the maximum amplitude of the normalized derivative is diminished, but not the R-peaks. The absolute value of the amplitude of the normalized derivative corresponds to the R-peaks location lies between 0.4 and 0.8 as shown in figure 3. Shannon energy gives more emphasis when the absolute value of the amplitude lies between 0.4 and 0.8 as shown in figure 2. The absolute value of the amplitude of normalized derivatives between 0.4 and 0.8 refers to the R-peaks. So to give more emphasize on R-peaks we have used Shannon energy envelope calculation.

Figure 2. Comparison of energy value for different envelope methods using normalized amplitude.

Figure 3. Steps for preparing the signal for Shannon energy envelope calculation.

After extracting Shannon energy envelope, the signal is passed through a zero shift moving average filter in order to get a smooth Shannon energy envelope [17]. The each step of the second stage is shown in figure 4. It shows that after Shannon energy we do not get smooth envelope as shown in figure 4(b), due to its false peaks may be considered as R-peaks. Figure 4(c) shows that a smooth envelope is an output of moving average filter.

A moving average filter is nothing but a moving window integrator. Mathematical expression of a moving average filter is expressed below.

$$y(nT) = \frac{1}{N} [a(nT - (N-1)T) + a(nT - (N-2)T) + \dots + a(nT)].$$

The length of the moving average filter is an important parameter for this detection method. Generally, the length of the moving average filter is taken approximately to the width of the QRS complex. If the length of moving average filter is too large then the integration takes the T wave also, and if the length of the moving average filter is smaller compared to the QRS complex, then for a single QRS complex, it produces more spikes which leads to an erroneous detection. In our proposed method, we have taken a length of 65 samples moving average filter.

The window length of the moving average filter for performing the average operation on the Shannon energy envelope results depend upon the sampling frequency (f_s) under which the measurements are taken. In this work the window length of the moving average filter for producing smooth envelope is $0.18 \times f_s$. The sampling frequency of

Figure 4. Extraction of smooth Shannon energy envelope.

Figure 5. Peak estimation logic without considering any amplitude threshold.

the used database (MIT-BIH arrhythmia database) is 360. So we have taken the window length of 65 ($360 \times 0.18 = 64.8 \cong 65$) samples moving average filter.

2.3 Peak estimation logic

In this stage, we introduce a different approach from the other R-peak detection technique based on Shannon energy. Manikandan *et al* [17] use Hilbert transform with Shannon energy for detection of R-peaks but the Hilbert transform requires more memory buffer. Zhu and Dong [18] use amplitude threshold in the decision stage for extraction for true R-peaks. To overcome these problems, our R-peak detection logic should be free from amplitude threshold. After Shannon energy envelope extraction stage, we get smooth peak signal but it contains both true R-peaks and false R-peaks. In this stage, we actually attenuate the false R-peaks and emphasize the true R-peaks. This is done by following steps.

It is natural that the amplitude value for true R-peaks will be quite high than the false peaks. So if we take the first order differentiation of the signal, then it stores the slope information of the true peaks but it reduces the slope information of the false peaks.

First the signal is amplitude normalized to unity. The normalized bipolar signal is converted to unipolar signal by squaring operation. The amplitude of false peaks is very low and the squaring operation will attenuate these peaks completely. As a result, the true R-peaks are amplified and false R-peaks are diminished.

Figure 6. Different signal processing steps of peak estimation logic stage.

Figure 7. Total signal processing steps of the proposed R-peak detector: (a) ECG signal (Record 205). (b) Filtered signal. (c) Output of normalize operation. (d) Smooth Shannon energy envelope. (e) Differentiation of Shannon energy envelope. (f) Output of square operation. (g) Output of moving average filter. (h) True detected R-peaks (red colour impulses).

Then, the signal is passed through a moving average filter in order to get a smooth peak signal. In our proposed technique, we use a moving average filter of length 85 samples. Now signal is ready for peak detection.

Figure 5 shows labelled diagram of peak estimation logic without considering any amplitude threshold. After squaring, the smooth signal cannot be achieved, to get smooth signal moving average filter operation is performed. The output of the moving average filter does not provide any unnecessary rising peaks. So no threshold operation is required. The locations of rising peaks are referred to the locations of true R-peaks as shown in last subplot of figure 5. Thus, no amplitude threshold value is required for detection of R-peaks.

This peak estimation logic stage is demonstrated in figure 6.

2.4 True R-peak detection

In this stage, the R-peaks are detected by applying peak finding algorithm. However, the detected peak locations are slightly different from the actual positions of the R-peaks in

Table 1. Performance of our proposed R-peak detection method using MIT-BIH arrhythmia database with length 30 min.

Data no.	TP	FN	FP	Se (%)	+P (%)	DER (%)	Acc (%)
100	2272	1	0	99.95601	100	0.044014	99.95601
101	1866	0	2	100	99.89293	0.107181	99.89293
102	2187	0	0	100	100	0	100
103	2084	0	0	100	100	0	100
104	2255	0	4	100	99.82293	0.177384	99.82293
105	2581	5	5	99.80665	99.80665	0.387447	99.61405
106	2000	3	5	99.85022	99.75062	0.4	99.60159
107	2136	1	0	99.95321	100	0.046816	99.95321
108	1762	5	8	99.71703	99.54802	0.737798	99.26761
109	2532	0	3	100	99.88166	0.118483	99.88166
111	2124	0	4	100	99.81203	0.188324	99.81203
112	2539	0	1	100	99.96063	0.039386	99.96063
113	1795	0	0	100	100	0	100
114	1879	0	4	100	99.78757	0.212879	99.78757
115	1954	0	6	100	99.69388	0.307062	99.69388
116	2405	5	1	99.79253	99.95844	0.24948	99.75114
117	1535	0	1	100	99.9349	0.065147	99.9349
118	2276	2	5	99.9122	99.7808	0.307557	99.69339
119	1985	2	4	99.89935	99.79889	0.302267	99.69864
121	1866	0	4	100	99.7861	0.214362	99.7861
122	2476	0	0	100	100	0	100
123	1520	0	5	100	99.67213	0.328947	99.67213
124	1628	0	8	100	99.511	0.4914	99.511
200	2610	0	4	100	99.84698	0.153257	99.84698
201	1962	1	2	99.94906	99.89817	0.152905	99.84733
202	2136	0	2	100	99.90645	0.093633	99.90645
203	2981	9	8	99.699	99.73235	0.570278	99.43296
205	2655	1	0	99.96235	100	0.037665	99.96235
207	1857	3	5	99.83871	99.73147	0.430802	99.57105
208	2948	7	3	99.76311	99.89834	0.339213	99.66193
209	3003	2	0	99.93344	100	0.0666	99.93344
210	2633	1	2	99.96203	99.9241	0.113938	99.88619
212	2748	0	0	100	100	0	100
213	3250	1	2	99.96924	99.9385	0.092308	99.90778
214	2264	0	2	100	99.91174	0.088339	99.91174
215	3362	1	0	99.97026	100	0.029744	99.97026
217	2208	0	1	100	99.95473	0.04529	99.95473
219	2154	0	0	100	100	0	100
220	2048	0	0	100	100	0	100
221	2427	0	1	100	99.95881	0.041203	99.95881
222	2482	0	1	100	99.95973	0.04029	99.95973
223	2606	0	0	100	100	0	100
228	2053	0	5	100	99.75705	0.243546	99.75705
230	2256	0	2	100	99.91143	0.088652	99.91143
231	1571	0	0	100	100	0	100
232	1780	0	4	100	99.77578	0.224719	99.77578
233	3070	8	2	99.74009	99.9349	0.325733	99.67532
234	2753	0	0	100	100	0	100
Total	109,474	58	116	99.95155	99.88416	0.164668	99.84131

ECG signal. So in order to find the real positions of R-peaks, the actual sample instant of R-peaks in the ECG signal is found by searching the maximum amplitude within ± 25 samples of the identified location in the previous step.

In figure 7, the total signal processing of the proposed method is shown, where in figure 7(h) red impulses indicate the detected R-peaks from the proposed method.

Table 2. Comparison of performance of our proposed method with other methods for detecting R-peaks using MIT BIH database.

Method	Total beats (TP+FN)	FN (beats)	FP (beats)	Se (%)	+P (%)	DER (%)	Acc (%)	Standard deviation of Acc (%)
Our proposed method	109,532	58	116	99.95155	99.88416	0.164668	99.84131	17.1
PSEE [18]	109,494	93	91	99.92	99.92	0.168	99.832	28.7
WT [12]	104,184	65	112	99.94	99.89	0.170	99.830	31.6
SEHT [17]	109,496	79	140	99.93	99.87	0.200	99.800	29.7
DOM [4]	109,809	58	166	99.95	99.85	0.204	99.796	25
WT [13]	109,428	220	153	99.80	99.86	0.340	99.660	–
WT [10]	110,159	322	120	99.89	99.70	0.402	99.599	56.5
PT [5]	109,809	507	277	99.54	99.75	0.712	99.288	20

Figure 8. Performance of the proposed method for Record 103 (Low amplitude QRS complexes and baseline drift).

3. Results and discussion

Performance of the proposed R-peak detection method is evaluated by using MIT-BIH arrhythmia database [19, 20]. The MIT-BIH Arrhythmia database contains 48 half-hour two-channel ECG recordings, obtained from 47 subjects studied by the BIH Arrhythmia Laboratory between 1975 and 1979. The sampling frequency of the digitized ECG signal is 360 samples per second per channel with 11-bit resolution over a 10 mV range.

The proposed method is implemented on MATLAB R 2012A.

To evaluate the performance of our proposed R-peak detection method, we require three parameters namely true positive (TP), false negative (FN) and false positive (FP)

Figure 9. Performance of the proposed method for Record 232 (long pauses).

Figure 10. Performance of the proposed method for Record 106 (continuously varying QRS complexes and tall T waves).

Figure 11. Performance of the proposed method for Record 105 (noisy ECG signal).

from detected R-peak. Here TP is the number of correctly detected R-peaks, FN is the number of missed R-peaks. FP is the number of noise spikes detected as R-peaks. Sensitivity (Se), positive predictability (+P), detection error rate (DER) and accuracy (ACC) can be computed by using TP, FN, and FP by using the following equations respectively.

$$Se = \frac{TP}{TP + FN} \times 100\%$$

$$+P = \frac{TP}{TP + FP} \times 100\%$$

$$DER = \frac{FP + FN}{TP} \times 100\%$$

$$Acc = \frac{TP}{TP + FP + FN} \times 100\%$$

The performance of the proposed R-peak detection method for 48 ECG recording of the MIT-BIH arrhythmia database is summarized in table 1. The

Figure 12. Performance of the proposed method for Record 108 (small R-peaks).

proposed method detects a total number of 109,474 true peaks. It also produces 58 false negative (FN) and 116 false positive (FP). The average accuracy for the proposed method is 99.841% while the individual detection accuracies vary from 99.26% to 100%.

In table 2, the performance of the proposed method on MIT-BIH arrhythmia database is compared with other existing methods. It shows that our proposed method has a better accuracy than the other methods like wavelet transform techniques, differential operation method, Pan-Tompkins algorithm, Shannon energy with Hilbert transform technique, Shannon energy technique and our proposed method has less standard deviation compared to other methods.

In MIT-BIH arrhythmia database, records 104, 108, 203, and 228 contain high-grade of noise. Therefore, the accuracy of the method for these records is comparatively weaker than other records. Record 103 contains low amplitude QRS complexes and baseline drift, the performance of the proposed method for this record is shown in figure 8.

Record 232 contains a number of long pauses in the ECG signal. Our proposed method has better response to this signal shown in figure 9.

Record 106 has tall T waves and it contains continuously varying QRS complexes. Detection performance of the expressed method for this signal is shown in figure 10.

Record 105 has noisy ECG signal. Detection performance of the expressed method for this signal is shown in figure 11.

Record 108 has small R-peaks ECG signal. Detection performance of the expressed method for this signal is shown in figure 12.

4. Conclusion

In this paper, a computer based automated R-peaks in an ECG signal detection method have been described. This method is a four-stage detection method which is based on the Shannon energy envelope. The detection process of the method is based on band pass filter, first order differentiation, amplitude normalization, Shannon energy envelope, zero shift moving average filter, and squaring. The decision stage is based on first order differentiation, amplitude normalization, squaring and moving average filtering. In this method, the peak finding logic has been improved in such a way that there is no such use of amplitude threshold and no need of prior knowledge of the past detected R-peaks.

In order to evaluate the performance and effectiveness of the proposed R-peak detection method, we have used standard MIT-BIH arrhythmia database which consists of 48 ECG records of 30 min each. The detection performance of the proposed methodology is evaluated in terms of sensitivity (Se %), positive predictability (+P %), detection error rate (DER %), and accuracy (Acc %) which are calculated from true peak (TP), false negative (FN), and false positive (FP). Our proposed method achieves an average accuracy of 99.84%, an average sensitivity of 99.95% and positive predictability of 99.88% by using MIT-BIH arrhythmia database. For being a simple mathematical operation based algorithm, the proposed method has a very less computation time which is suitable for real time application. Moreover for ECG signals, containing noise and artefacts like varying QRS morphology, low amplitude, long pauses our proposed method gives a better response than the existing methods.

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