ANALYSIS OF SEVERAL CHARACTERISTICS OF ECG SIGNAL FOR CARDIAC ARRHYTHMIA DETECTION

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ABSTRACT

A vast number of people in the world today suffer from various cardiac problems. As a result, studying the characteristics of the ECG signal is critical for diagnosing various heart disorders. The electrocardiogram (ECG) is a test that shows how strong the electric impulses in the heart are. PQRST waves are various waves that make up one cardiac cycle of an ECG signal. The amplitude and time intervals of PQRST waves are estimated for the learning of ECG signals in the attribute removal of ECG signal. The amplitudes and time intervals values of the PQRST segment can be used to determine the appropriate operation of the human heart. The majority of approaches and research for evaluating the ECG signal have been created in recent years. The Wavelet Transform, Support Vector Machines (SVM), Genetic Algorithm (GA), Artificial Neural Networks (ANN), Fuzzy Logic Methods, and other Signal Examination techniques are used in the majority of the systems. In this paper comparison of SVM, ANN, Neural Mode Decomposition (NMD) and Incremental Support Vector Regression (ISVR) methods are presents. From these methods, theISVR method shows better results. However, each of the algorithms and strategies listed above has its own set of benefits and drawbacks. The wavelet transform Db4 is utilized to extract different features from an ECG signal in this article. Matlab software is utilized to design the proposed system. The proposed algorithm is demonstrated in this work using the MIT-BIH Arrhythmia record, which is used to manually annotate and develop validation.

Keywords: *Electrocardiogram (ECG), Wavelet Transform Db 4, QRS Complex, median filter, Cardiac Arrhythmia*

1. Introduction

One of the most important organs in the human body is the heart. It uses blood to deliver oxygen to the patient's body. The heart functions like a muscle pump. The heart is connected to the rest of the body via a complicated network of arteries, veins, and capillaries. The Electrocardiogram (ECG) is a compilation of various biopotential signals from human heartbeats. The electrodes are placed on the patient's epidermis to capture these bio potential signals. It's a visual representation of the electrical activity of the heart's muscles. ECG aids in the transmission of information about the heart and cardiovascular system. It's an important and fundamental mechanism for treating heart problems. Any abnormal heart rate in the morphological sample is an indication of cardiac arrhythmia. The ECG assists in providing information about the heart and circulatory system.It is a helpful and important tool for determining the severity of cardiac disease. The electrical activity of the cardiac muscles is represented by the ECG waveform, which is made up of unique electrical depolarization–repolarization patterns.The ECG signal is a real-time recording of the direction and magnitude of the electric commotion caused by depolarization and repolarization of the atria and ventricles of the coronary heart.

An arrhythmia is defined as a problem with coronary cardiac charge or rhythm, or a change in the morphological pattern. The selecting process takes longer with the guide statement for evaluating the recorded ECG waveform. As a result, a detection and classification device based on Artificial Intelligence (A.I.) is used. The P-QRS-T waves make up one cardiac cycle in an ECG signal. An example ECG signal is shown in Figure 1. The ECG is mostly used to follow and analyses patients.

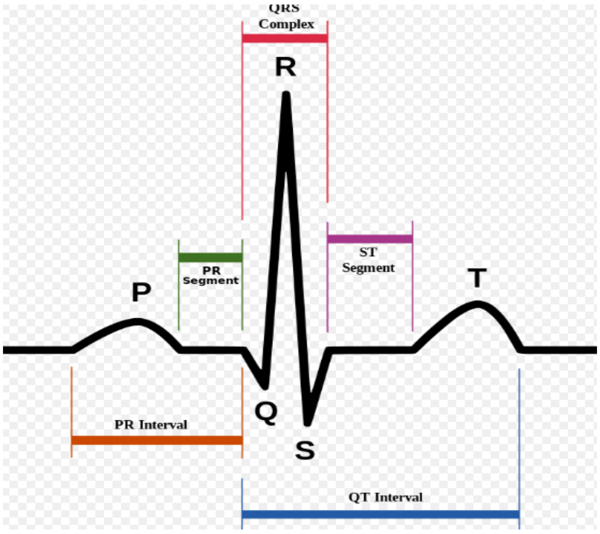

Figure 1. Shows a sample ECG signal with the P-QRS-T wave[24].

The characteristic retrieved from the ECG sign is critical in diagnosing heart illness. The development of accurate and time-saving technologies for automatic ECG feature extraction is critical. As a result, it's critical that the function extraction system works correctly. The goal of feature extraction is to find as few properties in an ECG signal as feasible that will allow for successful anomaly identification and green prognosis. This paper provides an overview of the technique and transformations used to extract various aspects from an ECG signal. The amplitude and duration of waves, intervals, and segments of the ECG signal are shown in table 1.

Table 1. Amplitude and period of waves, intervals and segments of ECG signal.

Sr no.	Features	Amplitude (mV)	Duration (ms)
1	P wave	0.1-0.2	60-80
2	PR-segment	–	50-120
3	PR-interval	–	120-200
4	QRS complex	1	80-120
5	ST-segment	-	100-120
6	T –wave	0.1-0.3	120-160
7	ST-interval	-	320
8	RR-interval	-	(0.4-1.2)s

The total performance of the ECG pattern category is greatly influenced by the characterisation energy of the capabilities generated from ECG data and the classifier's layout. The wavelet transform is a powerful technique for reading no stationary ECG indications because of its time–frequency localization features. The wavelet transform can be used to deconstruct an ECG sign in scale, allowing the key ECG waveform morphological descriptors to be separated from the original signal's noise, interference, baseline flow, and amplitude fluctuation.
The following is the structure of this paper: The proposed system's literature review is presented in section II. Section III examines the system's growth, as well as the several potential methods presented in this study. The simulation of the suggested system as well as the experimental findings are presented in section IV. MATLAB is used to run the simulation. Finally, part V brings this paper to a close.

2. Review of the Existing Literature

J.I. Williamset al.[1] performed out the measurement, which was examined separately by a group of cardiologists from the American Heart Association. The results of an analysis of a series of proposals aimed at standardizing quantitative ECG measurement are presented. These AHA suggestions have gained worldwide acclaim. Zhao et al [2] proposed a novel feature extraction method for accurate heart rhythm identification. The suggested classification method is made up of three parts: data pre-processing, feature extraction, and ECG signal categorization. The feature vector of ECG data is created by combining two different feature extraction algorithms. The coefficients of the transform are extracted as features of each ECG segment using the wavelet transform. At the same time, autoregressive modelling (AR) is used to grasp the temporal features of ECG signals. Finally, distinct ECG heart rhythms are classified using a support vector machine (SVM) with a Gaussian kernel. The correctness of the proposed approach was determined by computer simulations, which yielded a 99.68 percent overall accuracy.

The filtering approach based on moving averages reported by V. S. Chouhanet.al [3] produces smooth spike-free ECG output, which is suitable for slope feature extraction. The first step is to extract the slope feature from the filtered and drift-corrected ECG signal by processing and converting it so that the derived feature signal is greatly boosted in the QRS area and suppressed in the non-QRS region. The proposed method has a detection rate of 98.56 percent and a positive predictive value of 99.18 percent, respectively. S.C. Saxena et al. developed a modified mixed wavelet transforms technique in [4].

The approach was created to evaluate multi-lead ECG readings in order to diagnose heart problems. For QRS detection, a quadratic spline wavelet (QSWT) was utilised, while for P and T detection, the Daubechies six coefficient (DU6) wavelet was used. For the identification of various heart disorders, a process based on ECG data and a point scoring system has been developed. When both diagnostic criteria yielded the same results, the consistency and reliability of the discovered and measured parameters were confirmed. The comparison of several ECG signal feature extraction approaches is shown in Table 1. Ramli and Ahmad [5] explained correlation analysis for aberrant ECG signal feature extraction.

Their planned research looked on a technique for extracting essential features from ECG signals from a 12-lead system. They picked II as the basis for their entire study since it has representative properties for detecting prevalent cardiac problems. Cross-correlation analysis was used as the analysis technique. Cross-correlation analysis determines how similar two signals are and extracts the information contained in them. Their tests showed that the proposed technique could efficiently identify elements that differentiate between the numerous types of heart disorders studied as well as a normal heart signal.

Laurence et al. [6] provide a continuous wavelet transform-based technique for studying the respiratory sinus arrhythmia's non-stationary intensity and phase delay (RSA). The RSA is the cyclic variation of instantaneous heart rate at the frequency of breathing. Paced breathing or postural alterations, low respiratory frequencies, and quick changes have all been observed in studies of cardio-respiratory interaction during sleep. BekirKarhket et al. [7] used an artificial Neural network to assess ECG signals in the time domain and determine related arrhythmias, achieving a 95 percent success for arrhythmia recognition.

Chuang-chienetet al.[8] developed an efficient arrhythmia recognition algorithm based on the correlation coefficient in ECG signals for QRS complex detection. The correlation coefficient and RR interval were used to quantify arrhythmia similarity. Stefan Gradl et al. [9] investigated the Pan-Tompkins method for QRS detection, template generation and adaption, feature extraction, and beat classification. The MIT-BIH Arrhythmia and MIT-BIH Supraventricular Arrhythmia databases were used to validate the algorithm. The programme properly detected more than 98 percent of all QRS complexes. The overall sensitivity and specificity for aberrant beat detection were 89.5 percent and 80.6 percent, respectively.Wavelet transform and linear discriminate analysis were used to extract input features by J. Lee, K. et al.[10]. The accuracy of the proposed approach in detecting arrhythmias was 97.52, 96.43, 98.59, and 97.88 percent for NSR, SVR, PVC, and VF, respectively.

The wavelet transform and hidden Markova models were performed by Pedro R.Gomes et al. [11]. The experimental results in real data from the MIT-BIH arrhythmia data source show that it outperforms the standard linear segmentation. VahidTavakoliet.al [12] describe a novel ECG signal analysis approach based on non-uniform sampling. It is demonstrated that a newly developed method called Finite Rate of Innovation (FRI) may be used to better assess Left and Right Bundle Branch Block arrhythmias as well as normal signals, and that spline modelling can be used to better study different types of arrhythmia. As a result, a multi-stage technique for diagnosing and compressing ECG signals is provided, which is both faster and more accurate. The wireless and wearable sensor ECG system, hand held device with RF receiver, and arrhythmia algorithm were studied by Rune Fensli, et al.[13].

The continuous wavelet transform (CWT) was investigated by Khaledet.al. [14] for evaluating ECG signals and extracting desirable parameters such as arrhythmia. This approach distinguishes between Nomo cardia, Bradycardia, and Tachycardia with a clear threshold. The Pan Tomkins algorithm (it is implemented for the identification QRS complex on normal and arrhythmia databases and discrete WT) has been researched by V.Vijaya et al.[15]. The most common cause of mortality is cardiac arrhythmia. An algorithm for R Peak and QRS complex detection using WT has been developed and assessed using ECG feature extraction. A.R.Saheb et al.[16] investigated the construction of a heart diagnosis tool with few complex computations. The accuracy of the designed classifier is 98 percent, and it has been attained for three different arrhythmias, including RBBB, LBBB, and normal heart rhythm. The sequential probability ratio test has been modified by Szi-Wen Chen et al.[17]. Using this method, they were able to reduce the total rate error rate by 5% compared to the previous result.The feature extraction from the Pan-Tompkins algorithm and the Hierarchical system was studied by Heike Leutheuseret.al. [18]. Early detection of arrhythmic beats in the ECG signal may help identify patients who are at risk of sudden death due to coronary heart disease.

We chose the last expectation for organization because the algorithm predicted a characterization at each eighteenth information test.

The Physionet Challenge 2017 preparation dataset, which contains 8,528 single lead ECG accounts lasting from 9s to little over 60s, was used to cross-approve the results.
[24]

3. Resources and Techniques Employed

This section depicts the resources and strategies used to put the planned system into action.This section describes how the proposed method evolved.This approach has been evaluated on a training data set, and several modules are examined in this part. The block diagram of the suggested system is shown in Figure 2.

3.1 Input Image

Approach that has been proposed, the MIT-BIH Arrhythmia Database was used to test this approach.The suggested ECG classifier is trained and evaluated using the MIT/BIH arrhythmia database in this research. The database contains 48 facts, each of which has two-channel ECG signs for 30 minutes, chosen from 24-hour recordings of 47 people. Continuous ECG data are band pass filtered between one and one hundred Hz before being digitized at 360 Hz. The database includes annotation for both timing records and beat elegance data that has been verified by independent professionals.

The first 20 statistics (numbered between 100 and 124), which include representative samples of common medical recordings, are utilized to choose representative beats to be included in the common education data. Ventricular, junctional, and supraventricular arrhythmias are represented in the remaining 24 records (numbering between 200 and 234). Each ECG beat should be classified into one of the three heartbeat types: N (beats originating in the sinus mode), V (ventricular ectopic beats (VEBs/PVC), or Other (unclassified beats).

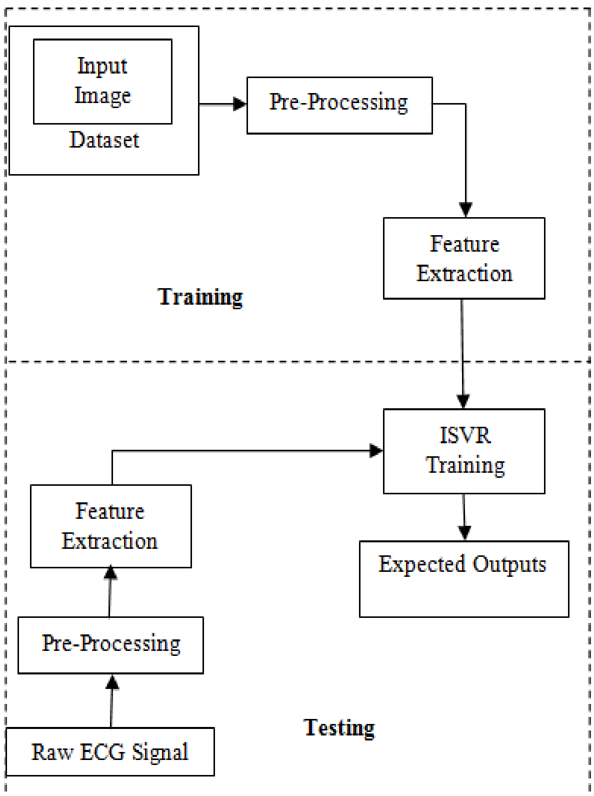

Figure 2. Block Diagram of Suggested System.

3.2 Preprocessing

Muscle noise is a significant issue in many ECG applications, particularly in recordings taken during exercise, because low amplitude waveforms can become entirely covered. Muscle noise is not removed by narrowband filtering, unlike baseline wander and 50/60 Hz interference, and presents a far more challenging filtering problem because the spectral content of muscle activity significantly overlaps that of the PQRST complex. Because the ECG is a repeated signal, procedures similar to those used to analyse evoked potentials can be utilised to decrease muscle noise. However, noise reduction by ensemble averaging is limited to one QRS shape at a time and needs the availability of multiple beats.

Frequency interference, baseline waft, electrode touch noise, polarisation noise, muscle noise, internal amplifier noise, and motor artefacts are all examples of sounds found in the ECG sign. Artifacts are noise introduced into ECG data as a result of electrode movement. The reduction of baseline wander is a common problem in ECG signal processing. To limit the variations in beat shape, baseline wander must be removed from the examination of the ECG sign.

In most types of ECG recordings, breathing and electrode impedance changes as a result of respiration are essential assets of baseline wander. The frequency content material of the baseline wander is usually in the sub-0.5Hz range. This baseline glide can be deleted without affecting or requiring the waveform's characteristics. To remove the baseline flow of the ECG signal, we apply median filters (2 hundred and 600 milliseconds) [21]. The steps are as follows:

A) To remove QRS complexes and P waves, the original ECG signal is treated using a mean clear out of 200 ms width.

B) To eliminate T waves, the resulting signal is next treated with a median filter with a width of 600 ms. The baseline of the ECG sign is carried by the sign as a result of the second filter operation.

C) A sign with baseline float removal can be obtained by subtracting the filtered sign from the authentic sign.

3.3 Discrete Wavelet Transform

The DWT's basic concept for a one-dimensional symbol is as follows:

A signal is separated into two halves, typically high and low frequencies. In the excessive frequency component, the signal's aspect additives are largely limited. The low frequency component is broken down further into two parts: excessive and coffee frequency. This process is repeated until the sign has completely disintegrated or has been halted using the application to hand approach. Normally, no more than five decomposition steps are computed for compression and watermarking software. The unique signal can also be reconstructed using the DWT coefficients. Inverse DWT is the name of the reconstruction method (IDWT). Before commencing the detecting procedure, selecting an appropriate wavelet is critical.

The wavelet to be used is determined by the type of sign to be evaluated. The wavelet that most closely resembles the signal shape is chosen. Harr, Daubechies, Biorthogonal, Coiflets, Symlets, Morlet, Mexican Hat, Meyer, and more wavelet families exist. As well as a variety of Real and Complex wavelets. We tested Daubechies 4 (Db4), Db5, Db6, rbio6.8, bior 5.5, and other variants. The Daubechies (Db6) Wavelet, on the other hand, has been proven to provide more accurate details. Furthermore, this Wavelet resembles QRS complexes, and its intensity spectrum is centred on low frequencies. As a result, for extracting the relevant features in our application, we used the Daubechies 6 (Db6) Wavelet. Figure 3 depicts the Db 4 Wavelet. To take part in all of these activities, the created programme first decomposes the purchased ECG signal into relevant Approximation and element coefficients throughout eight phases.ECGDe-noisingDespite the removal of low-frequency components from the standard sign, it may still have noise owing to high-frequency accessories.

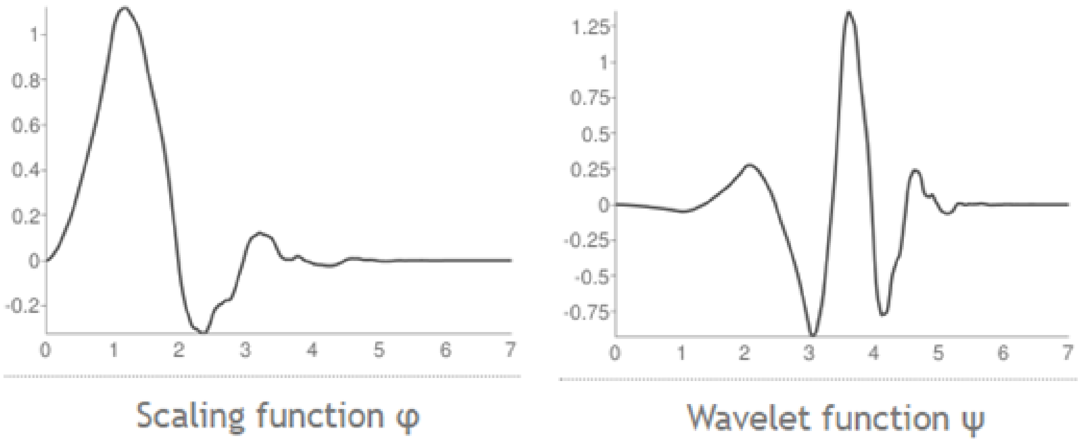

Figure 3 Coefficients of wavelet db4.

To remove noise from a signal, first determine which accessories are responsible for the noise, and then remove these recognized add-ons from the de-trended signal.

3.4 Extraction of Characteristics

The feature extraction technique is used to extract important data from the original signal. The diagnostic data from the ECG sign are extracted using the Feature extraction level in this article. It's helpful to become lost in the peaks and fine details of the sign. The initial stage in feature extraction is to find R top. In comparison to other leads, the R height inside the sign of the Modified Lead II (MLII) lead has the largest amplitude of all the waves.

The R point of the heartbeat, which is in general the place where the heartbeat has the greatest amplitude, is determined in the QRS complex detection. A regular QRS complex shows that the electric impulse has progressed normally from the bundle of his to the Purkinje community via the right and left bundle branches, and that the right and left ventricle have depolarized normally.

The QRS complex has the most energy between 3 and 40 Hz [21]. The Fourier Transform of the wavelets' 3-dB frequencies show that the majority of the QRS complex's energy is concentrated between scales 23 and 24, with the largest scale at 25. After removing noise (e.g., baseline wander), the ECG signal is squared and decomposed up to level five using the Db 4 wavelet, distinguishing approximate and detail coefficients.

The signal is then approximated by using the inverse Discrete Wavelet transform. Then, using the database annotation, a region of -300ms to +400ms around the R wave was selected to extract the number of QRS complex wavelet transform features. The DC offset was reduced, and the amplitude variance between files was abolished.The onset and offset of the QRS complex are used to compute the QRS width. After this QRS complex has been located, the next step is to determine the onset and offset parameters for each QRS complex, as well as to identify the QRS complex's component waves. At level three, the Daubechies (db4) are found to be more suitable for R peak detection.

3.5 Incremental Support Vector Regression (ISVR)

ISVR (Incremental Support Vector Regression) seems to be an effective machine learning system that works particularly well with normalized as well as binaries data.

Furthermore, because of its quadratic complexity in terms of the number of training instances, it cannot be used to train on big datasets, especially those that are extensive and require regular retraining.

To maximize SVR's validation set accuracy while using the fewest number of training examples, we suggest a straightforward two-stage greedy selection of training data [22].

Assume that one has a training data set $\{(x_1,y_1),.....,(x_l,y_l)\} \in x_i$ x,where x_i denote the training data and y_i denote the target BPs. The ISVR is to find $y=\langle\omega,\phi(x)\rangle \mathrm{H}+b,$ where ω and $\phi(x)$ denote the vectors obtained from reproducing Kernel Hilbert space H .

Therefore, we can minimize the following risks:

$$R = \frac{1}{2}\|\omega\|^2 + c\sum_{i=1}^{l} L(y_i, x_i, f). \tag{1}$$

Where $L(y_i, x_i, f)$ denotes the $\in$- loss function given by

$$L(y_i, x_i, f) = |y - f(x)|_{\in} = \max\left(0, |y - f(x)| - \in\right) \tag{2}$$

$$\min_{\omega,b,\xi,\tilde{x}i} \frac{1}{2}\|\omega\|^2 + c\sum_{i=1}^{l}\left(\xi^2 + \tilde{\xi}^2\right) \tag{3}$$

Subject to
$$\begin{cases} f(x_i) - y_i \le \in + \xi_i, i = 1, \dots, l \\ y_i - f(x_i) \le \in + \xi_i, i = 1, \dots, l \\ \xi_i, \tilde{\xi}_i \ge 0 \end{cases}$$

This problem can be solved by the Lagrangian theory as follows:

$$\omega = \sum_{i=1}^{l}(\hat{\alpha}_i - \alpha_i)\phi(x_i), \tag{4}$$

Where $\{\hat{\alpha}, \alpha\}, i = 1, \dots, l$ denotes the Lagrangian multipliers with respect to the constraints given in Eq(4), and the solution are given by

$$\max_{\alpha_i, \hat{\alpha}_i} \sum_{i=1}^{l} y_i(\hat{a}_i - \alpha_i) - \in \sum_{i=1}^{l}(\hat{a}_i + \alpha_i) - \frac{1}{2}\sum_{i,j=1}^{l}(\hat{a}_i - \alpha_i)(\hat{a}_j - \alpha_j) \left(K(x_i, x_j) + \frac{1}{c}\zeta i, j\right) \tag{5}$$

Subject to

$$\begin{cases} \sum_{i=1}^{l}(\hat{\alpha}_i - \alpha_i) = 0 \\ 0 \le \alpha_i, 0 \le \hat{\alpha}_i, i = 1, \dots, l, \end{cases}$$

Where $K(x_i, x_j) = \langle \phi(x_i), \phi(x_j) \rangle H$ and $\zeta_{i,j}$ represents the Kronecker symbol. This problem can be resolved using some disassemble methods of the ISVR.

4. Research Outcomes

This chapter entails the performance of the developed Wavelet Transformation Extraction of Different Features of ECG Signal for Cardiac Arrhythmia Detection.

4.1 MIT-BIH Dataset

The following is a summary of the data.

The ECG signals were received from the Private Hospital with real-time recording for research purposes.

The ECG signals database that was produced is described below.

1) ECG signals were obtained from 48 patients, 19 of whom were female (ages 23 to 89) and 29 of whom were male (ages 23 to 89). (age: 32-89).

2) There were 15 different forms of ECG signals: normal sinus rhythm, pacemaker rhythm, and 15 different types of cardiac dysfunctions (for each of which at least 10 signal fragments were collected).

3) A sampling frequency of 360 [Hz] and a gain of 200 [adu / mV] were used to record all ECG signals.

4) For the study, 1000 10-second (3600 sample) ECG signal fragments (not overlapping) were chosen at random.

5) The data is in Excel format, which we transform to MATLAB format.

The MIT/BIH arrhythmia database is used for training and performance evaluation of the proposed ECG classifier.The mathematical parameters calculated for MIT-BIH data are as follows.

4.2 Accuracy

The accuracy is determined using 240 examples acquired from Hospital. We selected 16 samples from each class in those samples. There are 15 classes in all.

Positive samples total (P) for each class: 16

Negative Samples in Total or Remaining 224th class (N)

True Positive (TP): Total samples of current class that were accurately retrieved (out of 16)

True Negative (TN): Total number of correctly retrieved samples from the remaining classes (Out of 224). The accuracy can be calculated by the equation

Accuracy=[TP + TN]/ [P + N] (6)

The Accuracy gained for each class is shown in the table 2. Each row represents a class, and each column represents the accuracy achieved using a suggested or current Classifier. The presented method is more accurate than other conventional methods. [21,22,23].

Table2. Comparison fo different methods for Accuracy.

Sr.No.	ISVR	SVM	ANN	NMD
1	0.995833	0.9875	0.979167	0.979167
2	0.9875	0.983333	0.979167	0.970833
3	0.9875	0.983333	0.983333	0.983333
4	0.9875	0.9875	0.9875	0.983333
5	0.991667	0.9875	0.983333	0.979167
6	0.991667	0.991667	0.983333	0.983333
7	0.991667	0.983333	0.983333	0.979167
8	0.995833	0.995833	0.991667	0.9875
9	0.991667	0.983333	0.979167	0.970833
10	0.991667	0.983333	0.975	0.975
11	0.995833	0.995833	0.991667	0.991667
12	0.995833	0.991667	0.9875	0.979167
13	0.9875	0.9875	0.9875	0.983333
14	0.995833	0.991667	0.983333	0.975
15	0.991667	0.991667	0.9875	0.983333

4.3 Sensitivity

Table 3. Comparison fo different methods for TPR.

Sr.No.	ISVR	SVM	ANN	NMD
1	0.9375	0.875	0.8125	0.8125
2	0.875	0.8125	0.75	0.6875
3	0.9375	0.9375	0.9375	0.9375
4	0.8125	0.8125	0.8125	0.8125
5	0.9375	0.9375	0.875	0.8125
6	0.9375	0.9375	0.875	0.875
7	0.9375	0.875	0.875	0.8125
8	0.9375	0.9375	0.9375	0.875
9	0.9375	0.875	0.8125	0.75
10	0.9375	0.875	0.8125	0.8125
11	0.9375	0.9375	0.875	0.875
12	0.9375	0.9375	0.9375	0.875
13	0.8125	0.8125	0.8125	0.75
14	0.9375	0.875	0.8125	0.75
15	0.9375	0.9375	0.9375	0.875

Sensitivity is determined using 240 samples taken from MIT-BIH.We took 16 samples from each class in those samples. There are 15 classes in all. Positive samples total (P) for each class: 16 Negative Samples in Total or Remaining224th class (N) True Positive (TP): Total number of correctly retrieved samples in current class (out of 16) True Negative (TN): Total number of correctly retrieved samples from the remaining classes (Out of 224)

The sensitivity gained for each class is shown in the table 3. Each row represents a class, and each column represents the accuracy achieved using a suggested or current Classifier. The presented method is more accurate than other conventional methods.

4.4 False positive rate (FPR)

False positive rate is determined using 240 samples taken from MIT-BIH.

We took 16 samples from each class in those samples. There are 15 classes in all. Positive samples total (P) for each class: 16 Negative Samples in Total or Remaining 224th class (N) True Positive (TP): Total number of correctly retrieved samples in current class (out of 16) True Negative (TN): Total number of correctly retrieved samples from the remaining classes (Out of 224).The False positive rate can be calculated by the equation

FPR=[FP] / [N] (7)

Table 4. Comparison fo different methods for False positive rate .

Sr.No.	ISVR	SVM	ANN	NMD
1	0.004444	0.008889	0.013333	0.013333
2	0.008889	0.013274	0.017621	0.022026
3	0.004484	0.004505	0.004505	0.004505
4	0.013216	0.013216	0.013216	0.013274
5	0.004464	0.004484	0.008929	0.013333
6	0.004464	0.004464	0.008929	0.008929
7	0.004464	0.008929	0.008929	0.013333
8	0.004444	0.004444	0.004464	0.008889
9	0.004464	0.008929	0.013333	0.017778
10	0.004464	0.008929	0.013393	0.013393
11	0.004444	0.004444	0.00885	0.00885
12	0.004444	0.004464	0.004484	0.008969
13	0.013216	0.013216	0.013216	0.017544
14	0.004444	0.00885	0.013274	0.017699
15	0.004464	0.004464	0.004484	0.008929

The false positive rate gained for each class is shown in the table 4. Each row represents a class and each column represents the accuracy achieved using a suggested or current Classifier. The presented method is more accurate than other conventional methods.

4.5 False Negative rate (FNR)

False negative rate is determined using 240 samples taken from MIT-BIH.

We took 16 samples from each class in those samples. There are 15 classes in all. Positive samples total (P) for each class: 16 Negative Samples in Total or Remaining224th class (N) True Positive (TP): Total number of correctly retrieved samples in current class (out of 16) True Negative (TN): Total number of correctly retrieved samples from the remaining classes (Out of 224).The False negative rate can be calculated by the equation

FPR=[FN] / [N] (8)

Table 5. Comparison of different methods for False negative rate .

Sr.No.	FNR_SVR	FNR_SVM	FNR_ANN	FNR_NMD
1	0	0.066667	0.133333	0.133333
2	0.066667	0.071429	0.076923	0.153846
3	0.117647	0.166667	0.166667	0.166667
4	0	0	0	0.071429
5	0.0625	0.117647	0.125	0.133333
6	0.0625	0.0625	0.125	0.125
7	0.0625	0.125	0.125	0.133333
8	0	0	0.0625	0.066667
9	0.0625	0.125	0.133333	0.2
10	0.0625	0.125	0.1875	0.1875
11	0	0	0	0
12	0	0.0625	0.117647	0.176471
13	0	0	0	0
14	0	0	0.071429	0.142857
15	0.0625	0.0625	0.117647	0.125

The False negative rate gained for each class is shown in the table 5. Each row represents a class, and each column represents the accuracy achieved using a suggested or current Classifier. The presented method is more accurate than other conventional methods.

4.6 Positive Predictive Value (PPV)

Positive Predictive Value is determined using 240 samples taken from MIT-BIH.

We took 16 samples from each class in those samples.

There are 15 classes in all. Positive samples total (P) for each class: 16 Negative samples in total or remaining 224th class (N) True Positive (TP): total number of correctly retrieved samples in current class (out of 16) true negative (TN): Total number of correctly retrieved samples from the remaining classes (Out of 224).The Positive Predictive Value can be calculated by the equation

$$PPV=[TP] / [TP + FP] \qquad (9)$$

The Positive Predictive Value gained for each class is shown in the table 6. Each row represents a class, and each column represents the accuracy achieved using a suggested or current classifier. The presented method is more accurate than other conventional methods. From the above results the ISVR method shows the better results than other method.

Table 6. Comparison of different methods for Positive Predictive Value (PPV)

Sr.No.	PPV_SVR	PPV_SVM	PPV_ANN	PPV_NMD
1	0.9375	0.875	0.8125	0.8125
2	0.875	0.8125	0.75	0.6875
3	0.9375	0.9375	0.9375	0.9375
4	0.8125	0.8125	0.8125	0.8125
5	0.9375	0.9375	0.875	0.8125
6	0.9375	0.9375	0.875	0.875
7	0.9375	0.875	0.875	0.8125
8	0.9375	0.9375	0.9375	0.875
9	0.9375	0.875	0.8125	0.75
10	0.9375	0.875	0.8125	0.8125
11	0.9375	0.9375	0.875	0.875
12	0.9375	0.9375	0.9375	0.875
13	0.8125	0.8125	0.8125	0.75
14	0.9375	0.875	0.8125	0.75
15	0.9375	0.9375	0.9375	0.875

5. Conclusion

This work suggests and investigates an ECG processing and analysis system with two principal applications. A fifteen-class ECG classification system that can distinguish between normal (healthy) and abnormal (unhealthy) ECG signals. The suggested ECG auto categorization system design is implemented using characteristics taken from the Local and MIT-BIH Datasets. The proposed ECG classification algorithms in this study correctly classified normal and abnormal ECG signals at a rate of 99 percent.

Proposed algorithm performed better than existing method. Proposed Architecture uses both time and frequency domain features for classification purpose. Due to use of higher order statistic our classification problem becomes simpler than traditional morphological Feature. Proposed algorithm delivered high performance even with smaller learning data.Modeling accuracy for ISVR Classification models can reach 98 percent. When they applied to imitate ECG signals, they were met with a lot of resistance.

In this paper comparison of SVM, ANN, Neural Mode Decomposition (NMD) and Incremental Support Vector Regression (ISVR) methods are presents. From these methods, theISVR method shows better results.

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