TS114 Project

Computer-aided analysis of electrocardiogram signals

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Introduction

The heart has always been an acurate indicator of one's health. Analyzing it and monitoring its activity is of utmost importance for clinicians. Thus this project's aim is to develop an automatic and user-friendly application capable of giving useful information about the heart activity, such as the cardiac rhythm or the detection of heart conditions, based on an electrocardiogram signal.

The project is divided in multiple parts. First, different ECG signals (from both ill and healthy people) were displayed and analysed under MATLAB to highlight their characteristics. Followed by the automated detection of the latter, as well as the identification of cardiac pathologies and the ECG signals denoising. Finally, the ultimate product is delivered with a Graphical User Interface (GUI) making it possible to the clinicians to use it easily and intuitively.

ECG visualization

In this first part, seven different ECG signals will be analyzed and displayed under MATLAB.

Three of these are records from healthy patient's heart, whereas the other four are from ill ones, each with different pathologies.

This part's aim is to study these signals in the frequency and time domain, starting by the latter, and underline the different caracteristics and properties of these.

1.1 Time display

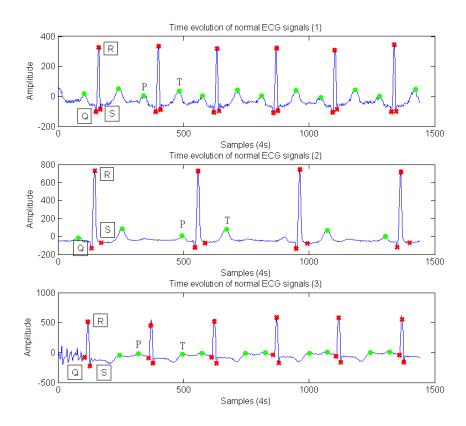


Figure 1.1: Normal ECG signals with the Q, R, S points and P, T waves highlighted

First, three seconds of each of the aforementioned normal ECG signals (i.e. from healthy heart) were plotted under MATLAB. On the figure above (Figure 1.1), their Q, R and S points are represented by red

dots, while their P and T waves are located by green dots.

The first signal displayed is noisier than the two others, although, its characteristic points and waves are easier to observe.

The second signal is uncluttered, but the S point is hardly recognizable.

Finally, the P and T waves of the third signal, which is a tad noisy, are nearly overlapping.

Based on these four seconds samples, the cardiac rhythm of each patients was computed and displayed in the table below.

Signal number	Cardiac rhythm (bpm)
1	91.7
2	53.0
3	86.4

According to the graphs and the computed cardiac rhythms, it can be assumed that the faster the heart-beat is, the noisier the ECG will be. Indeed, the first signal is the fastest heartbeat's ECG while being the noisiest. The same can be said of the third signal.

Moreover, the second signal represents the ECG of a heart with a slow heartbeat and since it is a normal ECG, it might be the heartbeat of an athletic person (otherwise it could be a heart condition like brady-cardia).

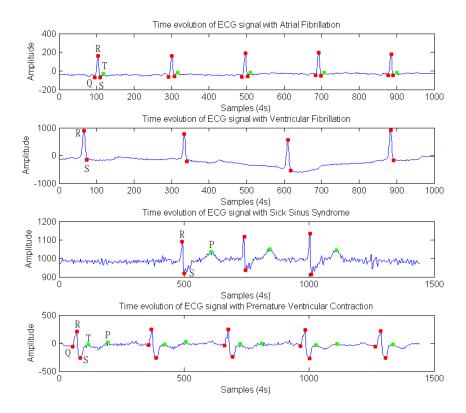


Figure 1.2: ECG signals with pathologies with the Q, R, S points and P, T waves highlighted

Then, the last four ECG signals associated with heart conditions were plotted under MATLAB.

The corresponding cardiac rhythms were computed and displayed in the table below.

Pathology	Cardiac rhythm (bpm)
Atrial Fibrillation	76.7
Ventricular Fibrillation	55.5
Sick Sinus Syndrome	84.7
Premature Ventricular Contraction	71.1

These pathologies have an impact on each ECG signals and useful characteristics are lost.

Compared to the normal signals, the Atrial Fibrillation's ECG signals lack P waves and has weak impulsions. In the same way, for the Ventricular Fibrillation, the signal depicts a slow heartbeat and hardly discernible P waves, as well as Q and S points.

The Sick Sinus Syndrome has a different effect on the signal, for it creates groups of P waves and R/S points with an absence of activity in between.

The Premature Ventricular Contraction pathology is the closest match, out of the four signals, to a typical one. But, as the pathology's name suggest, the P waves are ahead compared to a normal signal.

1.2 Frequency display

At first, the ECG power spectrum of the normal ECG signals. N samples were used, with $N = 15 \cdot F_s$.

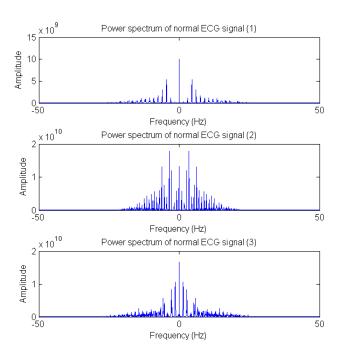


Figure 1.3: ECG power spectrum of the normal ECG signals

Since the original signals are periodic, peaks can be found every $f = \frac{n}{T_b}, \forall n \in \mathbb{N}$.

 T_b represents the number of beats per second. It is then possible to compute the different cardiac rhythms which are shown in the table below.

Signal number	Cardiac rhythm (bpm)
1	91.8
2	52.2
3	81.9

The results are pretty close to what was computed in the time domain.

Then, the signals associated with pathologies were plotted under MATLAB.

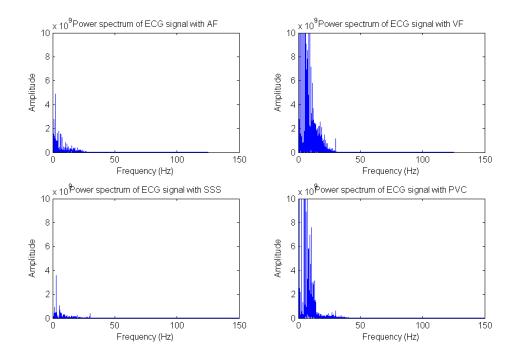


Figure 1.4: ECG power spectrum of the ECG signals associated with pathologies

These graphs were cut off, otherwise, since the component in f = 0 has a very high value, the scale follow suits and it is almost impossible to see the other components.

The differences between these power spectrum and those from the normal ECG, are firstly, as it was said above, the value of the f = 0 component.

jį Missing part į,į,

Out of the previous results, the cardiac rhythm were computed and displayed in the table below.

Pathology	Cardiac rhythm (bpm)
Atrial Fibrillation	76.8
Ventricular Fibrillation	57.6
Sick Sinus Syndrome	82.8
Premature Ventricular Contraction	72

Just as before, the results match those computed in the time domain.

Detection of P, QRS and T waves

2.1 Three different methods of R waves detection

The purpose of this section is to implement and compare three different methods in order to detect the R waves.

2.1.1 Method of local maxima

This method is rather straightforward. A window is set to work on a restricted number of samples. The signal is then cubed to make the R waves stand out.

A threshold r is chosen to only select the peaks corresponding to the R waves. r is a variable equal to a certain percentage of the maximum value contained in the ECG signals.

When the peaks are picked, the local maxima of the latter is selected and the corresponding times taken out. Finally, the R points are plotted on the original ECG at the previously found positions.

2.1.2 Method of the derivative

The derived ECG signals presents peaks with a local maxima followed by a local minima. Using the previous method, these different peaks are located with their beginning and ending times. The latter is transposed to the original ECG signals to get the different R waves domains. The R points correspond to the local maximum in each given domain.

2.1.3 Pan and Tompkins algorithm

As opposed to the previous methods, the Pan and Tompkins algorithm consists of different steps.

First, the ECG is signal is filtered by a band-pass filter, which is composed of a low pass-filter and a high pass-filter.

The resulting signal is filtered by a derivative signal to gain information about the QRS slope and then squared to make the local maxima stand out.

Then, the signal is integrated with a moving-window method. The width of the said window is N which is the empirically-determined duration of a QRS complex.

Finally, a threshold is applied on the integrated signal. Thanks to this step, the temporal location of the QRS complex can be extracted from the rising slope in the latter.

2.2 Q and S waves detection

Based on the R waves' location, it is now possible to detect the Q and S waves. They are respectively defined as the first minima preceding and following the R waves.

This definition is unfortunately too plain to be effective. If the signal happens to be a tad noisy, another minimum could be detected between the R wave and the actual Q or S wave.

2.3 P and T waves detection

Automatic identification of cardiac pathologies

Now that the P, Q, R, S and T waves detection was achieved, it can be put to a practical use. Indeed, in the following part, different pathologies identification methods will be studied and implemented.

3.1 Tachycardia/Bradycardia

The cardiac rhythm has been defined as the time between two R waves. Tachycardia (respectively bradycardia) is declared when the former is over 100 bpm (respectively under 60 bpm).

A simple method was used to compute the cardiac rhythm. At first, 4 seconds of the given ECG are isolated, the R waves are then detected and the time between them computed. All these times are summed and then divided by the number of R-R intervals to obtain an average time between the different R waves. The cardiac rhythm is finally computed based on the previous result.

Identifying basic arrhythmia pathologies now boils down to a simple comparison between the cardiac rhythm and the threshold defining these pathologies (60 bpm and 100 bpm in this case).

3.2 Heart rate variability

ECG denoising

Conclusion