

The parameters of the ECG signal are obtained by the wavelet decomposition dyadic tree. This tree decomposes the signal into the smooth (low pass) and detail (high pass) components. To estimate the said ECG parameters (QRS complex, T wave, P wave, locations and durations) in this work the following algorithm was used : Selecting a wavelet-transformed ECG block data, determination of the R wave location (as maxima) (scale 1) (which occurs at the zero crossing point between the most prominent maxima), determination of R-R intervals, as R-R distances, determination of Q, S points as the first zero crossing point before and after R wave, elimination of the QRS from the signal to obtain the other parameters, determination of the P wave location (as maxima) (scales 3,4) , and the P-Q distance, elimination of the P wave from the signal (same as step 5), determination of the T wave location (as the remained maxima) (scales 3,4), and S-T segments durations. This algorithm leads to determine the main parameters of an ECG signals. Were used over 27 files from the MIT-BIH database, signals containing normal sinus rhythms and signals with abnormalities in order to find the main parameters.

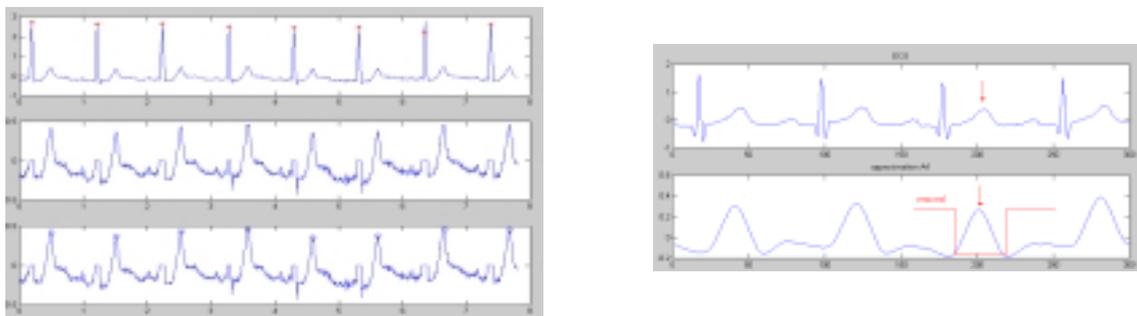


fig 5. R wave detection, QRS complex removal, T wave identification

3. Results

The results obtained (processing mainly ECG signals from normal sinus database) were compared with annotated files from ECG databases, and gave very promising results: R wave detection around 98%, R-R interval (HRV) determination 95% QRS complex detection over 91%, T wave detection/localization 88%, P wave detection/localization 78%. (27 files used from MIT-BIH database, signals were denoised and with the baseline drift removed)

4. Conclusions

The present study, based on biorthogonal wavelets, shown that the various morphologies of the ECG signals can be identified using wavelet decomposition at different scales.

5. References

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