Assignment 2 IM - ECG Analysis

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1 Introduction

Following the assignment request, two different systems have been developed: a detector for a PVC and a detector for Atrial Fibrillation occurencies. These two classifiers take as input the ECG data from a patient, extract the relevant features such as the R peaks, and output a binary value that reflects whether the events have occured or not.
The performances of the systems will be compared using sensitivity and specificity parameters.

2 Feature Extraction

The first module implemented in order to perform detection of both PVC and AF is a feature extracting function that allows the acquisition of the R peaks in the shape of an ECG signal. These peaks, in fact, are used to acquire the relevant information about an ECG signal, rapresenting each an atomic event of pulsation.
The function "detectPeaks()" takes as input the ecg raw data and the sampling frequence and outputs a vector of indexes corresponding to the peaks in the data. The code does some pre-filtering on the data itself before of computing the peaks, as it can be seen:

```
fs = frequence;

% analyse peaks
% lowpass filter
order = 4;
wc = 20;
fc = wc / (0.5 * fs);
[b, a] = butter(order, fc, 'low');
e1 = filtfilt(b, a, ecg);

% highpass filter
order = 4;
wc = 5;
fc = wc / (0.5 * fs);
[b, a] = butter(order, fc, 'high');
e2 = filtfilt(b, a, e1);

% differentiation
e3 = diff(e2);

```

```
% potentiation
e4 = e3.^2;

% moving average
timeWindow = 0.2;
N = timeWindow * fs;
b  = (1 / N) * ones(1, N);
a  = 1;
e5 = filter (b, a, e4);
```

Code 1: Pre-filtering

After the filtering is done, the peaks are computed:

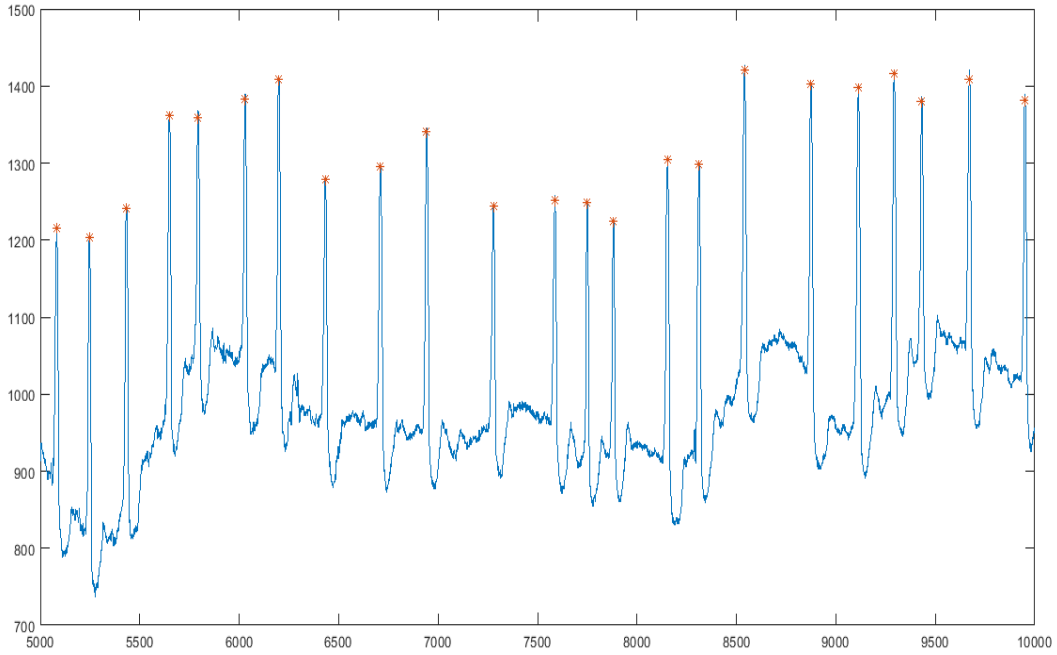
```
% find R peaks
threshold = 0.7 * mean(e5);
pause = 0.3;
indexPause = pause * fs;
peaks = [];
index = 1;
i = 2;
while i<=length(e5)
   if e5(i)==threshold || e5(i-1) < threshold && e5(i) > threshold
      peaks(index) = i;
      index = index + 1;
      i = i + indexPause;
   else
      i = i + 1;
   end
end

% set position
back = 0.2;
backIndex = back * fs;
for i=1:length(peaks)
   minIndex = peaks(i)-backIndex;
   stepBack = backIndex;
   if minIndex <= 0
      stepBack = backIndex + minIndex-2;
      minIndex = 1;
   end
   maxIndex = peaks(i)+backIndex;
   if maxIndex > length(ecg)
      maxIndex = length(ecg);
   end
   tempECG = ecg(minIndex:maxIndex);
   [~, maxIndex] = max(tempECG);
   peaks(i) = peaks(i) + maxIndex - stepBack-2;
```

```
35 end
36
37 peaks = peaks';
```

Code 2: Peak detection

The last code snippet has the task of readjusting the position of the found peaks. An example of the detected peaks can be seen in the next image, obtained plotting on a same axis the ecg data and the peaks.



3 PVC detection

3.1 Methods

Once the R peak of the ECG are detected, three different criterium have been chosen to perform PVC detection: anomalies in peak-to-peak (RR) intervals, QRS area comparison and hermite polynomial analysis.
The R-R regularity criterium has the following structure: it first detects the intervals between each couple of peaks, and then computes how much every interval duration is different from the mean duration. The code is the following:

```
% RR regularity
RRIntervals = zeros(length(ind)-1,1);
for i=2:length(ind)
    RRIntervals(i-1) = ind(i) - ind(i-1);
end
meanRR = mean(RRIntervals);
sdnnRR = calculateSDNNRR(peaks);

myPVCRR = zeros(length(RRIntervals)+1,1);
for i=1:length(RRIntervals)
   myPVCRR(i) = RRIntervals(i) / (meanRR + sdnnRR);
end
```

Code 3: RR criterium

The QRS area criterium makes the computation of the areas below the QRS complexes in the data, and then, like the R-R regularity criterium, compares it with the average area in the acquired data. Specifically the area was computed taking the 0.06 seconds around a peak and calculating the area below the resulting graph via the MATLAB method *trapz*(). This is the code implementing the criterium:

```
% QRS area
area = 0.06;
areaIndex = round(area * fs);

areas = zeros(length(ind),1);
for i=1:length(ind)
    minIndex = ind(i)-areaIndex;
    if minIndex <= 0
        minIndex = 1;
    end
    maxIndex = ind(i)+areaIndex;
    if maxIndex > length(ecg)
        maxIndex = length(ecg);
    end
    actualAreaECG = ecg(minIndex:maxIndex);
    areas(i) = trapz(actualAreaECG);
end

meanArea = mean(areas);

sum = 0;
for i=1:length(areas)
    sum = sum + (areas(i) - meanArea).^2;
end
```

```
25
26 sdnnArea = sqrt(sum / length(RRIntervals));
27 myPVCArea = areas(:) / (meanArea + sdnnArea);
```

Code 4: QRS area criterium

The Hermite criterium makes use of system modeling to consider the data as the output of an ar system. Then, with respect to the location of the poles of the actual system, if they are out of the (hyper-)circonference of radius 1, which is to say correspond to an unstable system, the considered event corresponds to a PVC occurrence. In order for the model to work, the peak considered should be necessarily a positive value, so a pre-processing of the peaks has been implemented in order to get coherent data.

To get the model for the ar system, the 8 values before and after a peak are taken into account, and so a window of 17 timesteps is obtained. Then, the MATLAB "ar" method is called with a desired order of 2. This has been chosen after considering the perfomances of the criterium with an order ranging from 1 to 8.

The roots of the system are found with the MATLAB function "roots()" and then their norm is computed to check if it is greater than 1. Actually, the threshold used is not 1, since after having performed some tests, that would have been too strong and therefore considered every sample as a non-PVC. Finally, the code is as follows:

```
1 % hermite
2 % Code to get positive peaks
3 for i=1:length(ind)
4     if(ind(i)>50 && ind(i)+50<length(ecg))
5         start=ind(i)-50;
6         finish=ind(i)+50;
7     elseif (ind(i)<50) % case first peak is before 50
8         start=1;
9         finish=ind(i)+50;
10    else % case for last peak
11        start=ind(i)-50;
12        finish=length(ecg);
13    end
14    A=ecg(start:finish);
15    [~,index]=max(A); % find positive peak
16    ind(i)=start+index; % code to reassign peak to positive values
17 end
18
19 myPVCHermit = zeros(length(ind),1);
```

```
for j=1:length(ind) % Take into account first and last case later
    a=8;
    b=8;

    minIndex = ind(j)-a;
    if minIndex <= 0
        minIndex = 1;
    end

    maxIndex = ind(j)+b;
    if maxIndex > length(ind)
        maxIndex = length(ind);
    end

    window=ecg(minIndex:maxIndex);

    model=ar(window,2);
    A=model.A;
    poles=roots(A);
    tmppvc=0;
    for i=1:length(poles)
        if norm(poles(i))>=0.98
            tmppvc=1;
            break;
        end
    end
    myPVCHermit(j)= tmppvc;
end
```

Code 5: Hermite criterium

3.2 Classification

As it can be noted, while the detection via RR regularities and QRS areas give a "classification" score, a decimal value that quantificates a certain probability of the event being a PVC, the hermite criterium output is a "simple" binary decision between PVC/not PVC. Thus, a specific criterium has to be chosen to combine the different results. Since the RR regularities or the QRS areas criterium have a really good specificity, if they report a PVC then the global output is set to be a detected PVC: this is why they are OR'd with the third criterium. The third OR case is a AND combination of two criteriums: the combination of RR and QRS scores if they are both quite high but not enough to be greater than 1 by themselves, and a detected PVC from the Hermite criterium. In fact, Hermite criterium has a sensitivity higher than RR+QRS, and as such it eventually compensates combining RR and QRS. All these considerations have

been done after experimental testing on specificity and sensitivity results that can be found in the "Result section". So, the resulting rule and thresholds are the following:

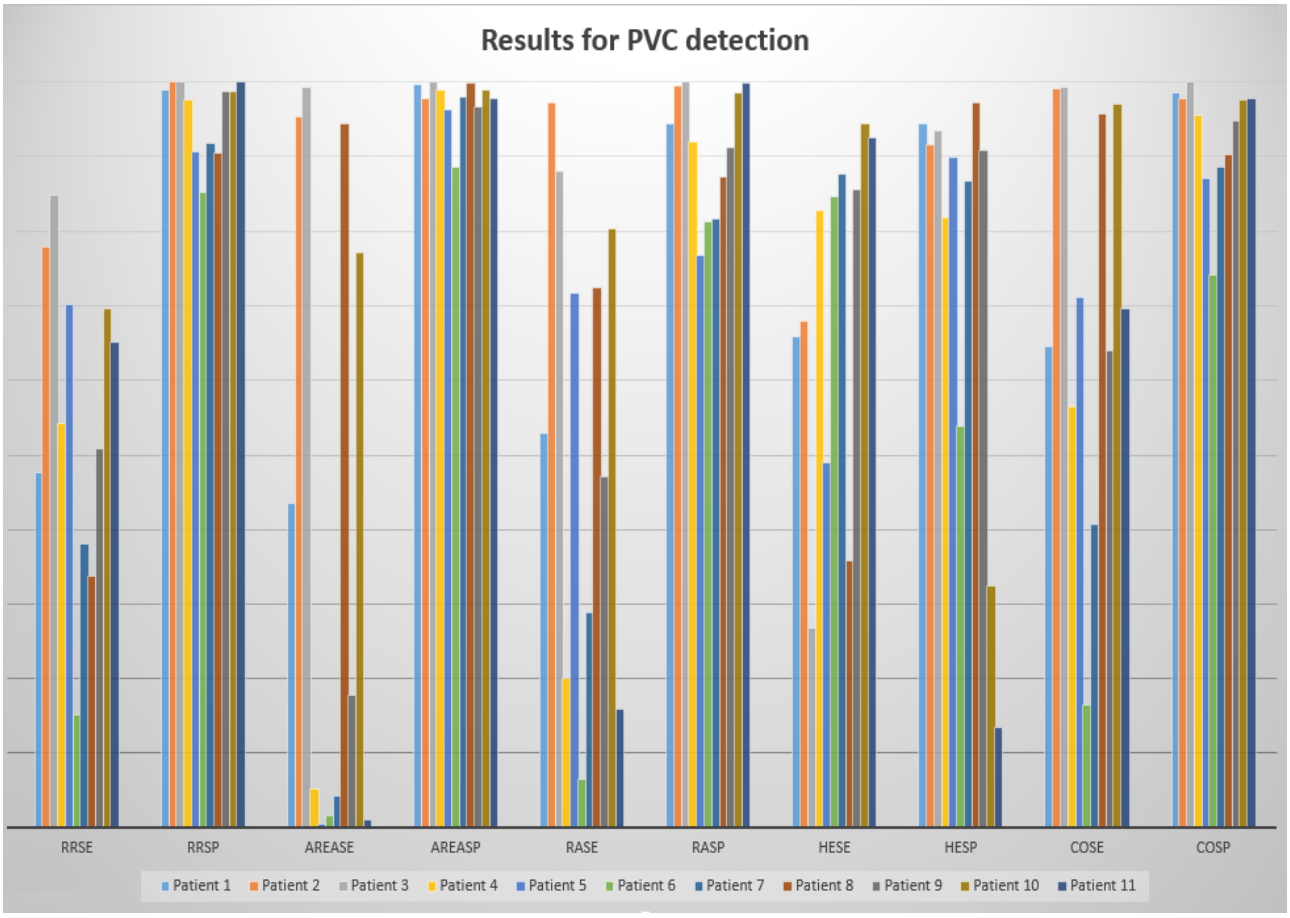
$myPVCRR(i) >= 1 || myPVCArea(i) >= 1 || (myPVCArea(i) + myPVCRR(i) >= 1.9) \&\& myPVCHermit(i) == 1$

3.3 Results

The following table shows the results obtained running the classifier on each patient with five different classification rules: the use of the three "simple" criteriums seen, as well as the combination of RR detection and QRS as well as the total combination of the five of them using the rule described in the last section. The last row of the table is obtained averaging the values: as it can be seen it gives a good insight over the efficiency of the implemented criteriums: the first three have a really good specificity but low sensitivity, while it's the opposite for the Hermite classifier. Finally, the combination of the three shows a definite and solid increase in both sensitivity and specificity.

	RR		QVC area		RR+QVC		Hermite		Combined	
	SE	SP	SE	SP	SE	SP	SE	SP	SE	SP
Patient 1	0.48	0.99	0.44	0.99	0.53	0.94	0.66	0.94	0.65	0.99
Patient 2	0.78	1	0.95	0.98	0.97	0.99	0.68	0.91	0.99	0.98
Patient 3	0.85	1	0.99	1	0.88	1	0.27	0.93	0.99	1
Patient 4	0.54	0.97	0.05	0.99	0.20	0.92	0.83	0.82	0.56	0.95
Patient 5	0.70	0.91	0	0.96	0.72	0.77	0.49	0.90	0.72	0.87
Patient 6	0.15	0.85	0.01	0.88	0.06	0.81	0.85	0.54	0.16	0.74
Patient 7	0.38	0.91	0.04	0.98	0.29	0.81	0.88	0.87	0.41	0.88
Patient 8	0.33	0.90	0.94	1	0.72	0.87	0.36	0.97	0.96	0.90
Patient 9	0.51	0.99	0.18	0.97	0.47	0.91	0.86	0.91	0.64	0.95
Patient 10	0.70	0.99	0.77	0.99	0.80	0.98	0.94	0.32	0.97	0.98
Patient 11	0.65	1	0.01	0.98	0.16	1	0.92	0.13	0.70	0.98
Average	0.55	0.96	0.40	0.97	0.53	0.91	0.70	0.75	0.70	0.93

A visual rapresentation of the performances can be seen in the following graph, where each bar is the result for a patient, while "clusters" of bars are representative of a criterium/metric.


Results for PVC detection
RRSE
RRSP
AREASE
AREASP
RASE
RASP
HESE
HESP
COSE
COSP
Patient 1
Patient 2
Patient 3
Patient 4
Patient 5
Patient 6
Patient 7
Patient 8
Patient 9
Patient 10
Patient 11

4 AF

For AF detection, two different criterium have been used: a

4.1 Methods

- RR intervals: same as for PVC just for all the different windows, than we compare the window values and try to get smart assumptions - frequency: just folllowing the slides -¿ output is shit TODO ??

4.2 Classification

For the AF detection we consider the difference of the RR intervals and the frequency for different areas of the ecg.

$sdnnRRWindows(i) < 0.9 \&\& lfhfWindows(i) < 0.4$

blabla TODO

4.3 Results

TODO Gerardo

5 Conclusions

5.1 PVC

MUITO BEM!!!

5.2 AF

0.5 * muito bem!