User Documentation: ECG Noise Extraction (ECG_NExT) Software

This software outputs an estimate of a noise signal collected as part of an electrocardiograph (ECG) recording. The objective is to obtain representative samples of noise and motion artifacts from ECG devices under test that can then be used for testing the robustness of ECG analysis algorithms. The algorithm for noise estimation is described in [1]. The core operation of this software is based on removing the instances of the QRS complex from the ECG so that an estimate of the noise component of the recording remains. Required inputs are the ECG signal recorded with noise/motion artifacts and locations (indices) of the R-peaks.

1. System Requirements

The software was developed in a MATLAB® 2021b environment and signal processing toolbox was extensively used when testing the software. Thus, the signal processing toolbox is a minimum requirement.

2. Data preparation (as in TestData.mat in the software package):

The following steps are taken to prepare data for processing and noise estimation by this software:

1. Obtain an ECG signal using the device under test – ECG_device – at a location where noise/motion artifacts are captured in the ECG signal.
2. Obtain a simultaneously recorded clean ECG signal from a standard site of collection, e. g., the chest area for reference – ECG_std.
3. Apply a bandpass filter to each ECG signal to obtain ECG_device_BP and ECG_std_BP. This is to clean the data from baseline wander and high-frequency noises as would normally be performed as part of data acquisition/processing for the device under test – typically in the range of 0.5 to 40 Hz.
4. Obtain the indices of the R peaks from, e. g., ECG_std_BP (or ECG_device_BP if it provides sufficient fidelity for reliable peak detection).

3. Input and Output of ECG_NExT

Inputs:

1. ECG_device with at least 30 seconds length (the TestData provides 60 seconds worth of data)
2. Bandpass-filtered ECG_device, ECG_device_BP.
3. R peak indices.
4. Sampling rate of 1 kHz. Both ECG_device and ECG_std should be sampled at this rate. The software will provide an error message if signals with a different sampling rate are provided.
5. Optional inputs of Blank and Blend parameters: This is a two-element array; the first element, the Blank parameter, specifies the length of data (in seconds) to be removed

from both sides of each R peak. The second element, the Blend parameter, specifies the length of data (in seconds) to be copied from neighboring segments of the removed data, crossfaded and inserted in the gap created by the process of removal of the area around the R peaks. The default values for Blank and Blend parameters are 40 and 60 milliseconds, respectively. This optional input allows the user to adjust for the differences in R-Peak widths within a data set. For more details please refer to [1].

Outputs:

1. Estimated Noise sample.
2. A time vector for demonstration purposes, e. g., plotting original signal and the estimated noise versus time on a shared horizontal axis. The removed segments from the original signals are accounted for in this time vector.

4. Algorithm Development Test Conditions:

The test signals used during the software development had the following characteristics:

1. Signals equivalent to ECG_Device and ECG_std were 60-second-long signals sampled at 1000 Hz.
2. The signals were passed through a Butterworth 5^{th} order bandpass filter with 0.5 and 40 Hz corner frequencies corresponding to low and high cutoff frequencies, respectively.

5. Software Package Content:

The software package includes the core function for ECG noise estimation (NoiseExtractClass.m). In addition, a sample wrapper/shell code is provided to run the function with a test set consisting of a 60-minute-long clean ECG sampled at 1000 Hz and a noise segment of the same length randomly generated from a Gaussian distribution.

References

1. Galeotti, L. and C.G. Scully, *A method to extract realistic artifacts from electrocardiogram recordings for robust algorithm testing.* Journal of Electrocardiology, 2018. **51**(6, Supplement): p. S56-S60.

Appendix I

Software Code

The shell code:

This is the primary module that runs the core software and initiates various function calls.

```
close all
clear
clc
% =+=+=+=+=+=+=+=+=+=+=+=+=+ User data +=+=+=+=+=+=+=+=+=+=+=+=+=+=+=+=+=+=+=+=+
% User data can be added here and should replace the test data below:
% =+=+=+=+=+=+=+=+=+=+ End of user data =+=+=+=+=+=+=+=+=+=+=+=+=+=+=+=+=+=+=+=+

% =+=+=+=+=+=+=+=+=+=+=+=+=+ Test data +=+=+=+=+=+=+=+=+=+=+=+=+=+=+=+=+=+=+=+=+
testDat = load('TestData.mat');
ECG_device = testDat.ECG_device;
ECG_device_BP = testDat.ECG_device_BP;
rPeaks = testDat.rPeaks;
fs = testDat.fs;

% =+=+=+=+=+=+=+=+=+=+=+ End of test data =+=+=+=+=+=+=+=+=+=+=+=+=+=+=+=+=+=+=+=+=+
NEWC = NoiseExtractClass;
[noiseEstimated, tVec] = NEWC.ecgNoiseExtractor(ECG_device, ECG_device_BP,
rPeaks, fs);

% Reduce noise record to same length as calculated noise record
NEWC.TestPlot(noiseEstimated, tVec, fs, ECG_device, testDat.ECG_std)
```

The main code:

```
classdef NoiseExtractClass
%{
This classdef construct includes the functions and sub modules necessary
to implement the ECG_NExT algorithm that obtains representative samples
of noise and motion artifacts from ECG devices under test.

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Galeotti, L. and C.G. Scully, "A method to extract realistic artifacts from
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that any derivative works bear some notice that they are derived from it,
and any modified versions bear some notice that they have been modified.
%}
    properties
        Property1
    end

    methods (Static)
        function [NOISE_Calc, tVec] = ecgNoiseExtractor(ECG_test, ...
                ECGtest_BP, rPeaksRef, fs, varargin)
            %     [NOISE_Calc, tVec] = ecgNoiseExtractor(ECG_test, ...
            % ECGtest_BP, rPeaksRef, fs)
            %
            % This function extracts noise from ECG recording as described
            % in Galeotti and Scully, JECG 2018.
            %
```

```
% Inputs:
% ECG_test: The ECG signal obtained from the device under test.
% ECGtest_BP: The bandpass-filtered version of ECG_test.
% rPeaksRef: Accurately and reliably detected R peak indices of
%            the ECG obtained.
% fs: Sampling frequency at which the ECG signals are sampled.
%
% Outputs:
% NOISE_Calc: The calculated noise using this algorithm/code.
% tVec: Contains the time references corresponding to the noise
% instances remained after removing areas surrounding the R
% peaks. This is only needed for demo purposes when plotting the
% noise and ECG under consideration against time is desired.
%
% [...] = ecgNoiseExtractor(..., CROSSFADE_PAR) accepts a
% two-element array of Blank and Blend parameters ([Blank Blend])
% specifying the amount of data be removed from around the R
% peaks (Blank in seconds) and the amount of data to be copied
% from the neighboring sides of the removed part (Blend in
% seconds) to be cross faded and used to fill the gap
% created after removing the data from around the R peaks.
% The default values for the Blank and Blend parameters are set
% to 40 ms and 60 ms, respectively.

% Check for sampling frequency:
% Although the code should work with any sampling frequency, the
% sampling frequency used during test and development of this
% software was 1000 Hz. We, therefore, make certain that the
% users provide signals with 1000 Hz sampling frequency.
if fs ~= 1000
    error('Please change the sampling frequency to 1 kHz.')
end

sigLen = length(ECGtest_BP);

% Track time:
tTemp = (0:sigLen - 1)/fs;

% Ensure consistent dimensionality of 1xn -- row vectors:
ECG_test = ECG_test(:)';
ECGtest_BP = ECGtest_BP(:)';
rPeaksRef = rPeaksRef(:)';

% Get residuals from test ECG after filtering (noise components
% outside bandwidth of bandpass filtered signal)
ECGtest_residuals = ECG_test - ECGtest_BP;

% Obtain median beat from filtered ECG signal
[MedBeat] = ComputeMedianBeat_long(ECGtest_BP, rPeaksRef, fs);

% Generate synthetic signal using the median beat:
[SynECG] = GenSynECG_blend(MedBeat, rPeaksRef, sigLen, fs);

% Obtain the noise component of the signal:
noiseRaw = ECGtest_BP - SynECG;

% Remove areas around the R-peaks in the obtained noise as well
```

```
    % as in the residuals obtained above and the time vector:

    % Handle optional inputs:
    if isempty(varargin)
        NOISE_Calc = RemRpkEffect(noiseRaw, rPeaksRef, fs);
        residualLessRpeaks = RemRpkEffect(ECGtest_residuals, …
          rPeaksRef, fs);
        tVec = RemRpkEffect(tTemp, rPeaksRef, fs);
    else
        NOISE_Calc = RemRpkEffect(noiseRaw, rPeaksRef, fs, …
          varargin{1});
        residualLessRpeaks = RemRpkEffect(ECGtest_residuals, …
          rPeaksRef, fs, varargin{1});
        tVec = RemRpkEffect(tTemp, rPeaksRef, fs, varargin{1});
    end

    % Add noise to residual from bandpass filtered signal:
    NOISE_Calc = NOISE_Calc + residualLessRpeaks;

    % Remove NaN instances:
    tVec = tVec(~isnan(NOISE_Calc));
    NOISE_Calc = NOISE_Calc(~isnan(NOISE_Calc));

% End of ecgNoiseExtractor function
end

function TestPlot(noiseEstimated, tVec, fs, ECG_raw, ECG_ref)
    % This is a supplemental function for demo purposes in case the
    % user wants to plot the obtained noise with the ECG under
    % consideration.
    %
    % Inputs:
    % noiseEstimated: The estimated noise by the ECG_NExT algorithm.
    % tVec: the vector of time references corresponding to the
    %       estimated noise samples in noiseEstimated.
    % fs: sampling frequency
    % ECG_raw: The ECG collected from the device under test.
    % ECG_ref: The clean reference ECG (if collected). If not
    %       provided, the ECG_raw will be used.

    % Handle optional input(s):
    if nargin < 5
        ECG_ref = ECG_raw;
    end
    figure
    tECG = (0:length(ECG_raw)-1)/fs;
    ax(1) = subplot(3,1,1);
    plot(tECG, ECG_ref)
    title('Reference ECG')
    ax(2) = subplot(3,1,2);
    plot(tECG, ECG_raw)
    title('Unfiltered Device ECG')
    ax(3) = subplot(3,1,3);
    plot(tVec, noiseEstimated)
    title('Estimated Noise')
    linkaxes(ax, 'x')
    xlabel('Time (sec)')
```

```
            set(gcf, 'units', 'normal')
            set(gcf, 'position', [0.1753    0.0977    0.5064    0.7831])

        % End of function
        end

    % End of method
    end
% End of classdef
end

function [sigOut, stitch] = RemRpkEffect(sigIn, Rpeaks, fs, varargin)
    % This function removes an area around R peaks and merges the segments
    % surrounding the removed area with crossfade of neighboring signal
    % segments around the removed area.
    %
    % Inputs:
    % sigIn: This input can represent an ECG, the estimated noise or the time
    % vector in this context as a row of samples.
    % Rpeaks: R peak indices
    % fs: sampling frequency
    %
    % Outputs:
    % sigOut: blended sigIn
    % stitch: midpoint of crossfade region (samples of sigOut).

    % Definition of blanklen and blendlen parameters:
    % The blanklen parameter is the length of signal (in milliseconds) being
    % removed from around of each R peak in final ECG noise; this is the part
    % really canceled before and after each Rpeak.
    % The blendlen parameter is the length of the neighboring signal segment
    % (in milliseconds) around the canceled part to be crossfaded when
    % stitching.

    % An "if" construct is included to make blanklen and blendlen parameters
    % optional and assign them default values.
    if isempty(varargin)
        blanklen = 40e-3;
        blendlen = 60e-3;
    else
        blanklen = varargin{1}(1);
        blendlen = varargin{1}(2);
    end

    % Convert to samples:
    blanklen = blanklen * fs;
    blendlen = blendlen * fs;

    % Create linear crossfade function
    W = linspace(1, 0, blendlen + 1);

    out_locs = [Rpeaks(:)-blanklen Rpeaks(:)+blanklen];
    blend_locs = [out_locs(:,1)-blendlen out_locs(:,2)+blendlen];

    % Adjust for Rpeaks too close to end of signal
    if blend_locs(end,2) >= length(sigIn)
        blend_locs(end,:) = [];
```

```
        out_locs(end,:) = [];
        Rpeaks(end) = [];
    end

    % Initialize the stitch vector
    rpkLen = length(Rpeaks);
    stitch = zeros(1, rpkLen);

    % Start after 1st beat to remove errors related to location of 1st beat
    sigOut(1:blend_locs(1,2)) = NaN;
    for n = 2:rpkLen
        % Copy from end of last blend region to start of new blend region
        segadd = sigIn(blend_locs(n-1,2)+1:blend_locs(n,1)-1);
        sigOut(length(sigOut)+1:length(sigOut)+length(segadd)) = segadd;

        % Blend cross fade region
        segadd = sigIn(blend_locs(n,1):out_locs(n,1)).*W + ...
            sigIn(out_locs(n,2):blend_locs(n,2)).*(1-W);

        % Keep track of stitch points:
        stitch(n) = length(sigOut) + floor(length(segadd)/2);
        sigOut(length(sigOut)+1:length(sigOut)+length(segadd)) = segadd;
    end
    stitch(1) = [];

    % Add data after last Rpeak
    segadd = sigIn(blend_locs(n,2)+1:end);
    sigOut(length(sigOut)+1:length(sigOut)+length(segadd)) = segadd;

% End of function
end

function [MedBeat] = ComputeMedianBeat_long(ECG, Rpeaks, fs)
    % Compute median beat
    %
    % Inputs: ECG, R-peak locations
    % Output: Median beat, correction

    % Get short median, it is used to adjust X and Y offset
    MedShort = ComputeMedianBeat_short(ECG, Rpeaks);

    shortl = floor(length(MedShort)/3);

    % Arbitrary area for the QRS complex to do time sync. For normal patients
    % should be <120ms, in this case let's do 50 ms each side.
    arbLen = 50e-3;
    arbLenSamp = arbLen*fs;
    qrsa = min(arbLenSamp, shortl-1) ;

    % Arbitrary max lag for crosscorr search in seconds.
    maxcorrlag = 20e-3;
    maxcorrlagSamp = maxcorrlag*fs;

    % Number of extra samples in each median beat (on the left double on the
    % right) should be at least as much as the cross-fade half duration
    extrabeat = 60;
```

```
    % Since sampling frequency at development time was 1000 samples/s:
    extrabeat = ceil(extrabeat*fs/1000);

    % Determine max RR interval to set segment lengths
    RRint = diff(Rpeaks);
    maxRR = max(RRint);

    % Get left and right segments, chop out a bit from 1/3>
    segl = floor(maxRR*0.35) + extrabeat + 1;
    segr = 2*segl;

    % Initialize
    ECGmatxy = NaN(length(Rpeaks),segl + segr + 1);
    yoffset = NaN(length(Rpeaks),1);
    xoffset = yoffset ;

    m = 1;

    % Catch cases where first Rpeak is too close to start of record
    while (m < length(Rpeaks)) && (Rpeaks(m) - segl - maxcorrlagSamp < 1)
        m = m + 1;
    end
    % Main loop on each qrs peak to fill ECG beat array
    while (m < length(Rpeaks)) && ...
            (Rpeaks(m) + segr + maxcorrlagSamp < length(ECG))

        % Calculate difference from short beat on Y
        shortdiff = MedShort - ECG(Rpeaks(m)-shortl:Rpeaks(m)+2*shortl);
        yoffset(m) = mean(shortdiff);
        [crosscorr, xcorrlag] = xcorr(ECG(Rpeaks(m)-qrsa:Rpeaks(m)+qrsa), ...
            MedShort(shortl-qrsa:shortl+qrsa), maxcorrlagSamp);
        [~, maxxcorri] = max(crosscorr);
        xoffset(m) = xcorrlag(maxxcorri);
        xoff = -xoffset(m);

        % Correct on X and Y.
        ECGmatxy(m,:) = (ECG(Rpeaks(m)-segl+xoff:Rpeaks(m)+2*segl+xoff))' + …
            yoffset(m);
        % Increase loop counter
        m = m + 1;

    % End while
    end

    % Compute median from ECG beat array
    MedBeat = median(ECGmatxy, 1, 'omitnan');
% End of function
end

function [MedBeat, ECGmat] = ComputeMedianBeat_short(ECG, Rpeaks)
    % Compute median beat
    %
    % Inputs: ECG, R-peak locations
    % Output: Median beat

    % Get RRintervals
    RRint = diff(Rpeaks);
```

```
    minRR = min(RRint);

    % Get left and right segments, chop out a bit from 1/3
    segl = floor(minRR*.29);
    segr = 2*segl;

    % Remove peaks too close to the edge
    if Rpeaks(1) < segl
        Rpeaks = Rpeaks(2:end);
    end
    if length(ECG) < Rpeaks(end) + segr
        Rpeaks = Rpeaks(1:end-1);
    end

    % Initialize and fill ECG beat array
    ECGmat = zeros(length(Rpeaks), segl + segr + 1);
    for m = 1:length(Rpeaks)
        ECGmat(m,:) = (ECG(Rpeaks(m)-segl:Rpeaks(m)+segr))';
    end

    % Compute median from ECG beat array
    MedBeat = median(ECGmat,1, "omitnan");

    % Ensure it is a row
    MedBeat = MedBeat(:)';
end

function [SynECG] = GenSynECG_blend(MedBeat, Rpeaks, ecgRefLen, fs)
    % Compute synthetic ECG by blending median beats.
    %
    % Inputs: MedBeat, Rpeak location, ecgRefLen (Length of reference ECG)
    % Output: synthetic ECG

    % Cross-fade over last xx elements
    % Half of the crossfade
    crosshalfn = 50;

    % Since sampling rate at development time was 1000 sample/s:
    crosshalfn = floor(crosshalfn*fs/1000);
    medl = floor(length(MedBeat)/3);

    % Add a dummy beat at the end to fill the full space
    Rpeaks(end+1) = Rpeaks(end)+floor(mean(diff(Rpeaks)));

    % Set total out recording duration to avoid overflow
    ECGlen = max(ecgRefLen, Rpeaks(end)+2*medl+2);

    % Create linear crossfade function
    W = linspace(1, 0, 2*crosshalfn + 1);

    % Create empty output.
    SynECG = NaN(1,ECGlen);
    m = 2 ;

    % If initial Rpeak is <= medl, delete and start with second Rpeak
    if Rpeaks(1) < medl
        Rpeaks(1) = [];
```

```
    end

    while m < length(Rpeaks)
        % Get segment duration for current beat (a beat is 1 segment to the
        % left and 2 segments to the right of QRS)
        segl = floor((Rpeaks(m) - Rpeaks(m-1))*1/3);

        % Merge point is 1/3 beat back.
        mergepoint = Rpeaks(m) - segl;

        % Getting pointers for extremes of crossfade region of output
        sl = mergepoint - crosshalfn;
        sr = mergepoint + crosshalfn;

        % Pointers for xfade region of median beat
        ml = medl + 1 - segl - crosshalfn;
        mr = ml + 2*crosshalfn;
        mx = length(MedBeat) - mr;

        % Check for errors:
        if ml < 1
            disp (['ERROR! median beat too short to crossfade! add some' ...
                ' margin to median beats!!! (there is a setting in ' ...
                'compute median beat function!)'])
        end

        % Crossfade on the left
        SynECG(sl:sr) = SynECG(sl:sr).*W + MedBeat(ml:mr).*(1 - W);

        % Stamp median beat outside the crossfade region
        SynECG(sr+1: sr+mx) = MedBeat(mr+1:end);

        % Increment counter
        m = m + 1;
    % End while
    end
    SynECG = SynECG(1:ecgRefLen);
% End function
end
```