# Code to use Maximal Overlap Discrete Wavelet Transform (MODWT) to Analyze ECG Data

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Mutli-scale Integrated Remote Sensing and Simulation (MINTS)

### **Load Data**

```
load('ECG_data.mat');
```

#### **Perform Wavelet Transform**

Use the maximal overlap discrete wavelet transform (MODWT) to enhance the R peaks in the ECG waveform. The MODWT is an undecimated wavelet transform, which handles arbitrary sample sizes.

Perform MODWT for 6 different scales. The rows of wt give the wavelet (detail) coefficents for each scale. Here we are using Symlets wavelet with 4 vanishing moments

Other wavelet families can be used. use waveletfamilies('f') to view all

```
% define number of levels to use
numLevels = 6;
% perform maximal overlap discrete wavelet transform (MODWT)
wt = modwt(ECG, 'sym4', numLevels);
```

## **Compare Coefficents for all MODWT Scales**

```
% get number of Wavelet Scales
[numScales, ~] = size(wt);

% create figure
fig = figure(1);
fig.Units = 'normalized';
fig.Position = [0 0 1 1];

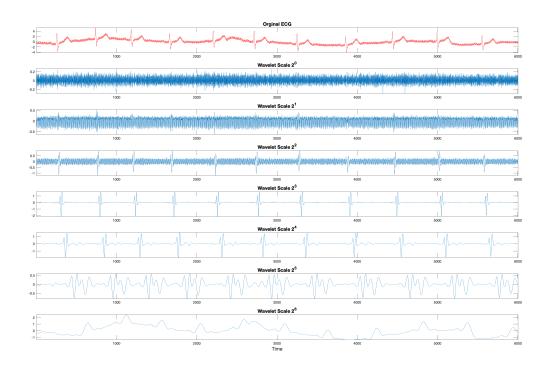
% plot original signal
subplot(numScales+1,1,1)
plot(ECG, 'r-')
title('Orginal ECG', 'FontSize', 14)
axis tight

% plot wavelet scale coefficents
for i=2:numScales+1
subplot(numScales+1,1,i)
plot(wt(i-1,:))
```

```
title(strcat('Wavelet Scale 2^', string(i-2)), 'FontSize', 14)
    axis tight
end

xlabel('Time', 'FontSize', 14)

% print figure to file
print('waveletTransform_signal_decomposition', '-dpng');
```



# **Reconstruct Signal via Inverse MODWT**

```
% create figure
fig = figure(2);
fig.Units = 'normalized';
fig.Position = [0 0 1 1];

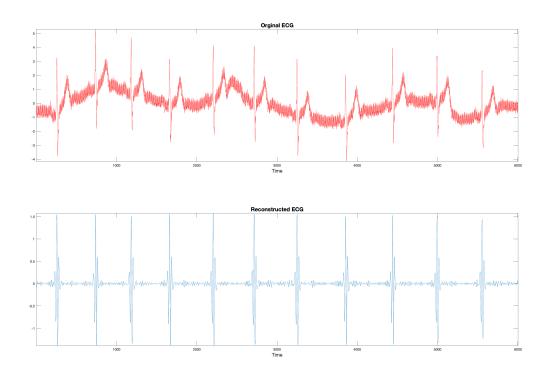
% initialize an array to store the coefficents corresponding to scale 2^4
recwt = zeros(size(wt));

% store coefficents corresponding to scale 4 in array
recwt(4,:) = wt(4,:);

% reconstruct signal using only scale 4 with inverse transform
recECG = imodwt(recwt,'sym4');

% compare original signal and reconstructed
% plot original signal
subplot(2,1,1)
plot(ECG, 'r-')
title('Orginal ECG', 'FontSize', 16)
```

```
xlabel('Time', 'FontSize', 14)
axis tight
% plot reconstructed signal
subplot(2,1,2)
plot(recECG)
title('Reconstructed ECG', 'FontSize', 16)
xlabel('Time', 'FontSize', 14)
axis tight
% print figure to file
print('waveletTransform_reconstructed_signal', '-dpng');
```



# Find R Peaks Using Reconstructed Signal

```
% define array of time indicies
t = 1:length(ECG);

% square reconstructed signal
recECG = abs(recECG).^2;
% find peaks in squared reconstructed signal
[qrspeaks,locs] = findpeaks(recECG, t, 'MinPeakHeight', 0.35, 'MinPeakDistance',100);
% create figure
fig = figure(3);
fig.Units = 'normalized';
fig.Position = [0 0 1 1];
% plot signal with detected r peaks using reconstructed signal from DWT
plot(t,ECG)
hold on
```

```
plot(locs,ECG(locs),'ro', 'LineWidth', 2)
xlabel('Time', 'FontSize', 14)
title('R Peaks Detected Using Reconstructed Signal from Wavelet Transform', ...
    'FontSize', 16)
axis tight
print('waveletTransform_rpeak_annotation', '-dpng');
```

