DSP Assignment 2 - Finite Impulse Response

Question 1 & 2

Chosen ECG file: ECG_msc_matric_5.dat renamed to ecg.dat

Number of taps was chosen to be 500, since in a theoretical system we can have a frequency resolution of 1 while in a practical scenario we must account for the transition width, therefore we must choose at least 2 times or 3 times the sampling rate given. So, we chose two times the sampling rate to give us 500 taps with a frequency resolution of 0.5.

For the bandstop design, we chose a cutoff frequency of 45Hz and 55Hz since the ideal bandstop must be designed to cover the values at 50Hz to be zero inorder to remove the 50Hz noise. Choice of cutoff frequency comes down to our chosen window function, the hanning window, which gives us good passband ripple rejection and an appreciable stopband ripple rejection. To give the window function ample room to begin its descent to 50Hz a cutoff frequency of 45 was chosen and 55hz was chosen as the second cutoff frequency to give the window function enough room to help return the impulse response back to a value of 1.

For the highpass design, a cutoff frequency of 0.5Hz was chosen since the ECG for a healthy person has a frequency of 1Hz, thus the frequency resolution must be 1Hz and DC occurs at 0Hz and therefore the cutoff must be between 0Hz and 1Hz. Additionally, we had to choose a window function that helped us achieve the best passband and stopband ripple rejection while keeping an eye on the transition width since a wide transition width near zero would not produce the best impulse response. If we were to choose the blackman window, we would achieve perfect passband and stopband ripple rejection but due to the large transition window, we would never reach the ideal impulse response value near zero as the window function would reach zero at a higher amplitude than desired whereas using no window functions we can note that we reach a more appreciable amplitude value near 0Hz while having passband and stopband ripples present. Our choice was narrowed to using the hamming or hanning window and to be consistent with our choice of window function, the hanning window was chosen.

For both the impulse response designs, we had to perform a reshuffle operation since the IDFT of the digital systems produce a non-causal system and in order to obtain a causal impulse response we had to reshuffle the coefficients, only then can we apply our chosen window functions to obtain our desired impulse responses.

We also noted that FIR doesn't perform convolution since convolution is a-causal and runs from -inf < t < +inf. FIR is casual and only has a finite number of taps in its process, therefore convolution is not suitable. However, we use a delay line system (ring buffer) to perform our

array multiplication and summation to simulate how a FIR system behaves and by feeding the delay line one sample at a time we simulated real time processing.

We used both designs and the ring buffer to plot the filtered ECG and compared it with our original ECG file, as seen in Figure 1. Note that there are 2 anomalies (2 massive spikes) in our FIR plot, these correspond to the filter starting up and are directly related to our relationship between our taps and sample frequency.

Figure 2 shows a zoomed-in heartbeat action of the FIR to show that it is PQRST intact.

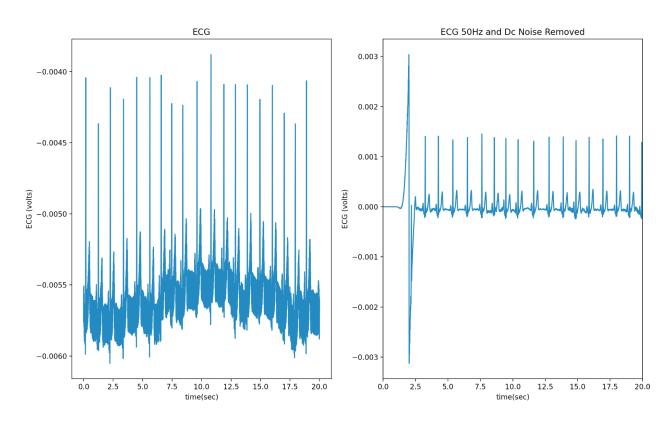


Figure 1: Plot showing the original ECG and the filter ECG

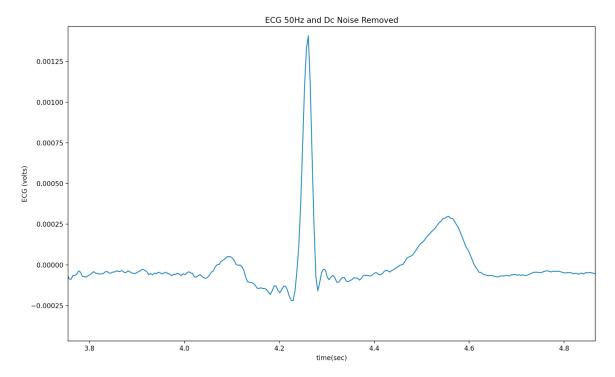


Figure 2: Shows PQRST intactness of the FIR signal

Question 3

We are tasked with removing the 50Hz noise on our ECG file using a Least Mean Square filter by emulating a sine wave at 50Hz as a noise signal and comparing our result with that of a bandstop filter at 50Hz.

The key components that define the LMS filter are the noise, learning rate ,error and the ever changing coefficients of the system. The dofilterAdapative function builds off the doFilter function, where the noise generated in the system is stored in a ring buffer so we can calculate the system output with the aid of the coefficients generated by the error function of the LMS filter. The error is calculated as the difference between our ecg data and the system output, then the coefficients of the system are updated by multiplying the error, learning rate and noise array. The error values are returned and used to plot the LMS filter.

To begin plotting the system, we must start with a very small learning rate, since we have to tweak the learning rate via trial and error before we are satisfied with the results of our filter. Figure 3 shows a direct comparison between our FIR filter with only the bandstop design applied coupled with a LMS filter at a learning rate of 0.001. We can note that both filters follow the same pattern with the bandstop filter elevated by a degree of 0.001 volts and it is shifted to

the right due to the action caused by the filter starting up whereas the LMS filter take a few milliseconds to learn, clearly seen in Figure (4), and then begins to apply the 50Hz removal.

Furthermore, Figures 5 and 6 shows the LMS filter at learning rates 0.000005 and 0.1 respectively where we can note at very low learning rates the system doesn't begin to apply the 50Hz removal as it learns at a very slow pace while at very high learning rates the system will completely diverge and fails to apply the 50Hz removal.

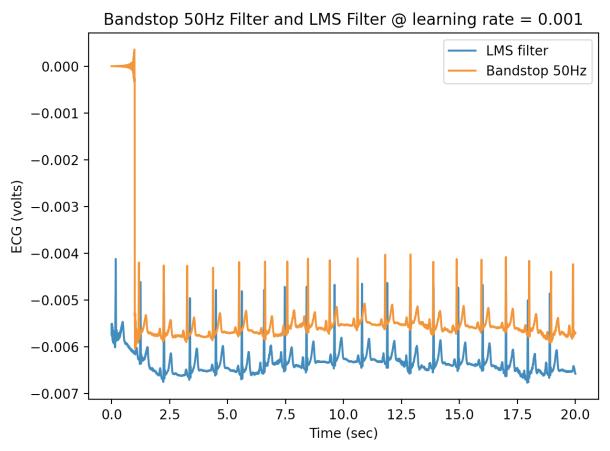


Figure 3: LMS FIlter and Bandstop 50Hz filter comparison

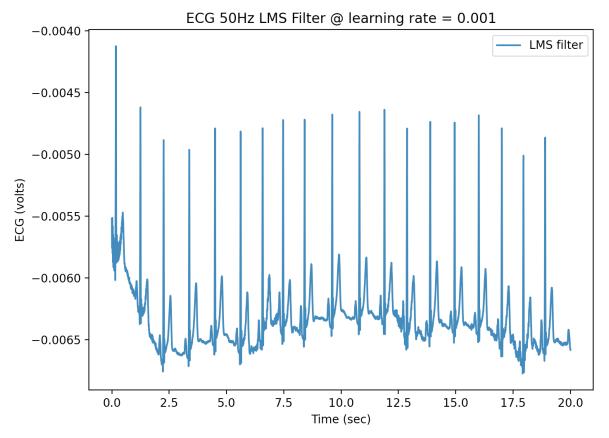


Figure 4: FIR FIlter with a learning rate of 0.001

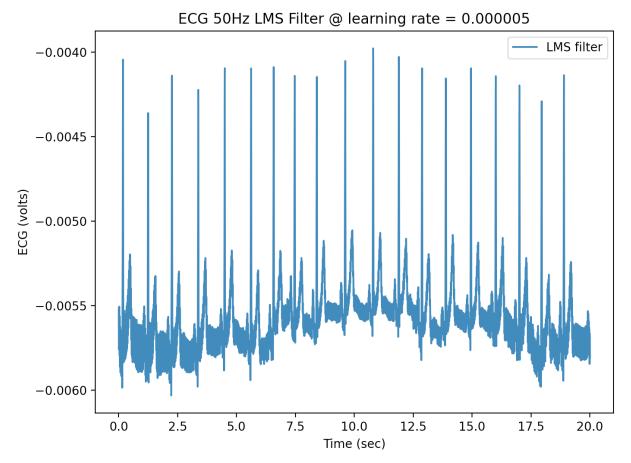


Figure 5: FIR FIlter with a learning rate of 0.000005

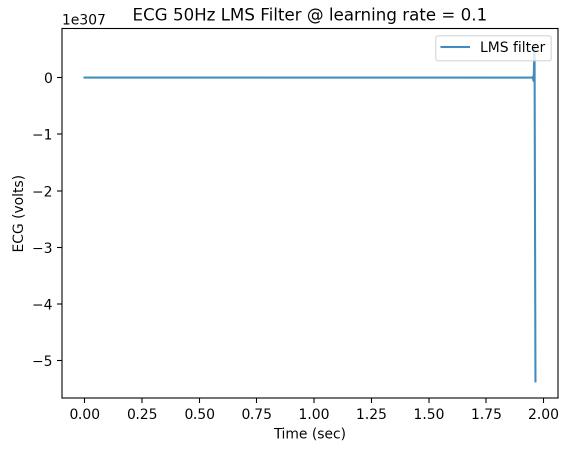


Figure 6: FIR FIlter with a learning rate of 0.1

Question 4

To perform the matched filter operation, we had to find an ECG action to be able to define our template as shown in Figure 7 and the process requires a signal that is both DC and 50Hz free so we used our previously found FIR signal as our input signal. Once our template was found, we time reversed it to create our coefficients in order to perform correlation on our FIR filter. Next, we used a sinc function (wavelet) to closely model our time reversed coefficients as seen in Figure 8. Once our wavelet was modelled, we used the ring buffer with our new wavelet and the previously found FIR signal to obtain a new FIR signal that was influenced by the sinc wavelet in order to improve our peak detection and Figure 9 shows both the original and the new FIR signals for comparison.

The wavelet coefficients were raised to a power of 5 in order to enhance our peak detection process as we were able to show that the difference between the highest and lowest peaks of the new FIR plot is far greater than that of the original filter. The difference between the peaks in the original FIR is 0.1397 - 0.000230 = 0.13947 while in the new FIR we have a peak difference of $(2.011*10^{-17} - 3.67*10^{-18})/10^{-17} = 1.644$ which is far greater than the difference in the original plot proving that our peak detection is working.

Next, we performed a threshold on our new FIR plot to obtain the R-peaks shown in Figure 10, so that we can remove the anomalies created by the filter starting up and only show the distinct peaks of the ECG. With the R-peaks plot, we can now define the the momentary heart beat for our FIR filter by calculating the time difference between two peaks using the following formulae:

Beats per minute = 60/(peak_time_2 - peak_time_1)

Momentary heart rate plot for both the original FIR and the wavelet influenced FIR are plotted in Figure 11.

Analysing Figures 9 & 11, we can observe that both waveforms closely follow one another, however there exists a group delay of 0.5s between the original FIR signal and the wavelet influenced FIR signal which means there was deviation from the linear phase when we applied the wavelet to our original FIR signal resulting in phase distortion. It can be concluded that using matched filter for peak detection will produce a similar plot to that of the original signal used with enhanced peak values but will carry a group delay.

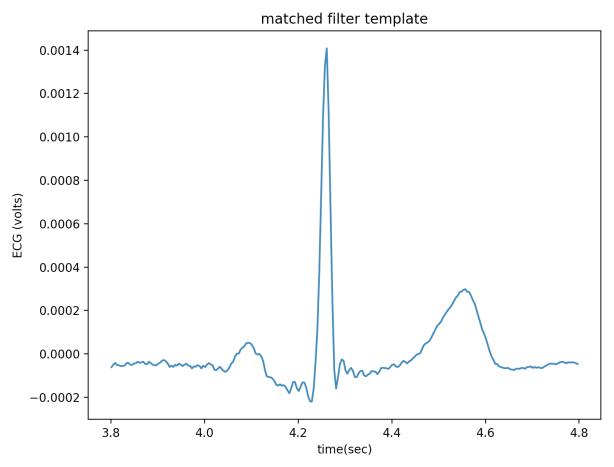


Figure 7: Matched filter template

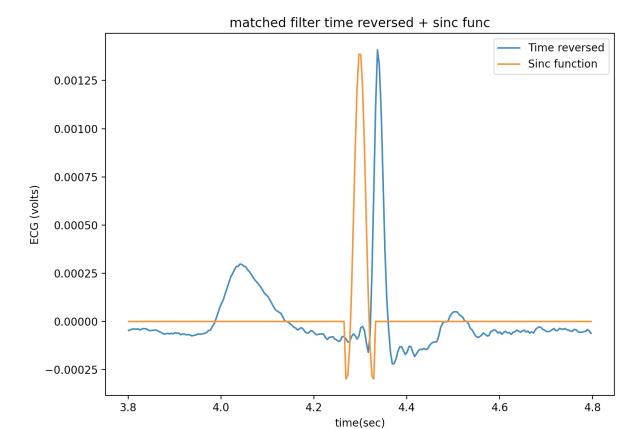


Figure 8: Time reversed template and Sinc function

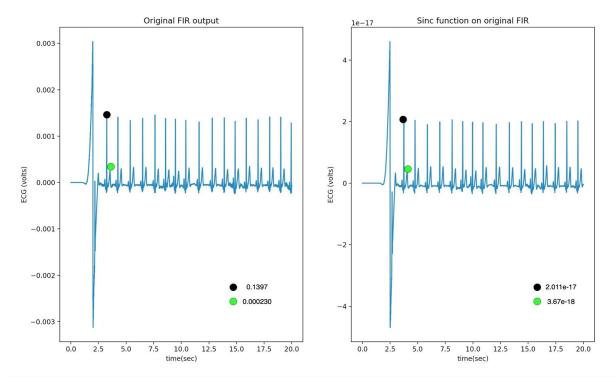


Figure 9: Comparison between the original FIR and the new FIR signal

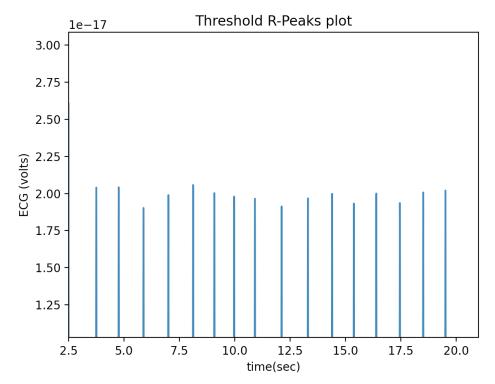


Figure 10: R-peak plot for the new FIR signal

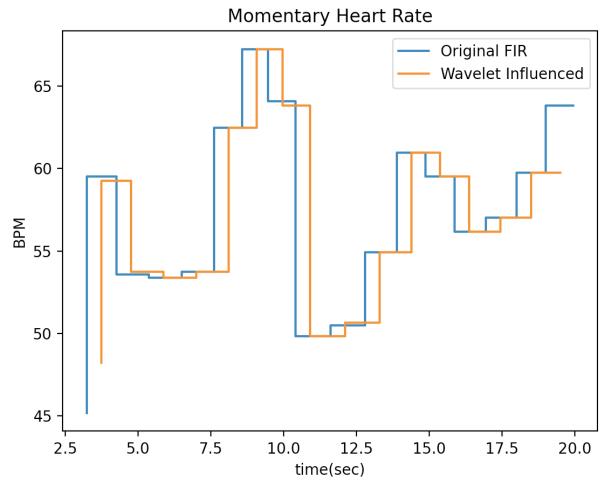


Figure 11: Momentary heart rate for both the original FIR and the new FIR signals

Appendix

hpbsfilter.py:

```
import numpy as np
import matplotlib.pyplot as plt
import firfilter
"""Reshuffle Function"""
def reshuffle(filter coeff):
  h = np.zeros(taps) # create an array to hold the impulse response values
  h[0:int(taps / 2)] = filter coeff[int(taps / 2):taps] # perform a
reshuffling action
  h[int(taps / 2):taps] = filter coeff[0:int(taps / 2)] # perform a
reshuffling action
  return h * np.hanning(taps) # return the impulse response with a window
function applied to it
"""50 HZ removal"""
def bandstopDesign(samplerate, w1, w2, itr):
   # frequency resolution =0.5
  M = samplerate * itr # calculate the taps
  X = np.ones(M) # create an array of ones to model an ideal bandstop
  X[w1:w2 + 1] = 0 # mirror 1 (set all values to 0)
  X[M - w2:M - w1 + 1] = 0 # mirror 2 (set all values to 0)
  x = np.real(np.fft.ifft(X)) # perform IDFT to obtain an a-causal system
  return x
"""DC noise removal"""
def highpassDesign(samplerate, w3, itr):
  # frequency resolution =0.5
  M = samplerate * itr # calculate the taps
  X = np.ones(M) # create an array of ones to model an ideal highpass
  X[0:w3 + 1] = 0 # mirror 1 (set all values to 0)
  X[M - w3: M + 1] = 0 \# mirror 2 (set all values to 0)
  x = np.real(np.fft.ifft(X)) # perform IDFT to obtain an a-causal system
  return x
```

```
# Q1 and Q2
"""Load data into python"""
data = np.loadtxt('ecg.dat')
"""Define constants"""
fs = 250 # sample frequency
t max = len(data) / fs # sample time of data
t data = np.linspace(0, t max, len(data)) # create an array to model the
x-axis with time values
practical = 2 # define by how much the taps are greater than the sampling rate
to account for transition width
taps = (fs * practical) # defining taps
"""Bandstop"""
f1 = int((45 / fs) * taps) # cutoff frequency before 50Hz
f2 = int((55 / fs) * taps) # cutoff frequency after 50Hz
"""Highpass"""
f3 = int((0.5 / fs) * taps) # ideal for cutting off DC noise
"""Call the function for Bandstop and Highpass"""
impulse BS = bandstopDesign(fs, f1, f2, practical)
impulse HP = highpassDesign(fs, f3, practical)
"""Reshuffle the time reversed coeff for highpass by calling reshuffle
function"""
h newHP = reshuffle(impulse HP)
"""Reshuffle the time reversed coeff for bandstop by calling reshuffle
function"""
h newBS = reshuffle(impulse BS)
"""Call the class method dofilter, by passing in only a scalar value at a time
which outputs a scalar value"""
# obtain FIR HP output when we couple the original ECG data with the highpass
over a ring buffer
fir HP = np.empty(len(data))
fi = firfilter.firFilter(h newHP)
for i in range(len(fir HP)):
   fir HP[i] = fi.dofilter(data[i])
# obtain FIR output when we couple the previously found FIR HP data with the
bandstop over a ring buffer
fir = np.empty(len(data))
po = firfilter.firFilter(h newBS)
for i in range(len(fir)):
  fir[i] = po.dofilter(fir HP[i])
```

```
"""Plot both the original ECG data set and new filtered data set """
plt.figure(1)
plt.subplot(1, 2, 1)
plt.plot(t_data, data)
plt.title('ECG')
plt.xlabel('time(sec)')
plt.ylabel('ECG (volts)')

plt.subplot(1, 2, 2)
plt.plot(t_data, fir)
plt.xlim(0, t_max)
plt.title('ECG 50Hz and Dc Noise Removed')
plt.xlabel('time(sec)')
plt.ylabel('ECG (volts)')
```

firfilter.py:

```
import numpy as np
class firFilter:
  def init (self, data):
      self.coeff = _data
      self.ntaps = len( data)
      self.buffer = np.zeros(self.ntaps)
      self.s offset = 0
  def dofilter(self, v):
       # ring buffer
      self.buffer[self.s offset % self.ntaps] = v
      output = 0
      for i in range(self.ntaps):
          output += self.buffer[(i + self.s offset) % self.ntaps] *
self.coeff[i]
      self.s offset += 1
      return output
  def dofilterAdaptive(self, signal, noise, learningRate):
      # ring buffer
      self.buffer[self.s offset % self.ntaps] = noise
```

```
output = 0
       for i in range(self.ntaps):
          output += self.buffer[(i + self.s offset) % self.ntaps] *
self.coeff[i]
       # update coefficients
       error = signal - output
       for k in range(self.ntaps):
           self.coeff[k] += error * learningRate * self.buffer[(k +
self.s offset) % self.ntaps]
       self.s offset += 1
       return error
Imsfilter.py:
import numpy as np
import matplotlib.pyplot as plt
import firfilter
"""Reshuffle Function"""
def reshuffle(filter coeff):
  h = np.zeros(taps) # create an array to hold the impulse response values
  h[0:int(taps / 2)] = filter coeff[int(taps / 2):taps] # perform a
reshuffling action
  h[int(taps / 2):taps] = filter coeff[0:int(taps / 2)] # perform a
reshuffling action
   return h * np.hanning(taps) # return the impulse response with a window
function applied to it
"""50 HZ removal"""
def bandstopDesign(samplerate, w1, w2, itr):
  # frequency resolution =0.5
  M = samplerate * itr # calculate the taps
  X = np.ones(M) # create an array of ones to model an ideal bandstop
  X[w1:w2 + 1] = 0 \# mirror 1 (set all values to 0)
  X[M - w2:M - w1 + 1] = 0 # mirror 2 (set all values to 0)
  x = np.real(np.fft.ifft(X)) # perform IDFT to obtain an a-causal system
  return x
```

"""Load data into python"""

```
data = np.loadtxt('ecg.dat')
"""Define constants"""
fs = 250 # sample frequency
t max = len(data) / fs # sample time of data
t data = np.linspace(0, t max, len(data)) # create an array to model the
x-axis with time values
practical = 2 # define by how much the taps are greater than the sampling rate
to account for transition width
taps = (fs * practical) # defining taps
f sine = 50 # noise signal frequency
1R = 0.001 # learning rate of the LMS filter
"""Bandstop"""
f1 = int((45 / fs) * taps) # cutoff frequency before 50Hz
f2 = int((55 / fs) * taps) # cutoff frequency after 50Hz
"""Call the function for Bandstop and Highpass"""
impulse BS = bandstopDesign(fs, f1, f2, practical)
"""Reshuffle the time reversed coeff for bandstop by calling reshuffle
function"""
h newBS = reshuffle(impulse BS)
"""Call the class method dofilter, where we only perform 50Hz removal to
compare with our LMS filter"""
fir BS = np.empty(len(data))
fi = firfilter.firFilter(h newBS)
for i in range(len(fir BS)):
  fir BS[i] = fi.dofilter(data[i])
# 03
lms = np.empty(len(data)) # create an empty array to store LMS filter results
time = np.linspace(0, t max, len(lms)) # create an array to model the x-axis
with time values
"""Call the dofilterAdaptive function from the class to compute the FIR
dataset"""
f = firfilter.firFilter(np.zeros(taps))
for i in range(len(data)):
   sinusoid = (np.sin(2 * np.pi * i * (f sine / fs)))
   lms[i] = f.dofilterAdaptive(data[i], sinusoid, lR)
"""Plot the LMS filter"""
plt.figure(1)
plt.plot(time, lms, label='LMS filter')
plt.plot(time, fir BS, label='Bandstop 50Hz')
plt.title('Bandstop 50Hz Filter and LMS Filter @ learning rate = ' + str(lR))
```

```
plt.xlabel('Time (sec)')
plt.ylabel('ECG (volts)')
plt.legend(loc='upper right')

plt.figure(2)
plt.plot(time, lms, label='LMS filter')
plt.title('LMS Filter @ learning rate = ' + str(lR))
plt.xlabel('Time (sec)')
plt.ylabel('ECG (volts)')
plt.legend(loc='upper right')
```

hrdetect.py:

```
import numpy as np
import matplotlib.pyplot as plt
import firfilter
"""Reshuffle Function"""
def reshuffle(filter coeff):
  h = np.zeros(taps) # create an array to hold the impulse response values
  h[0:int(taps / 2)] = filter coeff[int(taps / 2):taps] # perform a
reshuffling action
  h[int(taps / 2):taps] = filter_coeff[0:int(taps / 2)] # perform a
reshuffling action
   return h * np.hanning(taps) # return the impulse response with a window
function applied to it
"""Pull out one ECG action"""
def get ecgaction(dataset, dataset time, start, stop):
  data range = dataset[start:stop] # define the data points of interest
   time_range = dataset_time[start:stop] # define the time range in which
those data points occur
  return data range, time range # return both data and time arrays of the ecg
action
"""Create a sinc wavelet"""
```

```
def get wavelet(length, time reversed dataset, time range):
  data val = max(time reversed dataset) * np.sinc(length * 25) # define the
characteristics of the sinc wavelet
  data val[0:int(len(time range) / 2) - 8] = 0 # zero out all values of no
interest to
   # the left of the function's peak
  data val[int(len(time range) / 2) + 8:len(time range)] = 0 # zero out all
values of no interest to
   # the right of the function's peak
  return data val # return the wavelet data array
"""R Peak Threshold Function"""
def threshold(dataset):
  val = max(dataset[700:]) # 700 was picked as we want to avoid the anomalies
caused by the filter starting up
  highest volt = val + val / 2 # Dynamically set the max of the threshold
  lowest volt = val * 0.5 # Dynamically set the min of the threshold
  return lowest volt, highest volt # return the max and min threshold of the
dataset
"""Generate a list to store peak times"""
def get peaktime(dataset, upper, lower, r):
  data points = [] # create a list to store data points
  iter val = 0
  while iter val < (len(dataset)):</pre>
      if upper > dataset[iter val] > lower: # set the condition for data
storage
          data points.append(r[iter val]) # store data points
          iter val += 50 # we add 50 data points once we find our max point
to avoid the next closest peak from
          # overwriting our peak value
          iter val += 1 # we add 1 data point to move to the next iteration
value
   return data points # return the list of data points
"""Generate a list to store bpm fir wavelet values"""
```

```
def get bpm(dataset):
  data points = [] # create a list to store data points
   for iter val in range(len(dataset) - 1):
      data points.append(60 / (dataset[iter val + 1] - dataset[iter val])) #
perform the bpm calculation
   return data points # return the list of data points
"""50 HZ removal"""
def bandstopDesign(samplerate, w1, w2, itr):
  # frequency resolution =0.5
  M = samplerate * itr # calculate the taps
  X = np.ones(M) # create an array of ones to model an ideal bandstop
  X[w1:w2 + 1] = 0 # mirror 1 (set all values to 0)
  X[M - w2:M - w1 + 1] = 0 \# mirror 2 (set all values to 0)
  x = np.real(np.fft.ifft(X)) # perform IDFT to obtain an a-causal system
  return x
"""DC noise removal"""
def highpassDesign(samplerate, w3, itr):
  # frequency resolution =0.5
  M = samplerate * itr # calculate the taps
  X = np.ones(M) # create an array of ones to model an ideal highpass
  X[0:w3 + 1] = 0 \# mirror 1 (set all values to 0)
  X[M - w3: M + 1] = 0 \# mirror 2 (set all values to 0)
  x = np.real(np.fft.ifft(X)) # perform IDFT to obtain an a-causal system
  return x
"""Load data into python"""
data = np.loadtxt('ecg.dat')
"""Define constants"""
fs = 250 # sample frequency
t max = len(data) / fs # sample time of data
t data = np.linspace(0, t max, len(data)) # create an array to model the
x-axis with time values
practical = 2 # define by how much the taps are greater than the sampling rate
to account for transition width
taps = (fs * practical) # defining taps
```

```
"""Bandstop"""
f1 = int((45 / fs) * taps) # cutoff frequency before 50Hz
f2 = int((55 / fs) * taps) # cutoff frequency after 50Hz
"""Highpass"""
f3 = int((0.5 / fs) * taps) # ideal for cutting off DC noise
"""Call the function for Bandstop and Highpass"""
impulse BS = bandstopDesign(fs, f1, f2, practical)
impulse HP = highpassDesign(fs, f3, practical)
"""Reshuffle the time reversed coeff for highpass by calling reshuffle
function"""
h newHP = reshuffle(impulse HP)
"""Reshuffle the time reversed coeff for bandstop by calling reshuffle
function"""
h newBS = reshuffle(impulse BS)
"""Call the class method dofilter, by passing in only a scalar value at a time
which outputs a scalar value"""
# obtain FIR HP output when we couple the original ECG data with the highpass
over a ring buffer
fir HP = np.empty(len(data))
fi = firfilter.firFilter(h newHP)
for i in range(len(fir HP)):
   fir HP[i] = fi.dofilter(data[i])
# obtain FIR output when we couple the previously found FIR HP data with the
bandstop over a ring buffer
fir = np.empty(len(data))
po = firfilter.firFilter(h newBS)
for i in range(len(fir)):
  fir[i] = po.dofilter(fir HP[i])
# 04
"""Find the range in the FIR plot where an ECG action occurs and plot it"""
plt.figure(1)
plt.subplot(1, 2, 1)
template, ecgaction time = get ecgaction(fir, t data, 950, 1200) # call the
function to pull out one ecg action
plt.plot(ecgaction time, template)
plt.title("matched filter template")
plt.xlabel('time(sec)')
plt.ylabel('ECG (volts)')
```

```
"""Plot the time reversed version of the template """
plt.subplot(1, 2, 2)
time reversed coeff = template[::-1] # time reverse the template to obtain
desired coefficient values
plt.plot(ecgaction time, time reversed coeff, label='Time reversed')
plt.xlabel('time(sec)')
plt.ylabel('ECG (volts)')
"""Create and plot the sinc function"""
n coeff = get wavelet(np.linspace(-1, 1, len(ecgaction time)),
time reversed coeff, ecgaction time)
plt.subplot(1, 2, 2)
plt.plot(ecgaction time, n coeff, label='Sinc function')
plt.legend(loc='upper right')
n coeff = n coeff ** 5 \# Raised to the power of 5 to show the significant
difference between the highest peak and the
# smallest peak
""""Call the dofilter function in the FIR class with the wavelet data
and the filtered FIR data set to find the new fir data set influenced by a
wavelet"""
fir wavelet = np.empty(len(data))
fi = firfilter.firFilter(n_coeff)
for i in range(len(fir wavelet)):
  fir wavelet[i] = fi.dofilter(fir[i])
"""Plot both the original FIR and the new FIR"""
ecg time = np.linspace(0, t max, len(fir))
plt.figure(2)
plt.subplot(1, 2, 1)
plt.plot(ecg time, fir)
plt.xlabel('time(sec)')
plt.ylabel('ECG (volts)')
plt.title('Original FIR output')
plt.subplot(1, 2, 2)
plt.plot(ecg time, fir wavelet)
plt.xlabel('time(sec)')
plt.ylabel('ECG (volts)')
plt.title('Sinc function on original FIR')
"""Define and plot the R peak threshold """
```

```
plt.figure(3)
min thresh, max thresh = threshold(fir wavelet)
plt.plot(ecg time, fir wavelet)
plt.xlim(2.5) \# Limit the x-axis to start from 2.5 since we don't data want
the range of values at which our filter
plt.ylim(min thresh, max thresh) # Limit the y-axis between max threshold and
min threshold values
plt.xlabel('time(sec)')
plt.ylabel('ECG (volts)')
plt.title('Threshold R-Peaks plot')
"""Call the get peak time function to create a list of peak times for the
wavelet influenced FIR"""
peak time fir wavelet = get peaktime(fir wavelet, max thresh, min thresh,
ecg time)
"""Call the get bpm function to create a list of bpm values for the wavelet
influenced FIR"""
bpm fir wavelet = get bpm(peak time fir wavelet)
"""Define the original FIR R peak threshold """
min FIR thresh, max FIR thresh = threshold(fir)
"""Call the get peak time function to create a list of peak times for the
original FIR"""
peak time fir = get peaktime(fir, max FIR thresh, min FIR thresh, ecg time)
"""Call the get bpm function to create a list of bpm values for the original
FTR"""
bpm fir = get bpm(peak time fir)
"""Plot the momentary heart rate for the original fir and the fir wavelet"""
plt.figure(4)
plt.step(peak time fir[2:], bpm fir[1:], label="Original FIR")
plt.step(peak time fir wavelet[2:], bpm fir wavelet[1:], label="Wavelet
Influenced")
plt.xlabel('time(sec)')
plt.ylabel('BPM')
plt.title('Momentary Heart Rate')
plt.legend(loc="upper right")
plt.show()
```