
MyoMex Quickstart

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Before you begin, please read through README.txt and follow all steps for setting up the Myo Connect application, Myo SDK, and building the MEX function myo_mex.

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Before Using MyoMex

If you decided not to read through README.txt, let's at least show you the quickest possible way to get started.

```
install_myo_mex; % adds directories to MATLAB search path
% install_myo_mex save % additionally saves the path
```

```
sdk_path = 'C:\myo-sdk-win-0.9.0'; % root path to Myo SDK
build_myo_mex(sdk_path); % builds myo_mex
```

Evaluating mex command:

```
'mex -O -I"C:\myo-sdk-win-0.9.0\include" -L"C:\myo-sdk-win-0.9.0\lib" -lm
```

MEX-file 'myo_mex' built successfully!

MyoMex Usage

Before using MyoMex, you must decide how many Myos you'd like to use in this session. The MyoMex constructor argument countMyos specifies this value. Make sure that exactly countMyos devices are connected in the Myo Connect application or else MyoMex construction will fail. The Myo device(s) should also be worn on a human arm to avoid unexpected disconnection of the device from Myo Connect. If Myo Connect loses a device at any time, MyoMex will terminate and invalidate itself.

```
countMyos = 1;
```

Instantiate MyoMex

After constructing a new MyoMex instance, we'll inspect its properties.

```
mm = MyoMex(countMyos)
```

Warning: The class file for 'MyoData' has been changed, but the change can

applied because objects based on the old class file still exist. If you use those objects, you might get unexpected results. You can use the 'clear' command to remove those objects. See 'help clear' for information on how to remove objects.

```
mm =
```

```
MyoMex with properties:
```

```
myoData: [1x1 MyoData]
```

Notice that the only property of mm is a 1xcountMyos MyoData object. There is no device data stored by mm. The data from each physical Myo device is passed through mm to each element of mm.myoData.

Inspect MyoData

Since MyoData objects inherit from handle, we can get handles to the MyoData objects representing each physical device and use them directly.

```
m1 = mm.myoData(1);  
if countMyos == 2, m2 = mm.myoData(2); end
```

Now, we'll just continue this exercise with m1, but the exact same demonstration applies for m2 as well (if countMyos == 2).

The most recent data from Myo will be stored in the relevant properties of the MyoData object (i.e. quat, gyro, accel, emg, pose, etc.). The following is a list of all properties in MyoData.

```
pause(0.1); % wait briefly for the first data frame to come in
```

```
% data properties sampled on the IMU time base
```

```
m1.timeIMU  
m1.quat  
m1.rot  
m1.gyro  
m1.gyro_fixed  
m1.accel  
m1.accel_fixed
```

```
% data properties sampled on the EMG time base
```

```
m1.timeEMG  
m1.emg  
m1.pose  
m1.pose_rest  
m1.pose_fist  
m1.pose_wave_in  
m1.pose_wave_out  
m1.pose_fingers_spread  
m1.pose_double_tap  
m1.arm  
m1.arm_right  
m1.arm_left  
m1.arm_unknown
```

```
m1.xDir
m1.xDir_wrist
m1.xDir_elbow
m1.xDir_unknown
```

```
ans =
```

```
0.1330
```

```
ans =
```

```
-0.7107 -0.4577 -0.3240 -0.4248
```

```
ans =
```

```
0.4290 -0.3072 0.8494
0.9004 0.2201 -0.3752
-0.0716 0.9258 0.3711
```

```
ans =
```

```
35.7500 14.1875 23.8125
```

```
ans =
```

```
31.2058 26.3787 19.4100
```

```
ans =
```

```
-0.0718 0.7300 0.3325
```

```
ans =
```

```
0.0274 -0.0288 0.8044
```

```
ans =
```

```
0.1730
```

```
ans =
```

```
Columns 1 through 7
```

```
-0.0625 -0.1406 -0.2422 0 0.0156 0.2656 -0.1953
```

Column 8

-0.1641

ans =

65535

ans =

0

ans =

0

ans =

0

ans =

0

ans =

0

ans =

0

ans =

2

ans =

0

ans =

0

```
ans =  
  
1  
  
ans =  
  
2  
  
ans =  
  
0  
  
ans =  
  
0  
  
ans =  
  
1
```

Using Logged Data

As MyoData receives data from MyoMex, it's automatically accumulated in so-called `data_log` properties, i.e. `quat_log`, `accel_log`, etc. We refer to this as the streaming mode of a MyoData object. This status is indicated by the `isStreaming` property.

```
ml.isStreaming
```

```
ans =  
  
1
```

We can inspect the accumulation of the logs for example,

```
fprintf('%10s%10s\n','time','samples')  
for ii = 1:5  
    fprintf('% 8.2f%10d\n',...  
        ml.timeIMU,size(ml.quat_log,1));  
    pause(0.2);  
end  
fprintf('\n\n');
```

<i>time</i>	<i>samples</i>
0.17	9
0.37	19
0.57	29

```
0.77      39
0.97      49
```

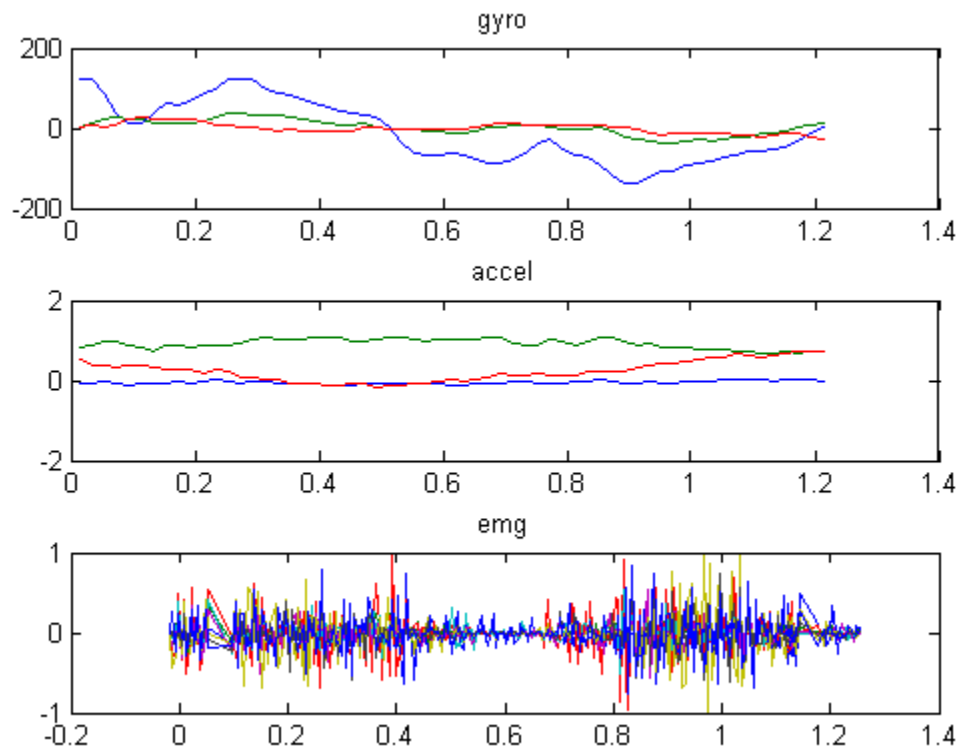
Although we can't stop the data from being passed to in MyoData, we can toggle streaming mode by using the methods `stopStreaming()` and `startStreaming()`.

```
m1.stopStreaming();
fprintf('Number of samples:           \t%d\n',length(m1.timeIMU_log));
pause(1);
fprintf('Number of samples after pause(1):\t%d\n',length(m1.timeIMU_log));
```

```
Number of samples:           61
Number of samples after pause(1): 61
```

Now we can plot some data taking care to use the correct time vectors.

```
figure;
subplot(3,1,1); plot(m1.timeIMU_log,m1.gyro_log); title('gyro');
subplot(3,1,2); plot(m1.timeIMU_log,m1.accel_log); title('accel');
subplot(3,1,3); plot(m1.timeEMG_log,m1.emg_log); title('emg');
```



If you'd like to clear the `data_log` properties to start a new logging trial, then you may use the `clear-Logs()` method,

```
% collect about T seconds of data
T = 5; % seconds
```

```
m1.clearLogs()
m1.startStreaming();
pause(T);
m1.stopStreaming();
fprintf('Logged data for %d seconds,\n\t',T);
fprintf('IMU samples: %10d\tApprox. IMU sample rate: %5.2f\n\t',...
    length(m1.timeIMU_log),length(m1.timeIMU_log)/T);
fprintf('EMG samples: %10d\tApprox. EMG sample rate: %5.2f\n\t',...
    length(m1.timeEMG_log),length(m1.timeEMG_log)/T);

    Logged data for 5 seconds,
        IMU samples:          251 Approx. IMU sample rate: 50.20
        EMG samples:          1000 Approx. EMG sample rate: 200.00
```

Finally, when you're done with MyoMex, don't forget to clean up!

```
mm.delete;
clear mm
```

Finally, take advantages of the following resources for additional information about MyoData!

```
% MyoMexGUI_Monitor
% properties MyoData
% methods MyoData
% help MyoData
```

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