

Bat2Matlab User Manual

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1 Introduction

Batlab is a custom built software program which provides a common interface for executing intracranial micro-electrode recording (MER) experiments exploring neural auditory pathways. This functionality includes presenting a wide variety of audio stimulus, storing the resulting MERs, and analyzing aspects of this data. Bat2Matlab is a set of Matlab tools for converting the Batlab data format into Matlab data structures. In addition, Bat2Matlab provides tools for visualization, analysis, and (in the near future) modeling of this data.

This document will cover:

- Batlab experiment structure
- Batlab data formats
- Importing audio data
- Importing Batlab XML
- Importing MER data
- Calculating spike times
- Generating histograms
- Visualizing the data
- Example

2 Batlab experiment structure

Experiments in Batlab have a hierarchical structure. An experiment consists of any number of tests. A test is usually performed on a single cell at a single depth. Often, tests are automated in Batlab and may include, for example, all of the stimulus presentations and MERs required to produce a frequency tuning for the cell. A test consists of any number of traces. Each trace includes multiple presentations of an identical stimulus and the recordings of the responses.

3 Batlab data formats

Batlab inputs and outputs a number of different audio formats. These include three obscure formats: kanwal (*.kanwal), call (*.call), and batcall (*.wav) formats as well as standard wave (*.wav) files. Data recorded during an experiment is saved into two file types. One is the metadata file (*.pst) and the other is the raw data file (*.raw). Each of these contains the data for each repetition of each trace of each test of the experiment. A separate program, Batgor¹, is used to export the *.pst and *.raw files to formats that can be imported by other analysis software packages (originally, it was intended to export to a format readable by Igor, hence the name). Bat2Matlab makes use of Batgor's XML output format. An XML schema for the Batgor XML format is included in Appendix A.

4 Importing audio data

A single Matlab interface is used for importing audio data in any of the formats listed above:

```
function [signal sample_rate] = ParseAudioData(file_path)
```

INPUT ARGUMENTS

file_path Path to the audio file.

OUTPUT ARGUMENTS

signal Audio signal, normalized to 1.

sample_rate The sample rate of the audio signal

This function automatically determines which type of audio file is being input (including discriminating between both *.wav formats).

5 Importing Batlab XML

```
function batlab_struct = BatlabXml2MatlabStructure(file_path)
```

INPUT ARGUMENTS

file_path Path to the Batlab XML file.

OUTPUT ARGUMENTS

batlab_struct Batlab metadata, converted to a Matlab structure

The resulting Matlab structure has the same hierarchical structure as the Batlab XML file. See Appendix B for the field names and organization of this structure.

6 Importing MER data

```
function test_data = ExtractRawData(experiment_data,  
                                     raw_data_filepath,  
                                     test_nums)
```

INPUT ARGUMENTS

experiment_data bat2matlab structure extracted from batlab XML file.

raw_data_filepath file path to the batlab *.raw file associated with

¹Much thanks to Ed Groth for adding the XML export functionality to Batgor that made Bat2Matlab possible.

	the batlab XML file.
test_nums	The number of the tests to extract data for.
OUTPUT ARGUMENTS	
experiment_data	bat2matlab structure with requested raw data included.

test_nums is a vector of numbers indicating which tests to extract the MER data for. Since the Matlab structure can grow very large, it is advised to only extract the test data needed for analysis.

7 Calculating spike times

```
function experiment_data = CalculateSpikeTimes(experiment_data,
                                              test_nums,
                                              peak_threshold,
                                              filter_cutoff)
```

INPUT ARGUMENTS	
experiment_data	bat2matlab structure
test_nums	The numbers of the tests to calculate spike times for.
peak_threshold	Theshold level used on the smoothed power of the MER signal for spike detection. Default: 0.2
filter_cutoff	Filter cutoff frequency (in Hz) used for smoothing the power of the signal for spike detection. Default: 1200
OUTPUT ARGUMENTS	
experiment_data	bat2matlab structure with requested spike times included

The default values should be suitable for most Batlab data. It is recommended, however, to perform a visual inspection of the spike calculations using the visualization tools (see below). The visualization tools make it easy to find appropriate values of `peak_threshold`, `filter_cutoff`, and `power_exponent`. In most cases, adjusting `peak_threshold` will suffice. If, however, if the MER signal is too smooth or not smooth enough, `filter_cutoff` can be increased or decreased, respectively.

8 Generating histograms

```
function experiment_data =
    GenerateHistograms(experiment_data,bin_width,test_nums)
```

INPUT ARGUMENTS	
experiment_data	bat2matlab structure
test_nums	The numbers of the tests to calculate spike times for.
bin_width	The width of the histogram bins (in milliseconds) Default = 1
OUTPUT ARGUMENTS	
experiment_data	bat2matlab structure with requested histograms included

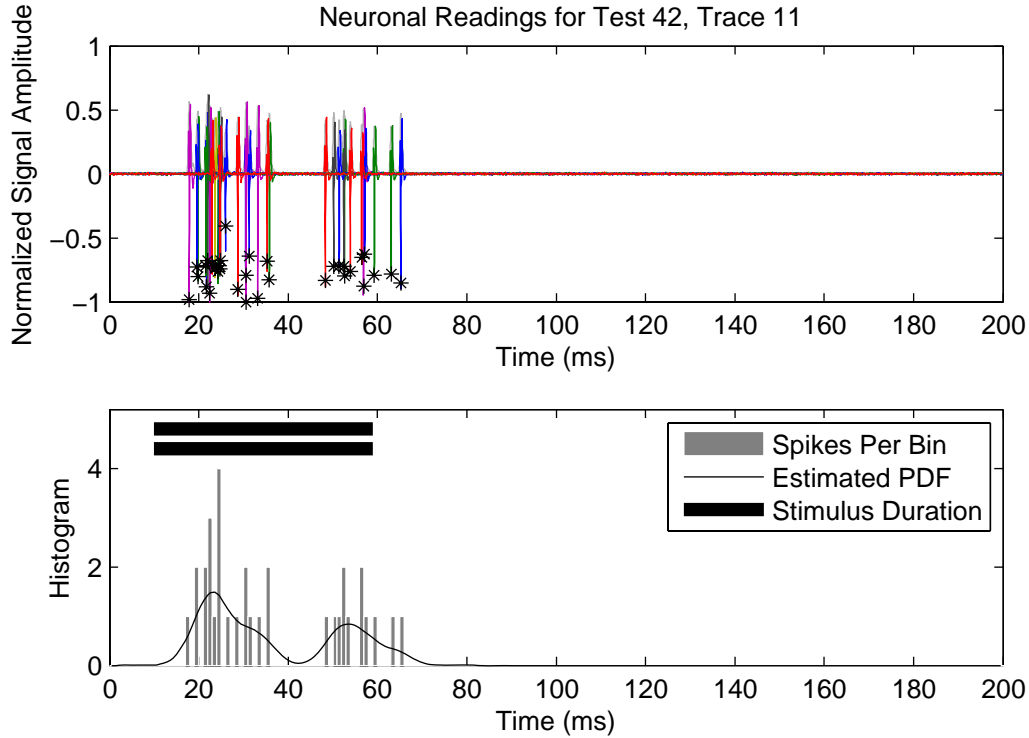


Figure 1: A typical plot using all of the plotting options

9 Visualizing the data

There is a single interface for visualizing the data contained in the Bat2Matlab structure:

```
function PlotData(experiment_data, test_num, trace_num, plot_flag)

INPUT ARGUMENTS
experiment_data    bat2matlab structure
test_num           The number of the test to display
trace_num          The trace of test test_num to display
plot_flag          A binary vector indicating the plotting features to
                    display
                    Default: [1 1 1 1 1 1]

    flag index 1    Plot the raw MER signals
    flag index 2    Plot the smoothed MER power signals (used for spike
                    detection)
    flag index 3    Plot the detected spikes
    flag index 4    Plot the spike histogram
    flag index 5    Plot the estimated spike histogram PDF (normalized
                    to the integrated histogram for viewing purposes)
    flag index 6    Plot the stimulus
```

By default, all of the viewing options are "on" (`plot_flag = [1 1 1 1 1 1]`). A typical output is shown in Figure 1.

The top plot shows the MERs. Each sweep of the trace is shown in a different color. The asterisks indicate the spikes that were calculated. They appear at the points of maximum signal

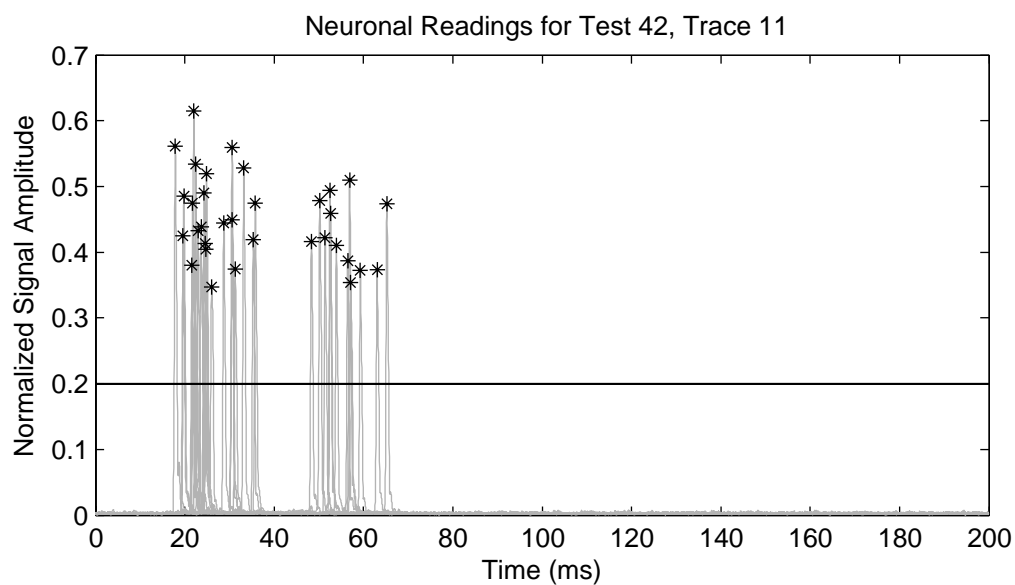


Figure 2: A plot showing the power signal used for spike detection and the detected spikes

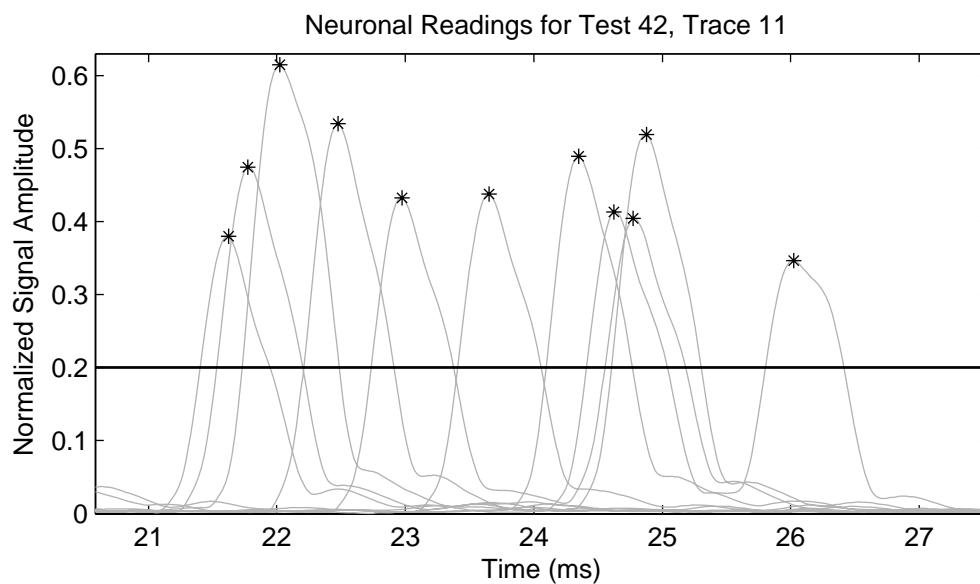


Figure 3: Zoomed in view of Figure 2

power, and hence don't necessarily occur at the minima or maxima of the spike. The power of the signals are indicated in grey (they are difficult to see with the MER signals overlaid).

The bottom plot of Figure 1 shows the histogram of the detected spikes in grey. The horizontal black lines indicate the duration of the stimulus in the test. In this case, the test consisted of two tones being presented. Each starts at 10 milliseconds and is 50 milliseconds in duration

By turning off the plotting of the spikes and the histogram features (e.g., by setting `plot_flag` to `[0 1 1 0 0 0]`), one can view the power of the MER signals, the power peaks used for spike detection, and the threshold used for the detection. The same data used in Figure 1 viewed in this format is shown in Figure 2. Figure 3 shows a zoomed in view of Figure 2. Here, it is clear that the peaks of the MER power are well above the default threshold and the noise floor is well below.

10 Example

The following example assumes that the Batlab XML file has already been generated from the PST file using the Batgor program. Both the XML and raw data files are needed to use Bat2Matlab. Running the example will generate the plots in Figures 1 and 2.

```
raw_data_filepath = 'bat 7b.raw';
xml_data_filepath = 'bat 7b-alltests.xml';
%Extract XML metadata and convert to Matlab structure
experiment_data = BatlabXml2MatlabStructure(xml_data_filepath);

%Specify which tests to extract raw data from
%This can be in the form [42 46] or [42:52] to specify multiple tests
test_nums_to_extract = 42;

experiment_data = ExtractRawData(experiment_data,raw_data_filepath,test_nums_to_extract);
experiment_data = CalculateSpikeTimes(experiment_data,test_nums_to_extract);
experiment_data = GenerateHistograms(experiment_data,test_nums_to_extract);

%Specify the trace number to view
trace_num = 11;
PlotData(experiment_data,test_nums_to_extract,trace_num,[1 1 1 1 1 1]);
PlotData(experiment_data,test_nums_to_extract,trace_num,[0 1 1 0 0 0]);
```

11 Appendix A: Batlab Metadata XML Schema

```
<?xml version="1.0" encoding="UTF-8" ?>

<xs:schema xmlns:xs="http://www.w3.org/2001/XMLSchema">
  <xs:element name="experiment">
    <xs:complexType>
      <xs:sequence>
        <xs:element ref="test" maxOccurs="unbounded" />
      </xs:sequence>
      <xs:attribute name="computername" type="xs:string" use="required" />
      <xs:attribute name="date" type="xs:string" use="required" />
      <xs:attribute name="who" type="xs:string" use="required" />
      <xs:attribute name="pst_filename" type="xs:string" use="required" />
      <xs:attribute name="program_date" type="xs:string" use="required" />
      <xs:attribute name="title" type="xs:string" use="required" />
    </xs:complexType>
  </xs:element>
```

```

<xs:element name="spikedata">
  <xs:complexType mixed="true">
    <xs:attribute name="units" type="xs:string" use="required" />
  </xs:complexType>
</xs:element>

<xs:element name="stimulus">
  <xs:complexType>
    <xs:attribute name="soundtype" type="xs:NMTOKEN" use="required" />
    <xs:attribute name="AMdepth" type="xs:decimal" use="optional" />
    <xs:attribute name="AMdutycycle" type="xs:decimal" use="optional" />
    <xs:attribute name="duration" type="xs:decimal" use="optional" />
    <xs:attribute name="delay" type="xs:decimal" use="optional" />
    <xs:attribute name="reverse_vocal_call" type="xs:NMTOKEN" use="optional" />
    <xs:attribute name="attenuation" type="xs:decimal" use="optional" />
    <xs:attribute name="frequency" type="xs:decimal" use="optional" />
    <xs:attribute name="AMrate" type="xs:decimal" use="required" />
    <xs:attribute name="vocal_call_file" type="xs:NMTOKEN" use="optional" />
    <xs:attribute name="AMtype" type="xs:NMTOKEN" use="required" />
    <xs:attribute name="phase" type="xs:decimal" use="optional" />
    <xs:attribute name="soundtype_name" type="xs:string" use="required" />
    <xs:attribute name="rise_fall" type="xs:NMTOKEN" use="optional" />
  </xs:complexType>
</xs:element>

<xs:element name="test">
  <xs:complexType>
    <xs:sequence>
      <xs:element ref="trace" maxOccurs="unbounded" />
    </xs:sequence>
    <xs:attribute name="length_in_raw_file" type="xs:positiveInteger" use="required" />
    <xs:attribute name="date" type="xs:string" use="required" />
    <xs:attribute name="testtype" type="xs:NMTOKEN" use="required" />
    <xs:attribute name="id" type="xs:NMTOKEN" use="required" />
    <xs:attribute name="comment" type="xs:string" use="required" />
  </xs:complexType>
</xs:element>

<xs:element name="trace">
  <xs:complexType>
    <xs:all>
      <xs:element ref="spikedata" />
      <xs:element ref="stimulus" />
    </xs:all>
    <xs:attribute name="window_detector_status" type="xs:NMTOKEN" use="required" />
    <xs:attribute name="level_detector_status" type="xs:NMTOKEN" use="required" />
    <xs:attribute name="run_number" type="xs:positiveInteger" use="required" />
    <xs:attribute name="level_voltage" type="xs:decimal" use="required" />
    <xs:attribute name="spike_enhancer_status" type="xs:NMTOKEN" use="required" />
    <xs:attribute name="record_duration" type="xs:positiveInteger" use="required" />
    <xs:attribute name="soundtype" type="xs:NMTOKEN" use="required" />
    <xs:attribute name="window_voltage" type="xs:decimal" use="required" />
    <xs:attribute name="is_control" type="xs:NMTOKEN" use="required" />
    <xs:attribute name="spike_enhancer_power_value" type="xs:decimal" use="required" />
    <xs:attribute name="length_in_raw_file" type="xs:positiveInteger" use="required" />
    <xs:attribute name="samplerate_ad" type="xs:positiveInteger" use="required" />
    <xs:attribute name="display_duration" type="xs:decimal" use="required" />
    <xs:attribute name="binwidth" type="xs:positiveInteger" use="required" />
    <xs:attribute name="peak_detector_status" type="xs:NMTOKEN" use="required" />
  </xs:complexType>
</xs:element>

```

```

<xs:attribute name="samplerate_da" type="xs:positiveInteger" use="required" />
<xs:attribute name="invert_raw_data_status" type="xs:NMTOKEN" use="required" />
<xs:attribute name="num_samples" type="xs:positiveInteger" use="required" />
<xs:attribute name="repetition_rate" type="xs:positiveInteger" use="required" />
<xs:attribute name="samplerate_type" type="xs:NMTOKEN" use="required" />
<xs:attribute name="stats_start" type="xs:positiveInteger" use="required" />
<xs:attribute name="stats_end" type="xs:positiveInteger" use="required" />
</xs:complexType>
</xs:element>
</xs:schema>

```

12 Appendix B: Bat2Matlab Structure

```

experiment.computername
  date
  program_date
  pst_filename
  title
  who
  test.comment
    date
    id
    length_in_raw_file
    offset_in_raw_file
    testtype
    trace.display_duration
      invert_raw_data_status
      is_control
      length_in_raw_file
      level_detector_status
      level_voltage
      num_samples
      peak_detector_status
      record_duration
      repetition_rate
      run_number
      samplerate_ad
      samplerate_da
      samplerate_type
      soundtype
      spike_enhancer_power_value
      spike_enhancer_status
      stats_end
      stats_start
      window_detector_status
      window_voltage
      stimulus
      raw_data
      spike_idx
      spike_times
      smoothed_data
      peak_threshold
      histogram
      histogram_bin_width
      histogram_bin_centers

```