
pybdf Documentation

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CONTENTS

1	Introduction	3
2	Download and Installation	5
2.1	Download	5
2.2	Installation	5
3	Usage	7
4	Bugs	9
4.1	Known issues	9
5	pybdf – Class to read BIOSEMI BDF files	11
6	Indices and tables	13

Contents:

INTRODUCTION

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pybdf is a python library to read BIOSEMI 24-bit BDF files.

DOWNLOAD AND INSTALLATION

2.1 Download

The latest release of pybdf can be downloaded from the python package index:

<http://pypi.python.org/pypi/pybdf/>

For developers: the source code of pybdf is hosted on github:

<https://github.com/sam81/pybdf>

2.2 Installation

2.2.1 Requirements

To install pybdf you will need:

- python
- numpy
- a fortran compiler

On all platforms, after having unpacked the archive you can install pybdf by running:

```
python setup.py install
```

Note that pybdf has been built and tested only on Linux. I don't have enough time to build and test binaries for Windows and Mac OS X, but if you try, please let me know how you get on with it.

USAGE

To open a bdf file you need to create a `bdfRecording` object as follows:

```
bdfRec = bdfRecording('res1.bdf')
```

you can then query the properties of the recording stored in the BDF header using the appropriate functions, which are fully described [here](#). Some examples are shown below.

Get the duration of the recording:

```
bdfRec.recordDuration
```

Get the sampling rate of each channel:

```
bdfRec.sampRate
```

Get the channel labels:

```
bdfRec.chanLabels
```

To read in the data use the following method:

```
rec = bdfRec.getData()
```

this returns a python dictionary with the following fields:

- `data` : an array of floats with dimensions `nChannels X nDataPoints`
- `eventTable` : a dictionary with the codes, indexes and durations of triggers
- `chanLabels` : a list with the channel labels

For example, to get the value of the first sample of the recording, in the first channel, you can type:

```
rec['data'][0,0]
```

the same sample value, but for the second channel, is stored in:

```
rec['data'][1,0]
```

The *eventTable* contains a list of the trigger codes for the experimental conditions:

```
rec['eventTable']['code']
```

as well as a list of the sample numbers (or indexes) at which they started in the recording, and a list of their durations in seconds:

```
rec['eventTable']['idx']  
rec['eventTable']['dur']
```

The `ActiveTwo` actually stores one trigger code for each recording sample rather than a list of trigger onsets and durations as the `eventTable` does. The “raw” trigger channel with one trigger code for each recording sample can be retrieved by passing the argument `trigChan = True` to the `getData()` function:

```
rec = bdfRec.getData(trigChan=True)
```

the “raw” trigger channel will then be returned in `rec['trigChan']`.

It is also possible to retrieve additional system codes (bits 16-23 of the status channel, see http://www.biosemi.com/faq/trigger_signals.htm), like CMS in/out-of range, battery low/OK. These are returned in `rec['sysCodeChan']` when the `sysCodeChan = True` argument is passed to the `getData()` function. No particular effort has been made to decode these system codes.

Other usage examples are provided in the ‘examples’ directory inside the pybdf source archive.

Beware that pybdf does not check that you have sufficient RAM to read all the data in a bdf file. If you try to read a file that is too big for your hardware, your system may become slow or unresponsive. Initially try reading only a small amount of data, and check how much RAM that uses. You can read only a portion of the data by passing the `beginning` and `end` arguments to the `getData()` function. For example, to read the first 10 seconds of the recording, use:

```
rec = bdfRec.getData(beginning=0, end=10)
```

BUGS

Please, report any bugs on github <https://github.com/sam81/pybdf/issues>

4.1 Known issues

The filename or filepath of the BDF recording is currently limited to 256 characters. That should be sufficient for most purposes.

PYBDF – CLASS TO READ BIOSEMI BDF FILES

INDICES AND TABLES

- *genindex*
- *modindex*
- *search*