

OCD Analysis Software (v 1.0)

Nick Charron - Huang Biophysics Lab

Rice University. 2017.

This software is designed to plot and process spectra obtained from the J-810 Jasco CD spectrometer in the Huang lab. All measurement files must be in ASCII format (*.txt files). This format can be chosen when saving the measured spectra in the SPECTRA MANAGER software provided by Jasco. File reading routines are currently optimized for J-810 ASCII output, though these can be modified to read any consistent text-based spectral data file.

The software is designed to work as a collection of tools, so that it may be used interactively. To start an interactive python session, type the following command at your shell or command prompt:

```
>_ python
Python 2.7.13 (default, Dec 17 2016, 23:03:43)
[GCC 4.2.1 Compatible Apple LLVM 8.0.0 (clang-800.0.42.1)] on darwin
Type "help", "copyright", "credits" or "license" for more information.
>>>_
```

Depending on your distribution and machine details, the output above may not be exactly the same (This output was generated on a 2015 Apple MPB with a brew install of python 2.7). Next, the tools must be imported:

```
>>> from ocd import *
```

This also imports the following python modules: numpy as np, and matplotlib.pyplot as plt. Loading a spectrum is done through the creation of an instance of the class ocd_spec. Providing a filename loads the corresponding spectral information (wavelength and CD signal). Optionally the class may be called without a filename to create an empty spectrum (no spectral information):

```
>>> myspec = ocd_spec("scan_01.txt") #class with filename
>>> emptyspec = ocd_spec()           #empty class
```

You can also use file dialogs to select the file from your current directory. On Windows systems, this functionality is supplied by the wxPython-Phoenix package, which interfaces with the system default file manager and pipes the selected filename to the creation of an ocd_spec instance. *On Linux and OSX systems, the same process is achieved through*

verbatim *lll* myspec = fs()

Once a spectrum is loaded, you have access to the following functions and variables:

ocd_spec.wl : returns a 1D numpy array of wavelengths used in the scan

ocd_spec.cd : returns a 1D numpy array of CD signals measured

ocd_spec.load(string) : loads wl and cd data from specified filename

ocd_spec.name : changes the name of the spectrum

ocd_spec.graph() : produces a simple graph of the spectrum

ocd_spec.renorm(float) : globally rescales the CD signal by specified factor

avg_signal([spec1, spec2, ...]) : returns an ocd_spec class with an average CD signal from spec1,spec2, ... The spectra must be passed to this function as a python list.

mult_graph(specs,types,colors,title) : produces a graph of multiple spectra with custom style options. Specs, types, and colors must all be lists of the same size. A legend will automatically be generated using the values of spec.name.

graph_series(specs,title,cmap) : produces a graph of a sequence of experiments - useful for showing spectral changes dependent on concentration. The color map can be chosen from the matplotlib package, though the default is Reds(). A legend will automatically be generated using the values of spec.name.

In this way, multiple spectra can be graphed, rescaled, and analyzed interactively at the user's discretion. Custom graphs beyond the scope of the simple **mult_graph()** function (advanced color schemes, plotstyles, line types, etc.) can be made by directly calling functions from the pyplot module:

```
>>> fig = plt.figure("Custom Plot")
>>> plt.plot(myspec.wl, myspec.cd, 'b--',linewidth=2)
>>> plt.xlim(220,240)
>>> plt.title("230 nm Peak - A State")
>>> plt.ylabel("CD [mdeg]")
>>> plt.xlabel("Wavelength [nm]")
>>> plt.show()
```