

Processing Data How-To v2.1

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1 Installation

There are several steps to installing and setting up the *olberg-tools* program. The first is to make download the *olberg-tools* files from <https://github.com/RoundTab1er/olberg-tools>. The *olberg-tools* folder should be added to the path.

The second step is to install PuTTY, a client which allows the Windows computer to communicate with the computer containing the combinato cluster processing program. Go to this link, <https://www.chiark.greenend.org.uk/~sgtatham/putty/latest.html>, download the 64-bit installer in the first box (labelled Package Files), and install.

The third step is to set up combinato (link: <https://github.com/jniediek/combinato>). Installation steps are listed on that website (THIS IS ALREADY DONE FOR THE IMAC NEXT TO THE NEW WINDOWS MACHINE).

The final step is to run the *setup.bat* in the *olberg_tools* folder. This copies an important file to the combinato directory on the remote machine.

Now you should be all set up. All *olberg-tools* scripts are contained in the *olberg – tools* folder.

2 Save LabChart Data

The first step is to save the data recorded in LabChart to a MatLab compatible *.mat* file. Remove all of the blocks that don't contain relevant data, and save as a MatLab file. You will be given the choice of which channels to export. Only two channels are needed: the AC-coupled intracellular trace (as of writing channel 4), and the AC-coupled extracellular trace (as of writing channel 2). Move this to your MatLab path (anywhere should be fine). In addition, make sure the *olberg_tools_master* folder is in your path.

3 Set up prefs

Open *olberg_tools_prefs* and set up all the prefs as desired. Each variable has a comment describing its function.

Editing the preferences for the combinato clustering program requires editing *options.py*, which is located in the combinato directory on the imac. The path is `/anaconda2/lib/python2.7/site-packages/combinato/combinato/options.py`. The explanations of what each variable does is located here: <https://github.com/jniediek/combinato/wiki/Details>.

4 Convert Data

The next step is to convert the data to the form combinato requires. Combinato requires a $1 \times n$ *data* vector and a sampling rate variable named *sr*. Run *olberg_tool_file_processing_master*. This creates a folder called *data* in the *matlab* folder, and creates a subfolder for the file being processed. Once you run the script there will be a file in *data/*`{file_being_processed}`, named as the file you specified, with *_processed* (eg. `2017-06-07-trimmed-processed.mat`) on the end.

5 Run Combinato (Clustering)

Next, run the *olberg_tools_combinato_master.sh* script. This logs into a remote system, runs combinato, and copies the output files back to your pc. It will ask you for the name of the file you are processing, minus the *.mat*. It will then prompt you for the password of the system you are logging into three times (hopefully this will be improved soon!).

6 Process output data

Now we convert the data (output as *.h5* files) to *.mat* files. Run *olberg_tools_analysis_master*.

7 Visualizing Data

This will graph the results! The script requires a file name, channel, block, and cluster id's, and the *intra/extra_col* variables simply designate the column where the *intra/extra* block data exists in the *block_data* array.