utilities

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1 crossings2edt

The crossings2edt program generates an .edt file from a .chan file, based on positive-going crossings of a specified threshold. This can be useful if there is one large, clean unit on the channel. One spike is placed in the .edt file for each place in the .chan file where there is a sample below threshold followed by zero or more samples at threshold, followed by a sample above threshold. The spike time is the time of the above-threshold sample rounded to the nearest .1 ms. Running the program with no arguments gives this usage message:

usage: crossings2edt whatever.chan threshold unit_code

whatever.chan is the input .chan file. The output .edt file will have the same base name, but with an .edt extension. Any existing file by that name will be overwritten without warning. threshold is a number between -32768 and 32767 chosen by the user, typically by reading it off the waveform display (using waveform.tcl) as the fifth number displayed at the top, when the cursor is positioned at what looks like an appropriate threshold level. unit_code is the cell ID that will be used for the spike train in the .edt file.

2 crossings2pos

The crossings2pos program generates a .pos file with spike times for use with the spikesorter, from a .chan file, based on positive-going crossings of a specified threshold. This can be useful if there is one large, clean unit on the channel. Running it with no arguments gives this usage message:

usage: crossings2pos [-m THR2] POS CHAN THR CLUSTER... > OUTPOS

The program is intended to be used with a POS file that has been generated by the spikesorter from CHAN, but for which simple threshold crossings would have done a better job. THR is a number between -32768 and 32767 chosen by the user, typically by reading it off the waveform display (using waveform.tcl) as the fifth number displayed at the top, when the cursor is positioned at what looks like an appropriate threshold level. CLUSTER... is a space-separated list of cluster numbers in POS that will be left out of OUTPOS. The first number in the list will be re-used in OUTPOS as the cluster number for the threshold-crossing spikes. If the old .pos file is deleted or renamed or moved, the new .pos file can be given the name of the old one, and then it can be viewed in the spikesorter. The waveform overlay for the new cluster will be wrong, as will scatterplots that include it, but everything else should work.

If a second threshold is specified with the -t option, a spike will not be placed at a positive-going crossing of the first threshold unless the signal in CHAN goes back below the first threshold without going above the second threshold.

An example session might look like this:

```
cd 2004-01-25_001

crossings2pos 2004-01-08_001_11.pos 2004-01-08_001_11.chan 28000 1 2 3 4 5 > new.pos

mkdir -p save

mv -i 2004-01-08_001_11.pos save

mv new.pos 2004-01-08_001_11.pos
```

$3 \quad dmx$

The dmx program is used by spikesort_control_panel to determine the channel labels and which ones are enabled, but it can also be used by itself on the command line.

Invoked as

```
dmx DMXFILE > OUTFILE
```

where DMXFILE is a DataMAX header or recording file, it will translate the header to text and write it to OUTFILE.

Invoked as

```
dmx DMXFILE labels
```

it will print a list of the channel labels to the screen.

Invoked as

```
dmx DMXFILE enabled
```

it will print a list of zeroes and ones to the screen, indicating which channels are enabled (1 = enabled).

If DMXFILE is a recordong file, invoking it as

```
dmx DMXFILE N
```

will extract channel N (the first channel is channel 1) from DMXFILE and write it to a .chan file in the current directory. The name of the .chan file will be the

same as DMXFILE with any extension deleted, and with a two-digit channel number and the .chan extension appended. Andy existing file by that name will be overwritten without warning.

$4 \, \text{dmx_split}$

The dmx_split programs splits a DataMAX recording into .chan files. When invoked without arguments, it gives this usage message:

```
usage: dmx_split dmx_file_name
```

dmx_split creates two directories named clean and split in the current directory if they don't already exist. For each enabled channel, it will create a .chan file in the split directory. The name of the .chan file will be the same as dmx_file_name with periods changed to underscores, and with an underscore, a two-digit channel number, and the .chan extension appended. The first channel is channel 1 (not 0). If any of the .chan files already exist, dmx_split will ask once whether to overwrite them. If the answer is no, it will exit immediately and no .chan files will have been written.

It will also create a header file with the same name as dmx_file_name, but in the clean directory, containing just the header from dmx_file_name. If the file already exists, it will be overwritten without warning. If you answered no to overwriting .chan files as described above, the header file will still be overwritten if it exists, but the new file will be empty.

5 info_file.pl

The info_file.pl script reads a coordinate spreadsheet file and a .nam file and writes an "info" spreadsheet file. It is normally invoked automatically by spikesort_control_panel after the user clicks on the MERGE button, so the user doesn't need to know about it, but it can be invoked by itself from the command line if the user wants to recreate the info file without rerunning the merge. When invoked without arguments, it gives this usage message:

```
usage: info_file.pl datadir prefix
```

```
info_file.pl will look for a coordinate spreadsheet file at /oberon/experiments/prefix/prefix.xls info_file.pl will look for a .nam file at datadir/prefix.nam info_file.pl will write an info file at /oberon/experiments/prefix/prefix_info_orig.xls
```

The coordinate spreadsheet file must be in Excel 95-2003 format, and the output info file will be as well. The coordinate spreadsheet is created by hand. The .nam file is generated by spikesort_control_panel after the user clicks the MERGE button. (spikesort_control_panel calls merge_edt which calls info_file.pl.)

The .nam file is an ASCII text file with Linux-format line endings (line feed). It has one line per cell, with three items per line:

- 1. the digital channel number
- 2. the cell name assigned by edt_merge
- 3. the merged channel number assigned by edt_merge

info_file.pl requires the coordinate spreadsheet to have the following exact text in the indicated cells:

Cell C11: DIGITAL CHANNEL

Cell **I11:** A/P Cell **J11:** R/L

Cell K11: DEPTH

Cell A3: EXPERIMENT: Cell A5: RECORDING:

Cell A7: DATE:

There can be a second column of data with the same headers in the same rows, but starting in column P. If there is no second column, the header cells must be empty.

The data must start in row 12, and the background of the data cells must be colored. The row following the data must not be colored.

info_file.pl also requires the following information in the indicated cells:

Cell D48: the digital channel number of the reference electrode

Cell D50: the A/P coordinate of the reference electrode Cell D51: the R/L coordinate of the reference electrode Cell D52: the depth coordinate of the reference electrode

At USF, the coordinate spreadsheet is generated from a template in the Excel workbook found from Windows at

 $\verb|\cisc3| experiments | templates | coordinate_template_workbook.xls| (or at the coordinate_template) | templates | template$

/oberon/experiments/templates/coordinate_template_workbook.xls from Linux). Instructions can be found at

\\cisc3\experiments\templates\coordinate_template_instructions.doc.

The output info file for each experiment contains the name and date of the experiment; the recording number; the names, channel numbers, and coordinates of each cell; and places for the results of AA and STA testing. The data in the info file is read into xanalysis for display and, from there, is written out for import into a database (at USF, that's the GAIA database).

info_file.pl makes two copies of the output file, one with a name of the form prefix_info_orig.xls as in the usage message, and the other with a name of the form prefix_info_curr.xls. If prefix_info_orig.xls already exists, it is overwritten without warning. If prefix_info_curr.xls already exists, a warning is written to the command line and the old version is backed up to a file with a numbered extension, starting with prefix_info_curr.xls.~1~. No existing backups are overwritten or deleted. If AA or STA information was written to the old prefix_info_curr.xls, it is copied to both new files.

The two arguments to info_file.pl tell it where to find the input files and where to put the output file as shown in the usage message above. (The /oberon/experiments directory shown in the usage message is the path as seen from Linux. As seen from the Windows systems at USF, the same directory is \\cisc3\experiments.) It does not matter what the current working directory is when info_file.pl is invoked.

If the user wants to run info_file.pl without writing the output to the official directory, the user can specify a different directory as a third argument, and the output file will be written there instead.

6 integrate

The integrate program is used by spikesort_control_panel to process channels for which "I" is specified, but it can also be used by itself on the command line. Invoke it as

usage: integrate FILE.chan ANALOG_ID [CUTCODE CUT_EDT [flip]]

to rectify and integrate the signal in FILE.chan and write the result to FILE.edt with the specified ANALOG_ID. Also writes FILE.bin, which is a big-endian copy of FILE.chan. If FILE.edt or FILE.bin exists, it will be overwritten. Also appends an I to FILE.status.

If CUTCODE and CUT_EDT are specified, regions will be left out of the integrated signal in FILE.edt. The regions to be left out are specified by an event code in

an .edt file. CUTCODE is the id of the event code, and CUT_EDT is the name of the .edt file. The region between the first and second event of the given id is left out, and between the third and fourth, etc. In general, the region between each odd numbered event (start marker) and the following even numbered event (stop marker) is left out. If "flip" is specifed, the regions that would have been left out are kept, and vice-versa. The content of the left-out regions has no effect on the integrated signal.

Before each kept region, the integrator is initialized with the mean of the integrated signal, so there is little or no start-up ramp.

The samples of the integrated signal are 5 milliseconds apart, and the sampling clock continues to run during the left-out regions, so all intervals between samples are a multiple of 5 ms.

7 split_merge_abf

The split_merge_abf programs merges .abf files and then splits them into .chan files. When invoked without arguments, it gives this usage message:

usage: split_merge_abf whatever.abf [channel]...

The whatever.abf file must end in a .abf extension, but the .abf can be upper or lower case (or even mixed). The file is assumed to have a 2048 byte header, which is ignored, followed by consecutive interleaved 16-bit samples from 16 channels in channel order 1,9,5,13,2,10,6,14,3,11,7,15,4,12,8,16. There must be a multiple of 16 samples in the file. After processing the first .abf file it looks for another file with the same name except with an additional upper or lower case 'A' before the .abf extension (if there are both, it uses the one with the upper case). This file must also have a 2048 byte header, the same channel order, and a multiple of 16 samples, and the samples from this file are appended to those from the first file. The program then increments the ASCII value of the extra letter and looks for another file in the same way. This continues untils it fails to find the next file.

If no channel numbers are specified, all 16 channels will be extracted to .chan files. The file names will be the same as whatever.abf, except with the .abf extension replaced with a two-digit channel number followed by a .chan extension. Any existing files with the same names will be overwritten without warning. If channel numbers are specified (in a space-separated list), only those channels will be extracted

8 tkss.tcl

Using tkss.tcl, the spike sorter can be run without using the spikesort_control_panel, simply by typing

tkss.tcl

at the command line. This brings up what the Spike Sorter User Guide call the "Spike Sorter Window". It will not dispatch spike sorts to other computers like the control panel does, but you can run multiple spikesorts in parallel on the same computer using the SPIKESORT button to start them one at a time. They will continue to run after you quit. You can kill all the spike sorts you are running by typing

killall spikesort

at the command line.

You can also split a DataMAX file or an .abf file into chan files using the SPLIT button. tkss.tcl will not show you raw data, but you can use waveform.tcl for that. It will not show you the cluster "diagram" either, but you can get that by typing

dot -Tps FILENAME.dot > FILENAME.ps; gv -scale=2 FILENAME.ps

And it will not bring up the .notes file, but you can do that by typing

gedit FILENAME.notes

It will not invoke integrate, digitize, rpls, cpls, trachpls, muph, or chan2hdt for you, but those can all be run by hand. And it will not merge the .edt files for multiple channels into a single .edt file. You would have to do that by changing unit numbers with tmover or sed (figuring out yourself what they should be), and by merging the resulting .edt files with the merge utility.

But it does have all the functionality described under "RESULTS" in the Spike Sorter User Guide.