Analyse des séquences de potentiels d'action tutorial

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Required software

The code will be written mostly in C. If you want a clear and quick introduction to this language, check Ben Klemens: Modeling With Data. To compile the code you will need a C compiler like gcc. If you are using Linux or MacOS it's in a package from your favorite distribution, if you are using Windows you will have to install Cygwin. The heavy computational work is going to be performed mainly by the gsl (the *GNU Scientific Library*) that is easily installed through your package manager (from now on, for windows users, the "package manager" refers to the one of Cygwin). The graphs will be generated with gnuplot.

Data used

We are going to use spike trains obtained from the antennal lobe–first olfactory relay–of locust, *Schistocerca americana*. These spike trains can be found on the zenodo-locust-datasets-analysis GitHub repository. You can also find there a complete description of the sorting procedure used to go from the raw data, that are available on zenodo, to the spike trains. We will mostly use the spike trains from experiment locust20010214 that can be found at the following address: https://github.com/christophe-pouzat/zenodo-locust-datasets-analysis/tree/master/Locust_Analysis_with_R/locust20010214/locust20010214_spike_trains.

Getting a spike train

We will start by downloading the spike train from unit 1 from group Spontaneous_1. This is done by typing in the shell:

```
wget https://raw.githubusercontent.com/christophe-pouzat/\
zenodo-locust-datasets-analysis/master/Locust_Analysis_with_R/\
locust20010214/locust20010214_spike_trains/\
locust20010214_Spontaneous_1_tetB_u1.txt
```

This "spike train" contains in fact the result of 30 consecutive continuous acquisitions, each 29 s long with a 1 s gap in between, as is made clear in the detailed sorting description of this data set.

Making the "observed counting process" plot

We first get a gnuplot script, aspa_ocp.gp, from the github repository:

```
wget https://raw.githubusercontent.com/christophe-pouzat/\
aspa/master/gnuplot/aspa_ocp.gp
```

We then get a graph showing the observed counting process associated with the train by typing in the shell:

```
cat aspa_ocp.gp locust20010214_Spontaneous_1_tetB_u1.txt | \
gnuplot -persist
```

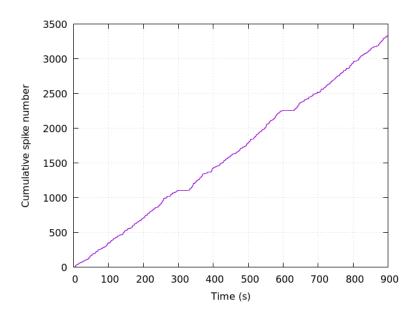


Figure 1: The observed counting process associted with $locust20010214_Spontaneous_1_tetB_u1.txt$ spike train

Notice the two "long" horizontal sections, these are due to the presence of recording noise during trials 11 and 21 (these trials were skipped during spike sorting).

C code

First test

To test that everything is properly set, we start by downloading the Makefile

```
wget https://raw.githubusercontent.com/christophe-pouzat/\
aspa/master/code/Makefile
```

as well as the header file aspa.h and the two C source files, aspa_single.c and aspa_single_test.c:

```
wget https://raw.githubusercontent.com/christophe-pouzat/\
aspa/master/code/aspa.h
wget https://raw.githubusercontent.com/christophe-pouzat/\
aspa/master/code/aspa_single.c
wget https://raw.githubusercontent.com/christophe-pouzat/\
aspa/master/code/aspa_single_test.c
```

We can then compile the test file (aspa_single_test.c) with:

```
make aspa_single_test
```

And we run it on the spike train we downloaded previously with:

```
cat locust20010214_Spontaneous_1_tetB_u1.txt | \
aspa_single_test
```

We should see the number of spikes (3331), the time of the first spike (0.290975 s), the time of the last one (898.15 s) printed followed by a bunch of statistics concerning the inter spike intervals (isi) distribution.

Notice that I've assumed here that the current directory is on your executable path, if not you can either add to the path with:

```
export PATH=.:$PATH
or use:
cat locust20010214_Spontaneous_1_tetB_u1.txt | \
./aspa_single_test
```

Checking memory leaks

One of the big advantages of C is that it let's you do efficient memory management; but this advantage comes at a cost: you have to do the memory management yourself meaning among other things that you should free the memory you allocate. A nice tool for checking that a code

does actually free everything it allocates is the Valgrind program. Once we have installed it, we can use it with:

Checking that there is no major mistake

We can give a regular sequence (with one spike every second, using the seq function) and see what it gives with (the spike times are entered here in sample points assuming a sampling rate of 15 kHz and a recording duration of 60 seconds):

```
seq 1 15000 900000 | aspa_single_test
```