Zerone tutorial

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1 Building instructions

Zerone is available as a Linux command line application and as an R package.

1.1 Downloading

We recommend that you use git to keep Zerone updated. You can clone the repository from Github with the following command on a standard terminal.

```
git clone git@github.com:gui11aume/zerone
```

Note that this requires that you already have a Github account and that the computer you are working on has an SSH key registered on GitHub. If this is not the case, follow the instructions from https://help.github.com/articles/generating-ssh-keys/.

Alternatively, if you prefer not to use git, you can download the source code from https://github.com/guillaume/zerone/archive/master.zip with the following commands.

```
wget https://github.com/guillaume/zerone/archive/master.zip
unzip zerone-master.zip
mv zerone-master zerone
```

This should create a directory named zerone.

1.2 Compiling

To build Zerone, execute the following from the zerone directory.

```
cd zerone
```

This should succeed on most Linux systems because make is available by default. If this is not the case, you can obtain it by typing sudo apt-get install make on the Ubuntu terminal.

Calling make should create an executable called zerone.

1.3 Testing

To check that the building was successful, test Zerone with the following commands.

```
cd src/test
make test
cd ../..
```

If it passes the tests without any error message, then everything went fine and you are done with the build. If not, something went wrong. In this case, you can explain how to reproduce the problem on https://github.com/guillaume/zerone/issues.

Installing Rzerone

To install the Rzerone R package, simply run this command from the zerone directory.

R CMD INSTALL Rzerone

Note that you need to have R installed on your computer. If this is not the case, run the command sudo apt-get install r-base on Ubuntu.

2 Zerone basics

2.1 Running Zerone

To run Zerone, you have to specify the files that contain the mapped reads of the ChIP-seq experiment you want to discretize. These can be in BED, SAM/BAM and GEM (.map) formats. You can include as many negative control files and as many experimental replicates as you need. Just enter controls after the -0 or --mock option, and targets after the -1 or --chip option.

For example, you can type in the following commands from the zerone directory.

```
./zerone --mock file1.bam,file2.bam --chip file3.bam,file4.bam
```

Where file1.bam and file2.bam are negative controls done without antibody, and file2.bam and file3.bam are two experimental replicates.

Note that path expansion will not work when using comma separated file names, so if you want to use path names starting with \sim , you can simply specify the mock and chip options as many times as needed, as shown below.

```
./zerone -0 file1.map -1 file2.map -1 file3.map
```

Enter the option -h or --help for usage instructions and --version to print the version number.

2.2 Output

Running Zerone as shown before should produce an output like the following.

```
# QC score: 0.732
```

advice: accept discretization.

chrX	1	300	0	0	0	0
chrX	301	600	0	0	0	0
chrX	601	900	0	0	0	0
chrX	901	1200	0	0	0	0
chrX	1201	1500	0	0	0	0
\mathtt{chrX}	1501	1800	0	0	0	0
chrX	1801	2100	0	0	0	0
chrX	2101	2400	0	0	0	0

The first two lines contain the result of the **quality control**. It consists of a quality score followed by an advice to either **accept or reject** the discretization, if the score is positive or negative, respectively. If the score is higher than 1 (or lower than -1), the advice is extremely reliable.

The rest of the lines contain the discretization proper. The first three columns specify the chromosome, start and end positions of each window.

The fourth column represents the **enrichment**. Zerone classifies each window into one of three possible states. States 0 and 1 represent two types of background signal. State 2 represents an **enriched window**.

The fifth column contains the read count of all the control profiles summed together, and the rest of the columns contain the read count of each of the target profiles.

2.3 List output

With the -1 or --list-output option, Zerone produces an alternative output in which only enriched windows are shown. Also, all contiguous windows are merged together.

```
# QC score: 0.732
# advice: accept discretization.
chrX
        2759401
                     2763000
        2772001
chrX
                     2786100
chrX
        3526801
                     3555900
chrX
        3526801
                     3555900
chrX
        3561301
                     3595500
chrX
        3622201
                     3625200
        3734101
                     3741900
chrX
chrX
        6966901
                     6996300
chrX
        7000501
                     7003800
```

3 Troubleshooting

In case Zerone crashes, start by recompiling it in debug mode. To do so, run the following commands from the zerone directory.

```
make clean
make debug
```

Then repeat the actions that triggered the crash and contact guillaume.filion@gmail.com attaching the debug information.