## Note S1 Tool availability and implementation

Smash++ is implemented in C++ language and is available at [1]. This tool is able to find and visualize rearrangements in sequences, FASTA and FASTQ files; although, it is highly recommended to use sequences as input. In the following sections, we describe installing and running Smash++.

#### S1.1 Install

In order to install Smash++ on Linux, run the following commands:

```
git clone https://github.com/smortezah/smashpp.git
cd smashpp
cmake .

make
```

#### S1.2 Run

A reference file and a target file are clearly mandatory to run Smash++ (without visualization). Running

```
1 ./smashpp
```

provides the following guide:

```
SYNOPSIS
2
     ./smashpp OPTIONS... -r REF-FILE -t TAR-FILE
3
4
   SAMPLE
5
   DESCRIPTION
6
7
     Mandatory arguments
     -r, --ref FILE
8
                                 reference file (Seq/Fasta/Fastq)
9
     -t, --tar FILE
                                                (Seq/Fasta/Fastq)
                                 target file
10
11
     Options
     -v, --verbose
                                 more information
12
     -1, --level INT
                                 level of compression: [0, 5]
13
     -m, --min INT
                                 min segment size: [1, 4294967295]
14
15
     -nr, --no-redun
                                 do NOT compute self complexity
                                 Entropy of 'N's: [0.0, 100.0]
     -e, --ent-n FLOAT
16
     -n, --nthr INT
                                 number of threads: [1, 8]
17
     -fs, --filter-scale S\,|\,M\,|\,L\, scale of the filter:
18
19
                                 {S|small, M|medium, L|large}
                                 window size: [1, 4294967295]
     -w. --wsize INT
20
     -wt, --wtype INT/STRING
                                 type of windowing function:
21
22
                                 {O|rectangular, 1|hamming, 2|hann,
23
                                 3|blackman, 4|triangular, 5|welch,
24
                                 6|sine, 7|nuttall}
     -d, --step
25
                    INT
                                 sampling steps
26
     -th, --thresh FLOAT
                                 threshold: [0.0, 20.0]
27
     -sb, --save-seq
                                 save sequence (input: Fasta/Fastq)
28
     -sp, --save-profile
                                 save profile (*.prf)
     -sf, --save-filter
                                 save filtered file (*.fil)
29
30
     -ss, --save-segment
                                 save segmented files (*-s_i)
     -sa, --save-all
                                 save profile, filetered and
31
32
                                 segmented files
     -h, --help
                                 usage guide
33
34
     -rm, --ref-model k,[w,d,] ir, a,g/t, ir, a,g:...
35
     -tm, --tar-model k, [w,d,]ir,a,g/t,ir,a,g:...
                                 parameters of models
36
37
                           (INT) k: context size
38
                           (INT) w: width of sketch in log2 form,
                                     e.g., set 10 for w=2^{-1}0=1024
39
                           (INT) d: depth of sketch
40
41
                           (INT) ir: inverted repeat: {0, 1, 2}
```

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```
42
                                        0: regular (not inverted)
43
                                        1: inverted, solely
44
                                        2: both regular and inverted
45
                           (FLOAT) a:
                                        estimator
46
                           (FLOAT)
                                        forgetting factor: [0.0, 1.0)
                                  g:
47
                             (INT) t:
                                       threshold (no. substitutions)
```

The arguments "-r" and "-t" are used to specify the reference and the target, respectively, which are highly recommended to have short names. Level of compression, that is an integer between 0 and 5, can be determined with "-l". By setting "-m" to an integer value, only those regions in the reference file that are bigger than that value can be considered for compression. Triggering "-nr" makes the tool not to perform the reference-free compression (self-complexity computation) part.

In implementation of the reference-based compression, we have replaced 'N' bases in the references and the targets with 'A's and 'T's, respectively. On reference-free compression, they are replaced with 'A's, in both references and targets. If a user tends to replace 'N' bases in a sequence with a normal distribution of 'A', 'C', 'G' and 'T's, he/she can employ GOOSE toolkit [2]. Note that we have set the entropy of 'N's to 2.0, by default, but it is possible for the user to set them to another value of interest, by "-e" option.

Building different finite-context models can be done in the multi-threaded fashion, setting "-n" to an integer. To find similar regions in the reference and the target, information profile (obtained by compression) needs to be filtered, of which the scale can be set as S (small), M (medium) or L (large). Size of the window and type of the windowing function, described in ??, can be set by "-w" and "-wt" options, respectively. Instead of considering the complete profile information, the user is able to make samples of it by steps of which size can be determined by "-d". For the purpose of segmenting the filtered information profile, the average entropy of reference-based compression is used as the threshold, by default. However, this threshold can be altered by "-th" option.

Smash++ accepts FASTA and FASTQ files as input, in addition to sequences. In these cases, the input files are converted to sequences and then processed further. It is possible to save these sequences by "-sb" option.

After obtaining the information profile, Smash++ filters it and then removes it, by default. However, it is possible to save the profile by "-sp" option. The same thing happens to the filtered file, i.e., it is segmented and then is removed. But, the user can use "-sf" to save the filtered file. Also, the segmented files can be saved using "-ss". The user can save all the information profile, filtered and segmented files, by triggering "-sa" option.

For the purpose of compression, either reference-based or reference-free, it is recommended to use "-l" option, since it configures the models automatically. However, using "-rm" and "-tm", the user would be able to manually configure the reference model, for reference-based compression, and the target model, for reference-free compression. Parameters of the models are described in detail in section ??.

Running Smash++ (without visualization), positions of the similar regions in the reference and the target, and also complexity of the regions is saved in a "\*.pos" file. To visualize this file, one can type

```
./smashpp -viz
   which gives
   SYNOPSIS
1
      ./smashpp -viz OPTIONS...
2
                                    -o SVG-FILE
3
4
   SAMPLE
5
6
   DESCRIPTION
7
     Mandatory arguments:
8
     POS-FILE
                                 positions file, generated by
Q
                                 Smash++ tool (*.pos)
10
      Options:
11
     -v, --verbose
                                 more information
12
```

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```
-o, --out SVG-FILE
                               output image name (*.svg)
13
                               reference name shown on output. If name
14
     -rn, --ref-name STRING
                               has space, use "s, e.g. "Seq label".
15
                               Default: name in header of position file.
16
17
      -tn, --tar-name STRING
                               target name shown on output
     -vv, --vertical
18
                               vertical view
     -nn, --no-nrc
19
                               do NOT show normalized
20
                               relative compression (NRC)
21
     -nr, --no-redun
                               do NOT show self complexity
     -ni. --no-inv
                               do NOT show inverse maps
22
     -ng, --no-reg
23
                               do NOT show regular maps
     -1, --link
24
                     INT
                               type of the link between maps: [1, 6]
25
          --color
                      INT
                               color mode: [0, 2]
     -c,
      -p, --opacity
26
                     FLOAT
                               opacity: [0.0, 1.0]
      -w, --width
27
                      TNT
                               width of the sequence: [15, 100]
     -s,
28
          --space
                      INT
                               space between sequences: [15, 200]
29
     -f,
          --mult
                      INT
                               multiplication factor for
30
                               color ID: [1, 255]
     -b, --begin
                      INT
                               beginning of color ID: [0, 255]
31
32
     -rt, --ref-tick INT
                               reference tick: [1, 4294967295]
                               target tick: [1, 4294967295]
33
     -tt. --tar-tick INT
34
     -th, --tick-human 0|1
                               tick human readable: 0=false, 1=true
     -m, --min
35
                     INT
                               minimum block size: [1, 4294967295]
     -h, --help
36
                               usage guide
```

The output of Smash++ visualizer is an "SVG" file, which its name is determined by "-o" option. By default, it is named "map.svg". Names of the reference and the target, which are going to be printed on the output image, can be altered by "-rn" and "-tn", respectively. They are by default the names written in the positions file. To have a vertical view of the image, instead of the default horizontal view, one can use "-vv" trigger.

Smash++ performs reference-based and reference-free compressions to calculate the normalized relative compression (NRC) and redundancy (self complexity), respectively. If the user is not interested in showing them, he/she can turn them off by "-nn" and "-nr" triggers. In addition, Smash++ considers both regular and reverse complement maps by default in its calculations. Triggering "-ni" and "-ng" will stop showing inverted and regular maps, respectively.

Options "-l", "-c", "-p", "-w", "-s", "-f" and "-b" can be used to change the appearance of the image. Assigning integers to "-rt" and "-tt" options will change the tick sizes of the reference and the target, respectively. Smash++ prints the sizes on axes in human readable format, e.g., 1K, 2M, etc. However, it can be triggered by "-th" option. Note that, here, 1K is equivalent to 1000 and not 1024. Finally, by setting "-m" to an integer value, only the regions that are bigger than that value will be illustrated.

### S1.3 Example

This section guides, step-by-step, employing Smash++ to find rearrangements.

# Install Smash++ and provide the required files

First, we install Smash++:

```
git clone https://github.com/smortezah/smashpp.git
cd smashpp
cmake .

make
```

Then, we copy Smash++'s binary file into example/ directory and go to that directory:

```
cp smashpp example/
cd example/
```

References 4

# References

[1] M. Hosseini, D. Pratas, and A. J. Pinho. Smash++. [Online]. Available: https://github.com/smortezah/smashpp

 $[2]\,$  D. Pratas. Goose. [Online]. Available: https://github.com/pratas/goose