

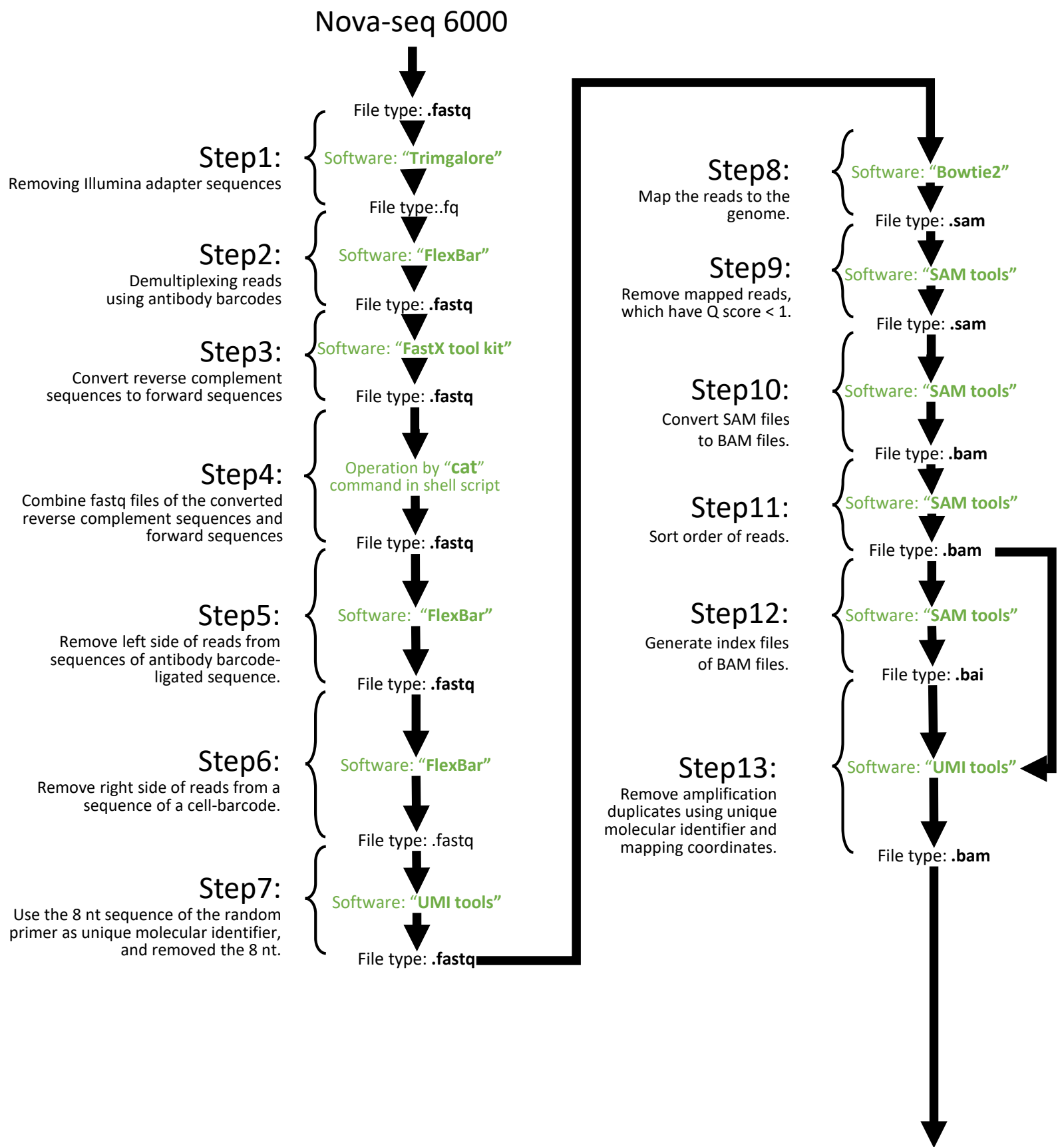
Page 1 Outline of the data analysis and data flow in the manuscript

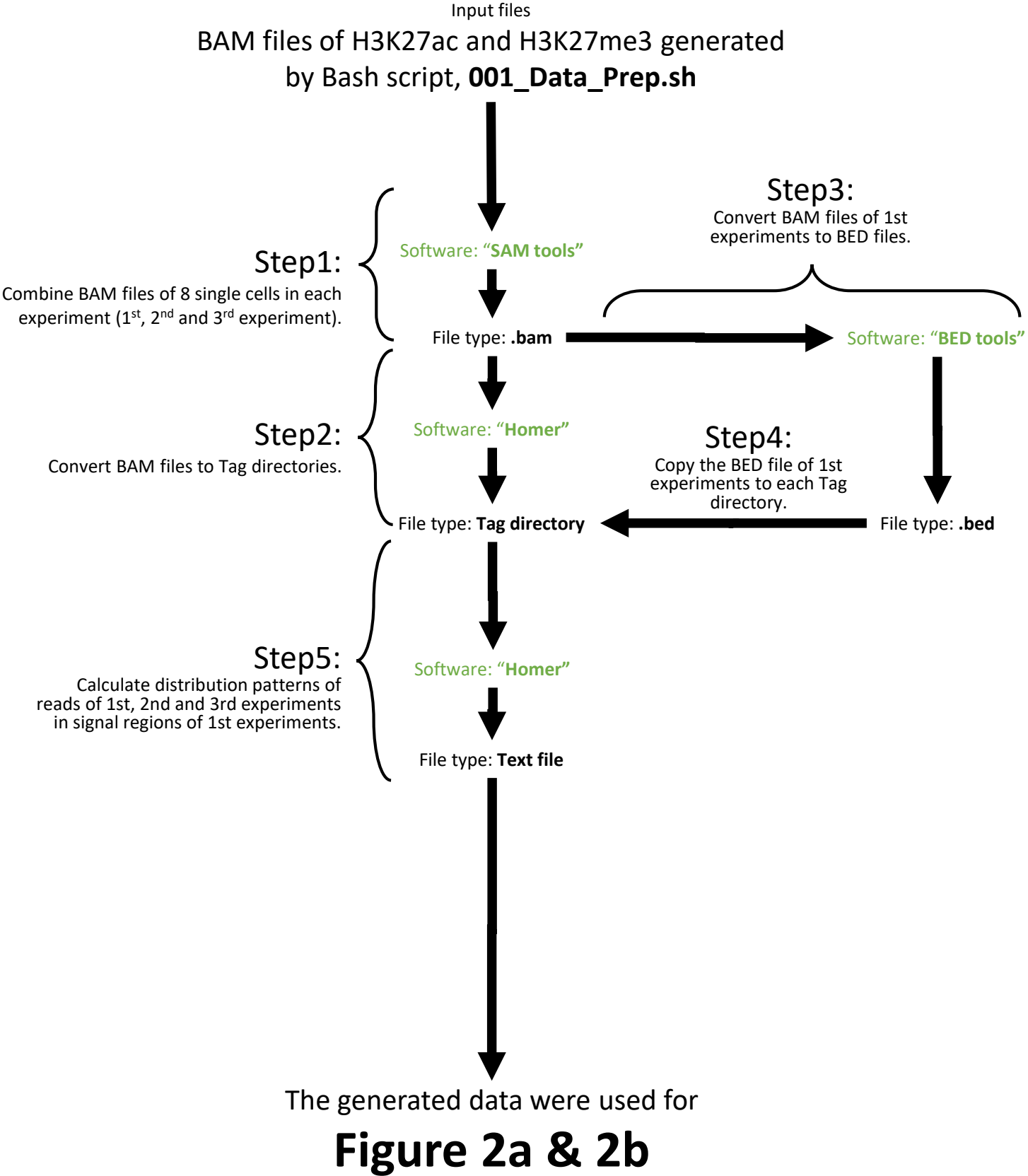
A high throughput sequence

Nova-seq 6000

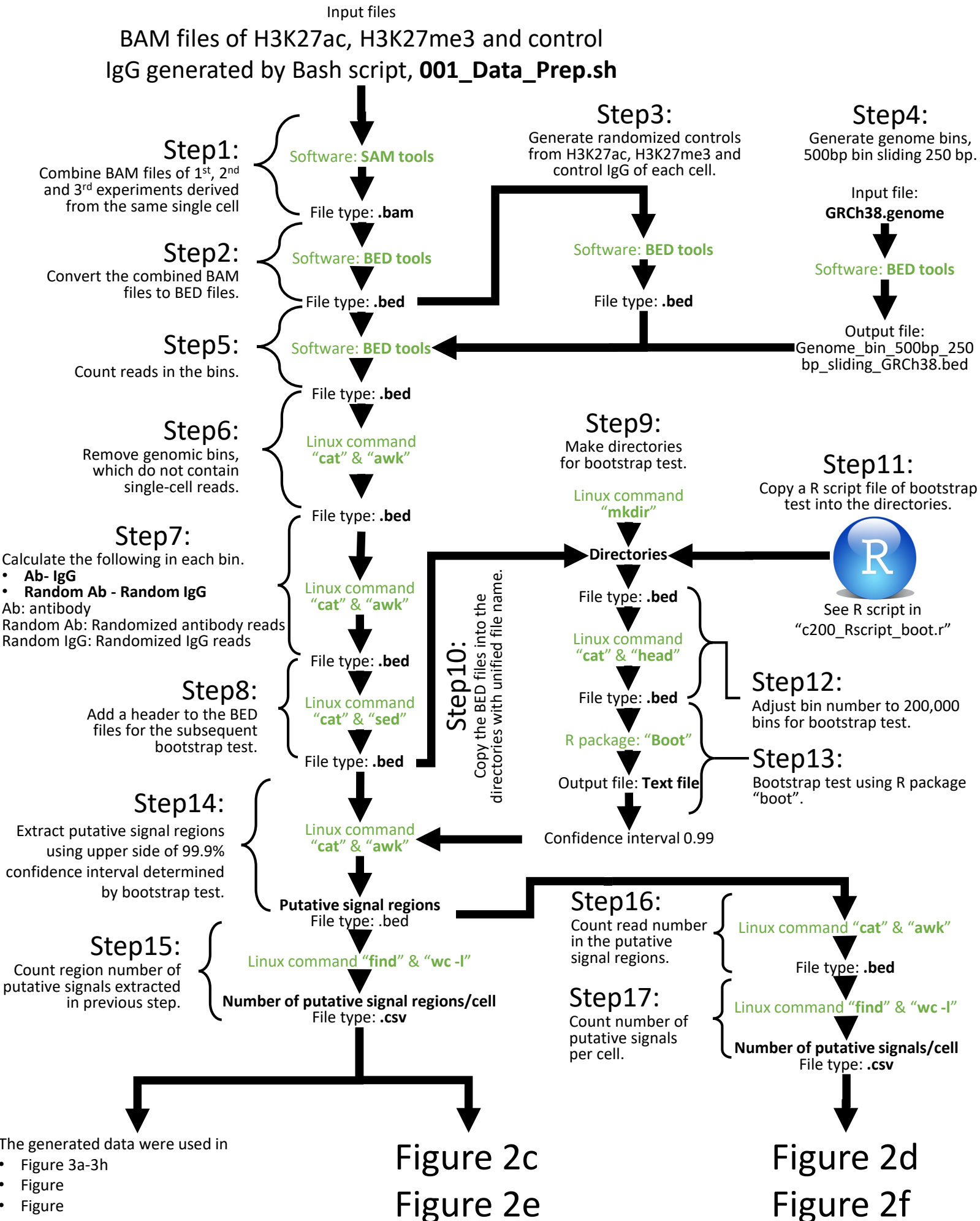
Output reads (Fastq files)

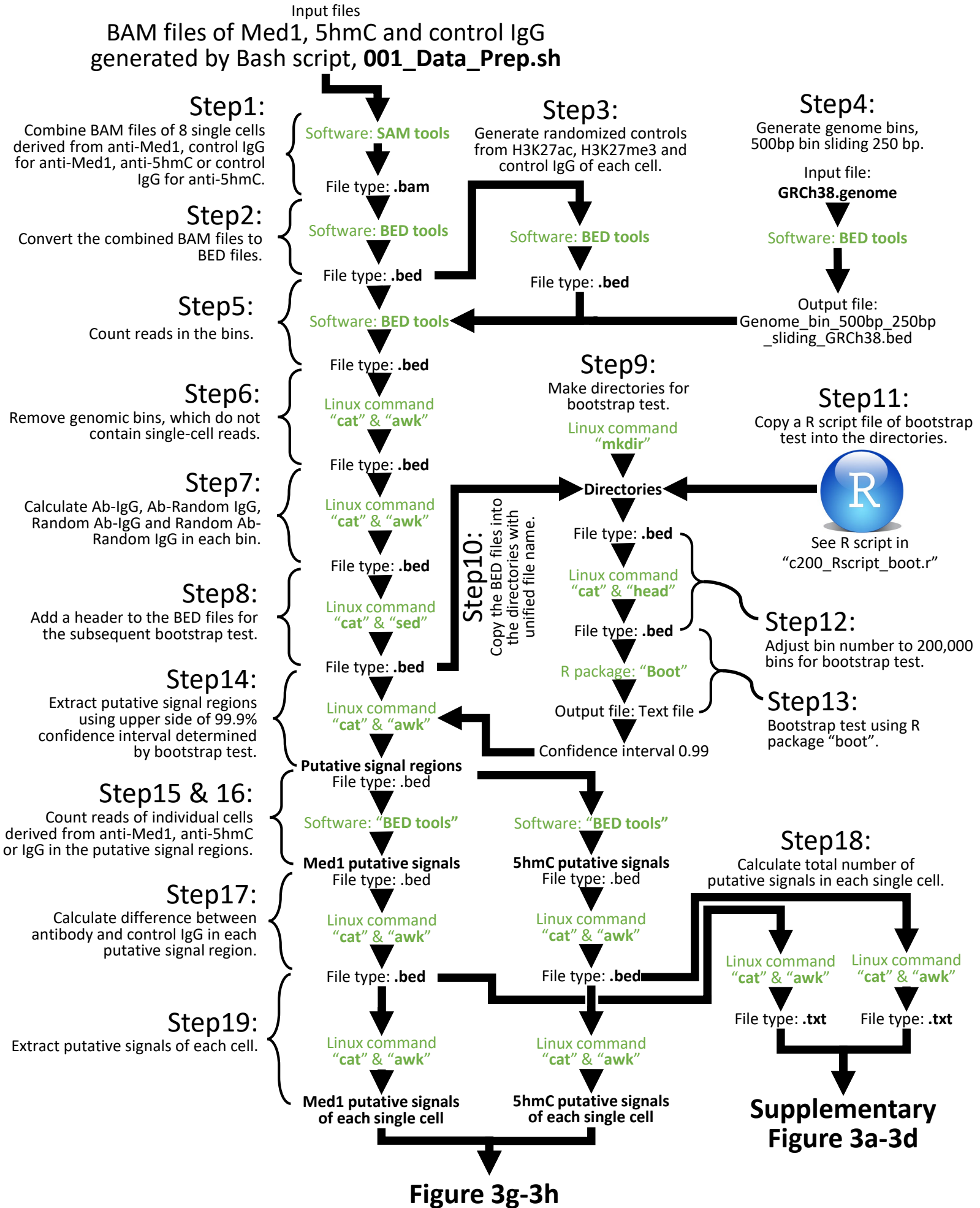


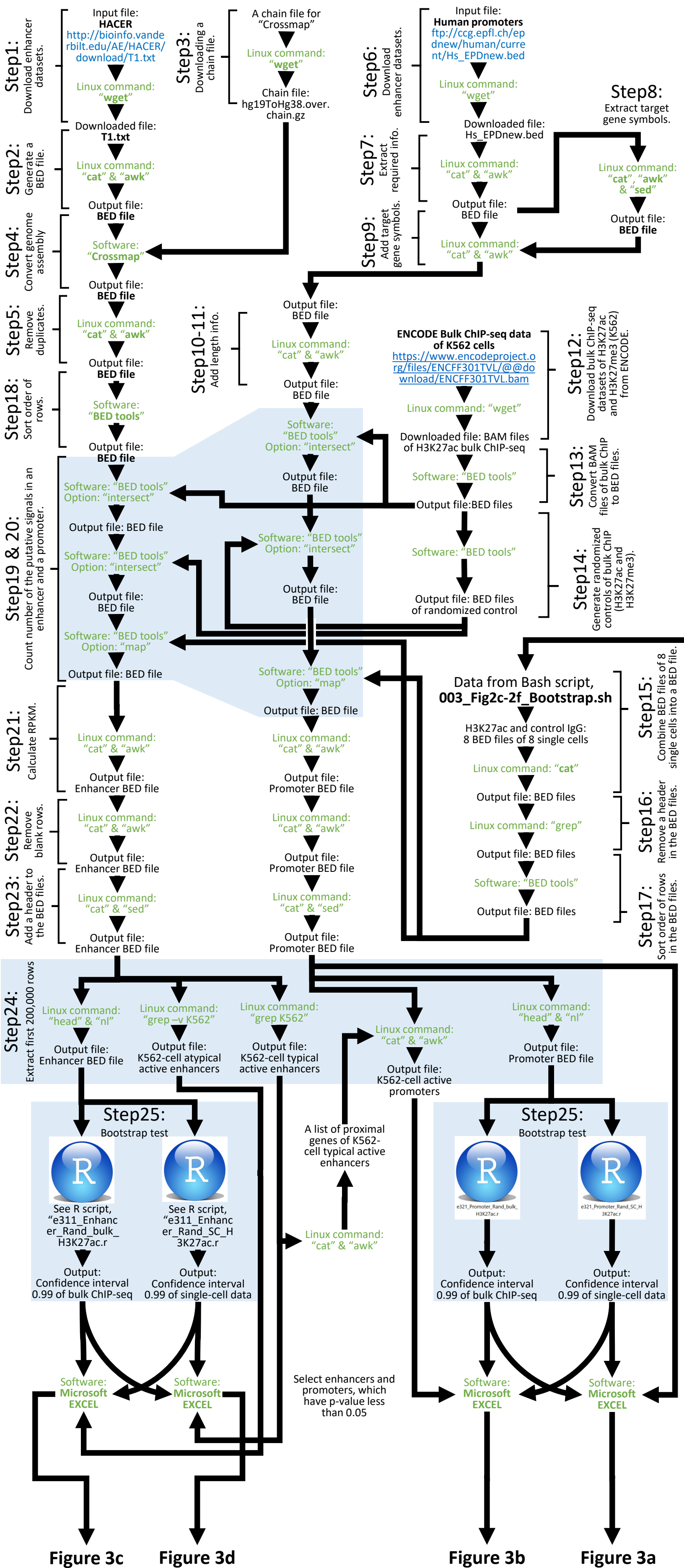


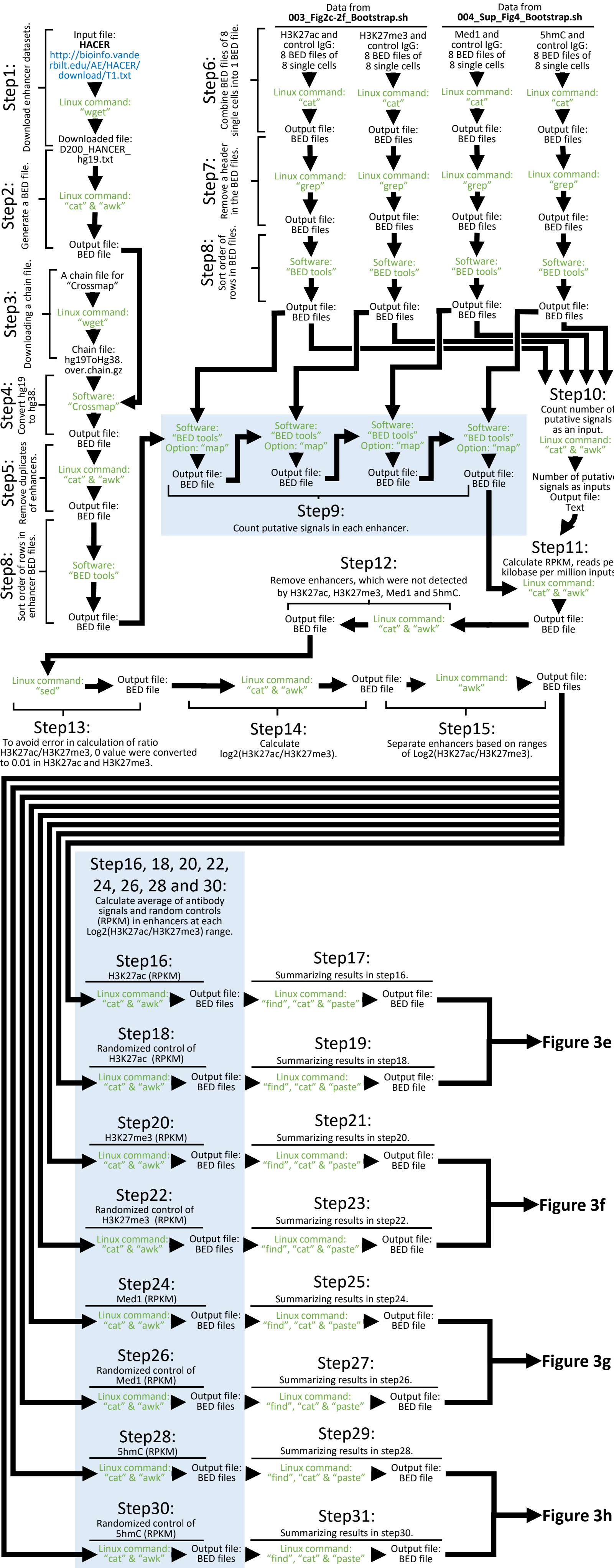


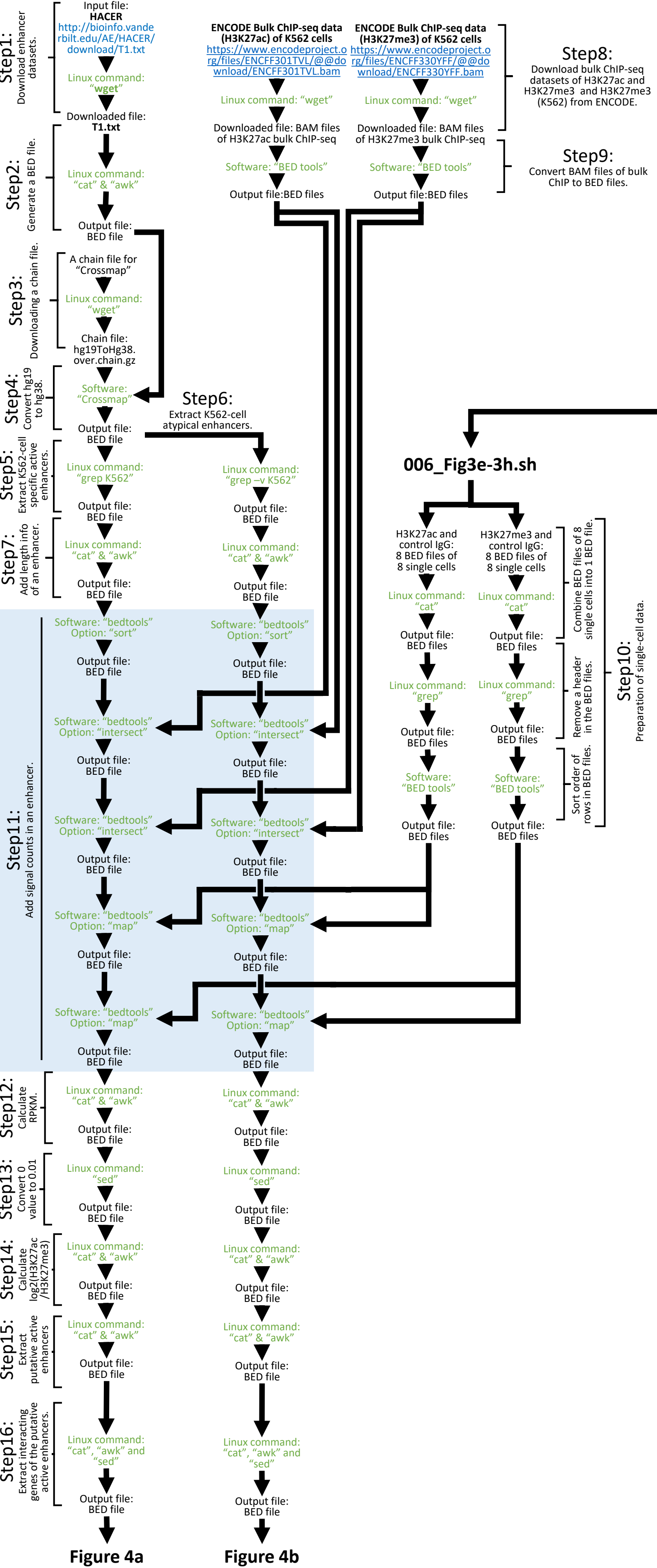
Bash script: 003_Fig2c-2f_Bootstrap.sh

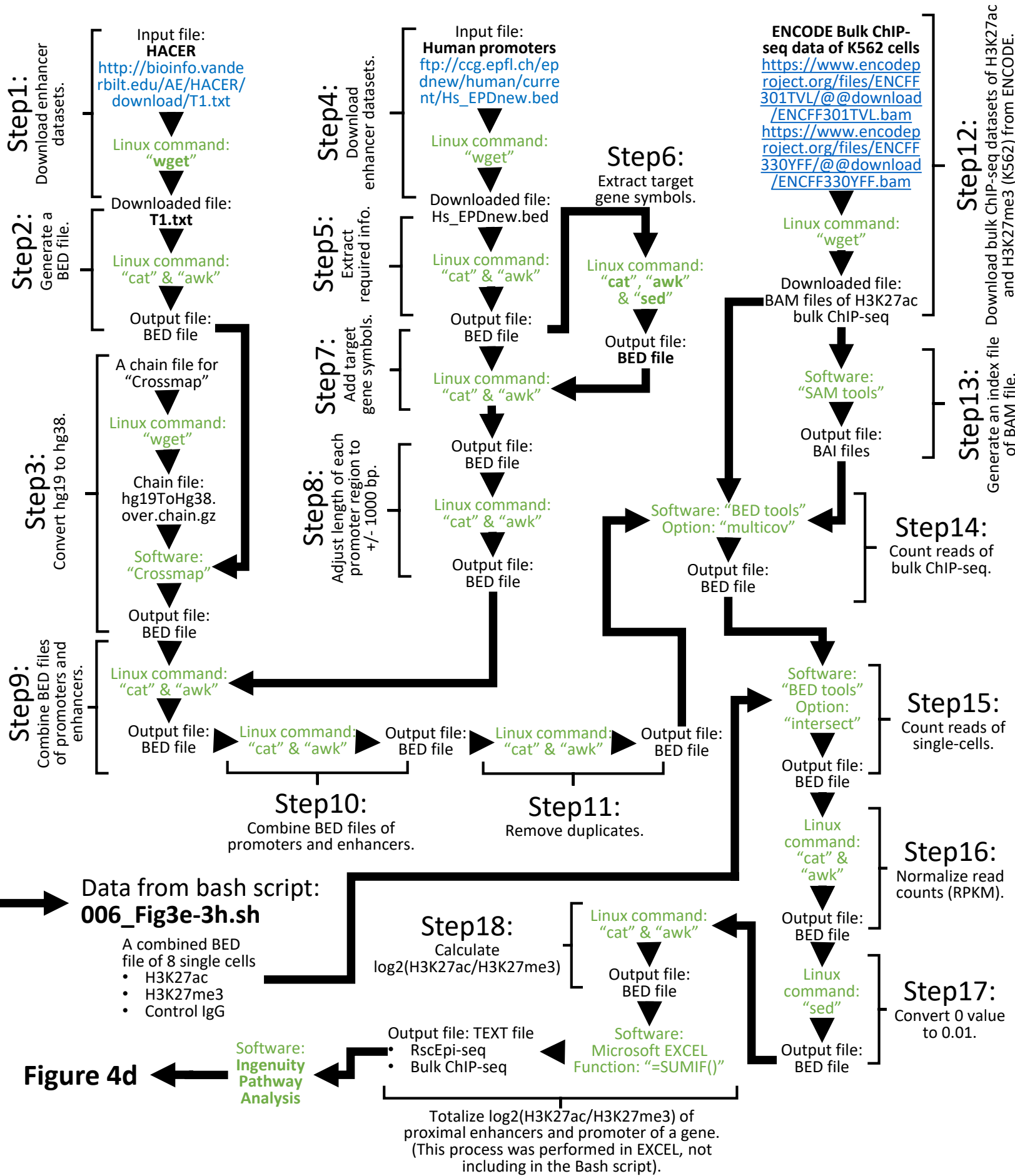














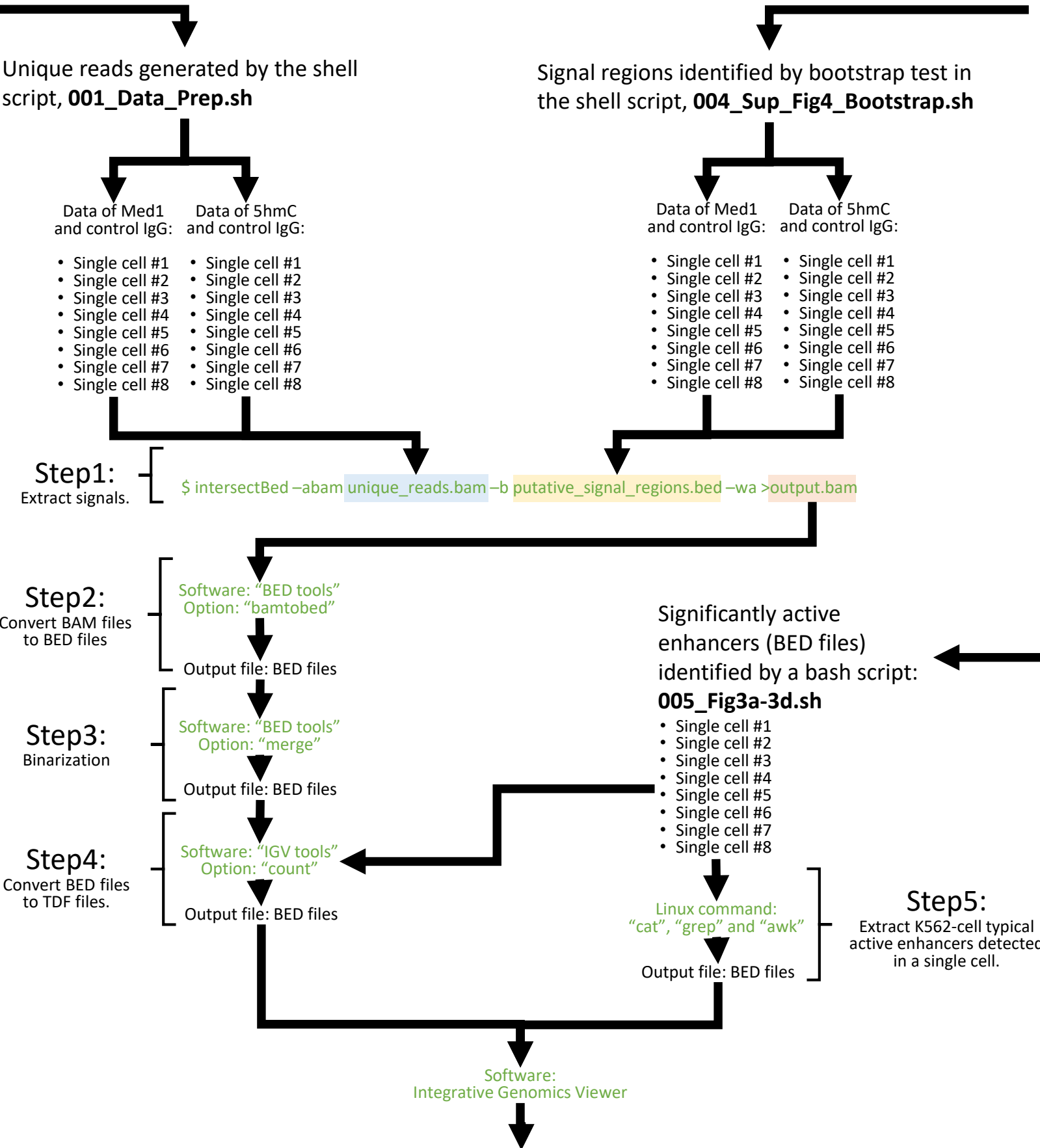


Figure 5b-5e and Supplementary Figure 5a-5c