
Supplementary Text SII

DEFAULTS: &defaults

UMI: 4

BARCODE: 6

GAP: 3

CDNA: 37

LAYOUT: src/barcodes.old.txt

PHYX:

GENOMESPIKERIBO: src/ebwt/hg19_ercc92_ynbA_u13369/ref

TRANSCRIPT: src/ebwt/hg19_refGene/ref

FLUCTUATION: 0.05

DIFFEXP: 0.05

LIBS:

SARMB:

<<: *defaults

FASTQS:

- src/150221_SN670_0304_AC655YACXX-lane7_NoIndex_L007_R1_001.fastq.gz
- src/140528_SN653_0292_AC4WG4ACXX-lane6_NoIndex_L006_R1_001.fastq.gz

SARMC:

<<: *defaults

FASTQS:

- src/150221_SN670_0304_AC655YACXX-lane8_NoIndex_L008_R1_001.fastq.gz
- src/140528_SN653_0292_AC4WG4ACXX-lane8_NoIndex_L008_R1_001.fastq.gz