

versions of software/tools used in RNA seq.

SRA version used sropath: 3.0.3 (3.0.2)
fastqc/0.12.1-gcc-13.2.0
RSEM: v1.3.1
bowtie2: 2.5.1
Samtools: 1.17
DESeq2: 1.50.2
R: 4.5.2