

versions of software/tools used in RNA seq.

SRA version used srapath: 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