

Next Generation Sequencing Data Analysis with R, spring 2023

Student Name: 张子栋

Student ID: 2020317210101

Submission Date: 6.16.2023

Project Title	对 NGS 数据的 ChIP-seq 分析			
R Packages used	RSamtools ChIPseeker			
Other tools used	LaTeX FastQC Trommomatic Bowtie2 MACS2			
Data type used*	ChIP-seq	RNA-seq	Hi-C	Other
	✓			
Group member	Name	Contribution		
1	张子栋	ALL		