

The Bioinformatics Core

(Office of Research Core Administration)

Maximizing Insights: Enhancing ATAC-Seq Experiment Design and Data Analysis

Topics of Discussion

- -Different Methods for Assessing Chromatin Accessibility
- -Experimental Design of ATAC-Seq
- -Pre-alignment QC
- -Post-alignment QC
- -Peak Calling
- -Peak Annotation
- -Differential Accessibility Analysis

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