Unveiling Cellular Complexity: Exploring the Analysis of Single-Cell RNA Sequencing Data

Topics of Discussion

-Advantage of Single Cell RNASeq

-10X and ParseBioscience Method

-Mapping

-Review Mapping & QC report

-Ambient RNA Removal

-Data Normalization

-Clustering

-Cluster Optimization

-Cluster Annotation

-Differential Composition Analysis

-Trajectory Analysis

-Pathway Analysis

-Cell-Cell Communication

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