

RNAseq plus

DECIDE TO GO FOR HIGH IMPACT

Complement your single cell transcriptomics data with bulk RNA sequencing data to validate your findings and reach for high impact publications.

COMBINE TO SHINE

Discover with Single Cell RNAseq



Explore innate heterogeneity

Single cell transcriptomics enable insights into sample heterogeneity providing more granular transcriptomic profiles of your samples.

Validate & Scale with Bulk mRNAseq



Low cost

We can process up to 20 bulk RNAseq samples for the same cost as 1 single cell RNAseq samples



Identify rare cell subpopulations

Rare cell subpopulations are lost in bulk RNAseq experimental settings, but may provide valuable information.



Lenient sample requirements

Sample requirements are less strict than for scRNAseq experiments, which gives you more flexibility to use all available samples.



Look into cellular states

Draw more informed conclusions from your transcriptomics data based on the state of each cell.



Increase cohort size

Easily include more samples in this experimental setting, to validate the data obtained in your scRNAseq research.



Get in touch with us to discuss your case