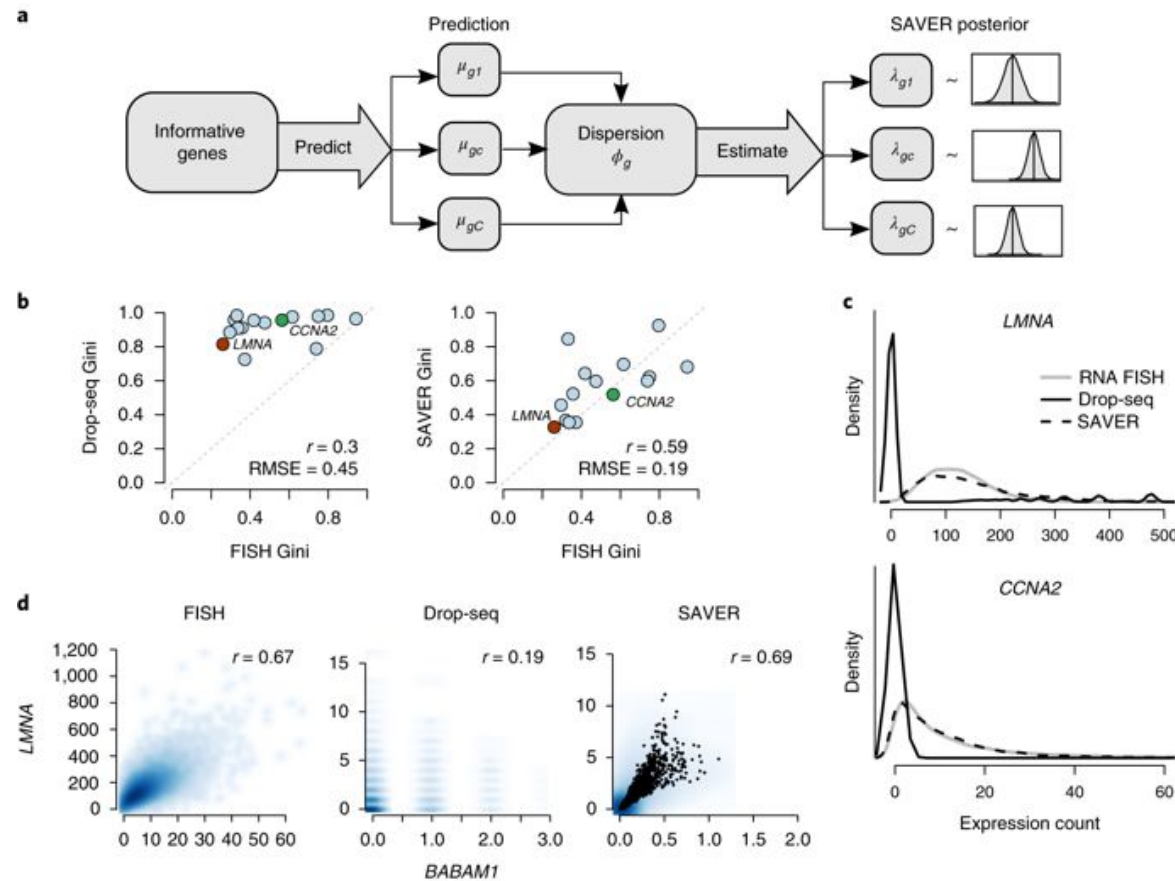


SAVERg: gene expression recovery for single-cell RNA sequencing

Yiqiu Tang

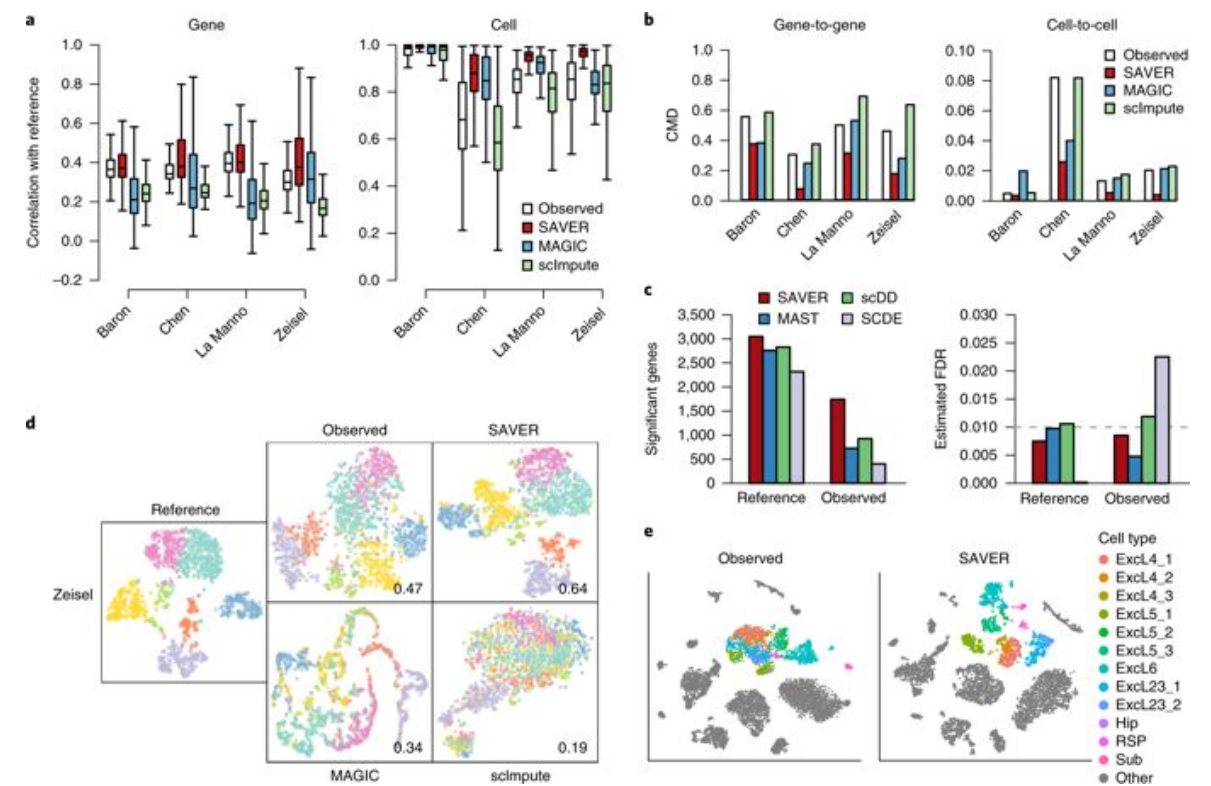


RNA FISH validation of SAVER results on Drop-seq data

What i will be doing...

cluster analysis, trajectory analysis,
differential expression analysis...

SAVER (Single-cell Analysis Via Expression Recovery) which is an expression recovery method for unique molecule index-based single-cell RNA sequencing data that borrows information across genes and cells to provide accurate expression estimates for all genes



Evaluation of SAVER by down-sampling and cell clustering.