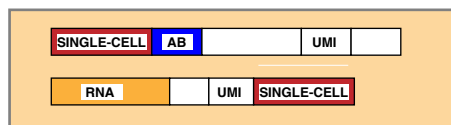
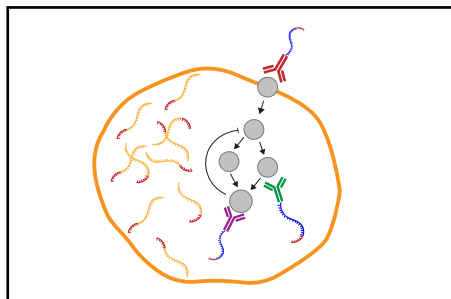
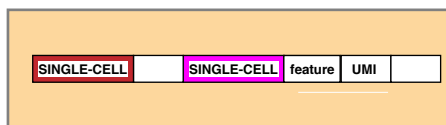
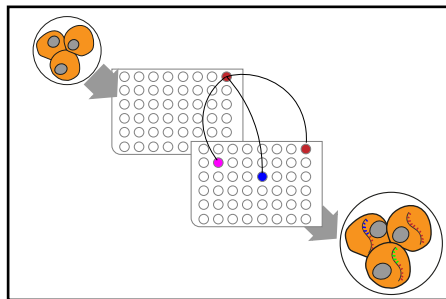


a

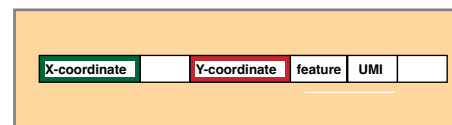
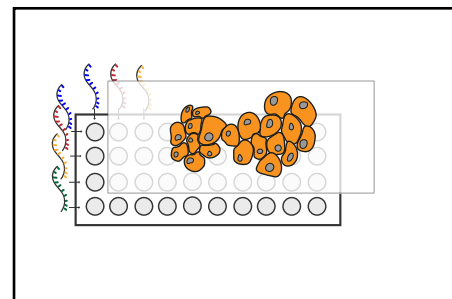
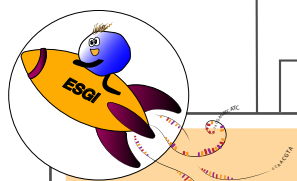
multi-modal single-cell data
(indices for RNA and protein)



combinatorial indexing data
(array of indices for single-cells)

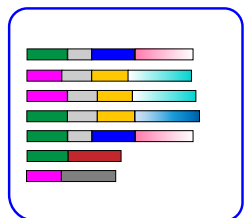


spatial single-cell data
(indices for x/y-coordinates)

**b**

Input

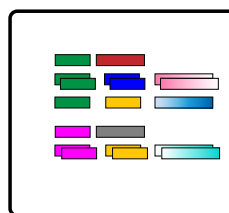
FASTQ



demultiplex
debarcode fastq-reads

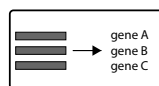


count
single-cell features (collapse UMIs)



STAR

map RNA-sequence to reference genome



Output

features

