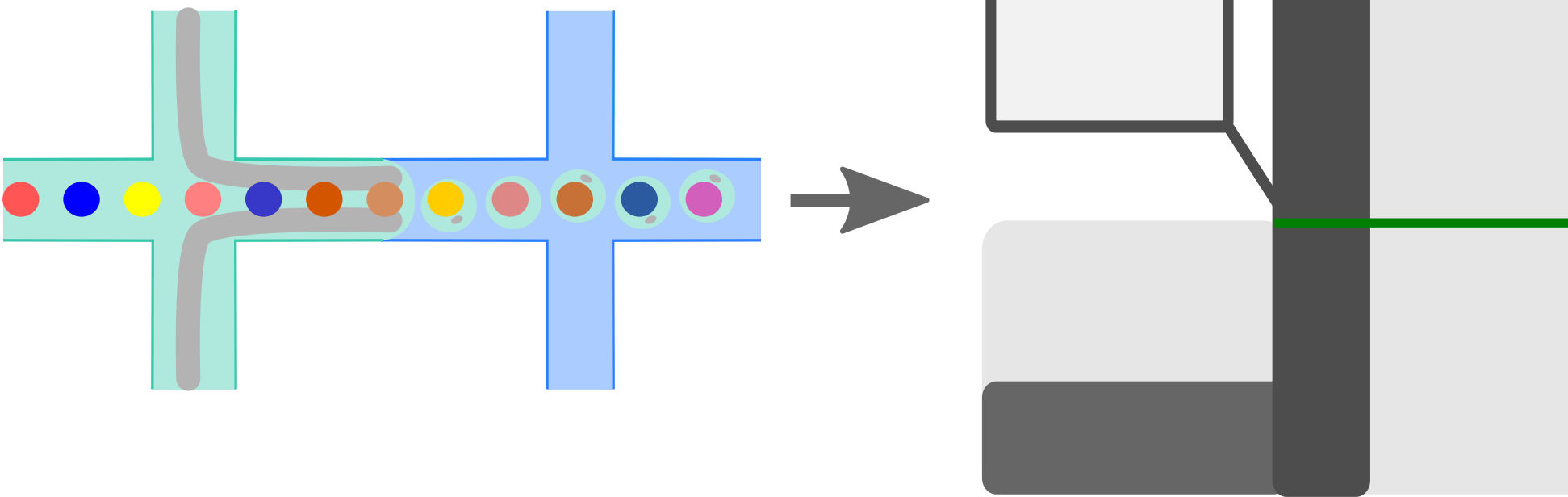


SEQUENCING

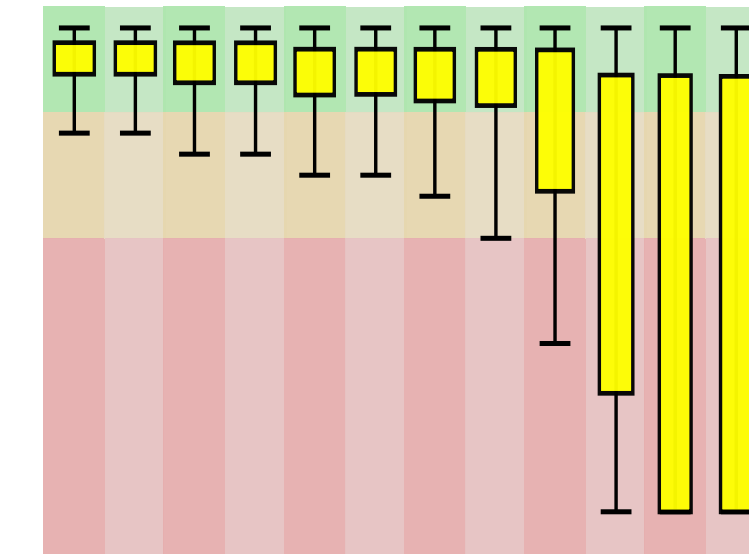


- 10x genomics

INPUT FASTQ



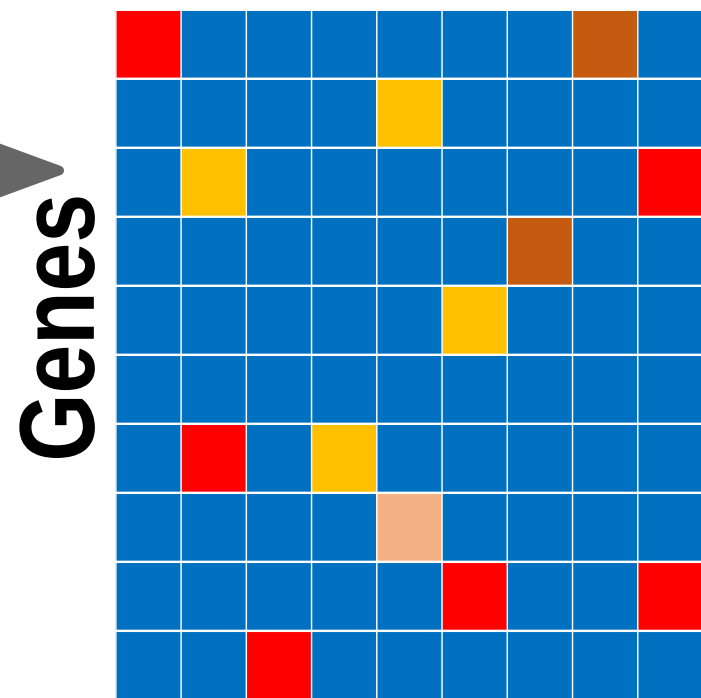
Preprocessing



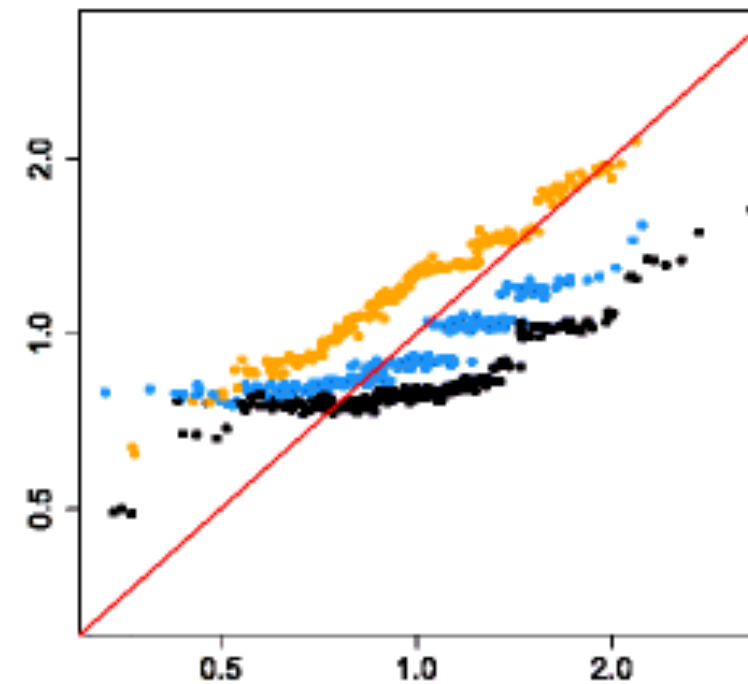
- STARSOLO
- Cell Ranger

raw Count Matrix

cells



Normalization



- STARSOLO
- Cell Ranger