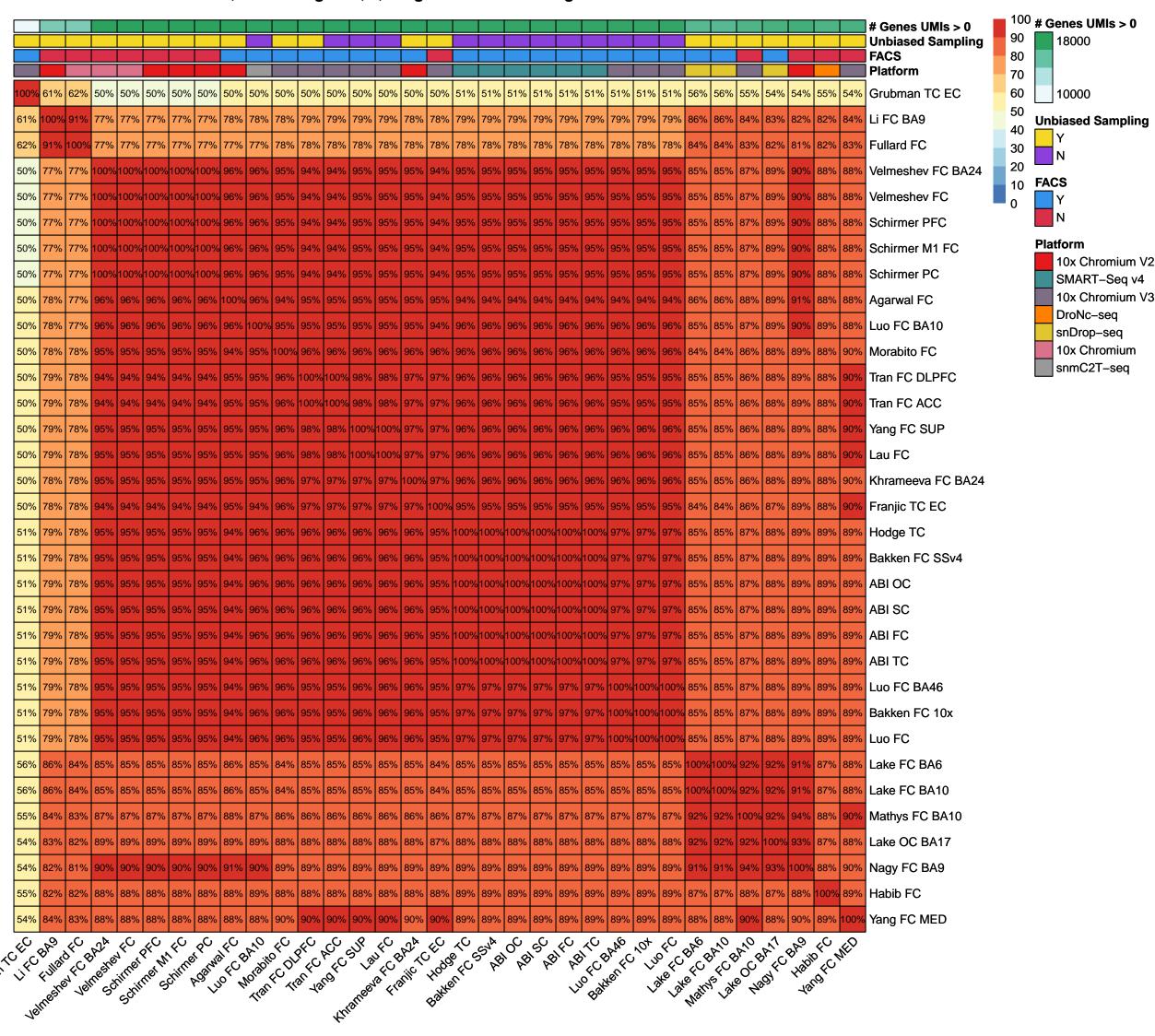
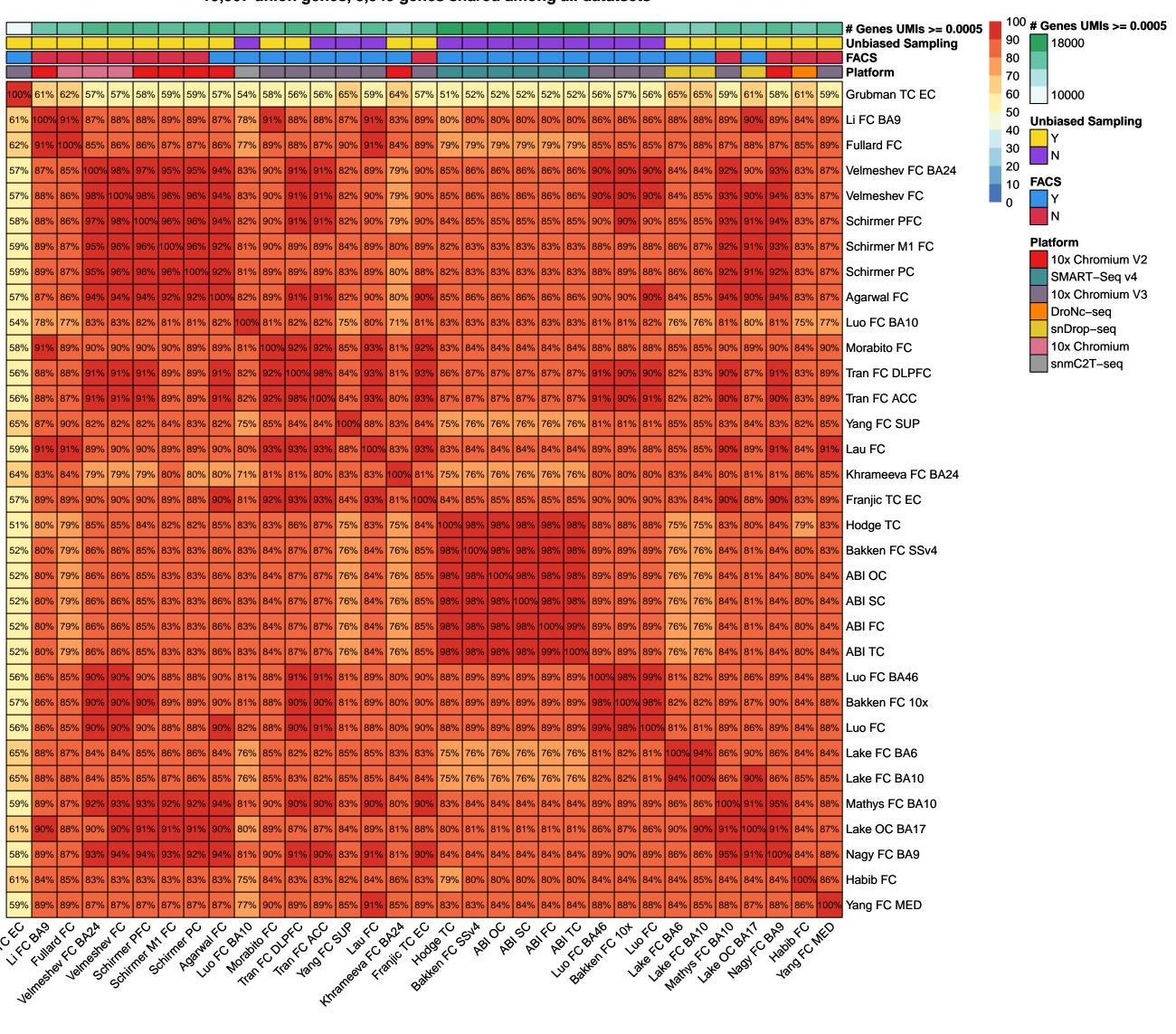
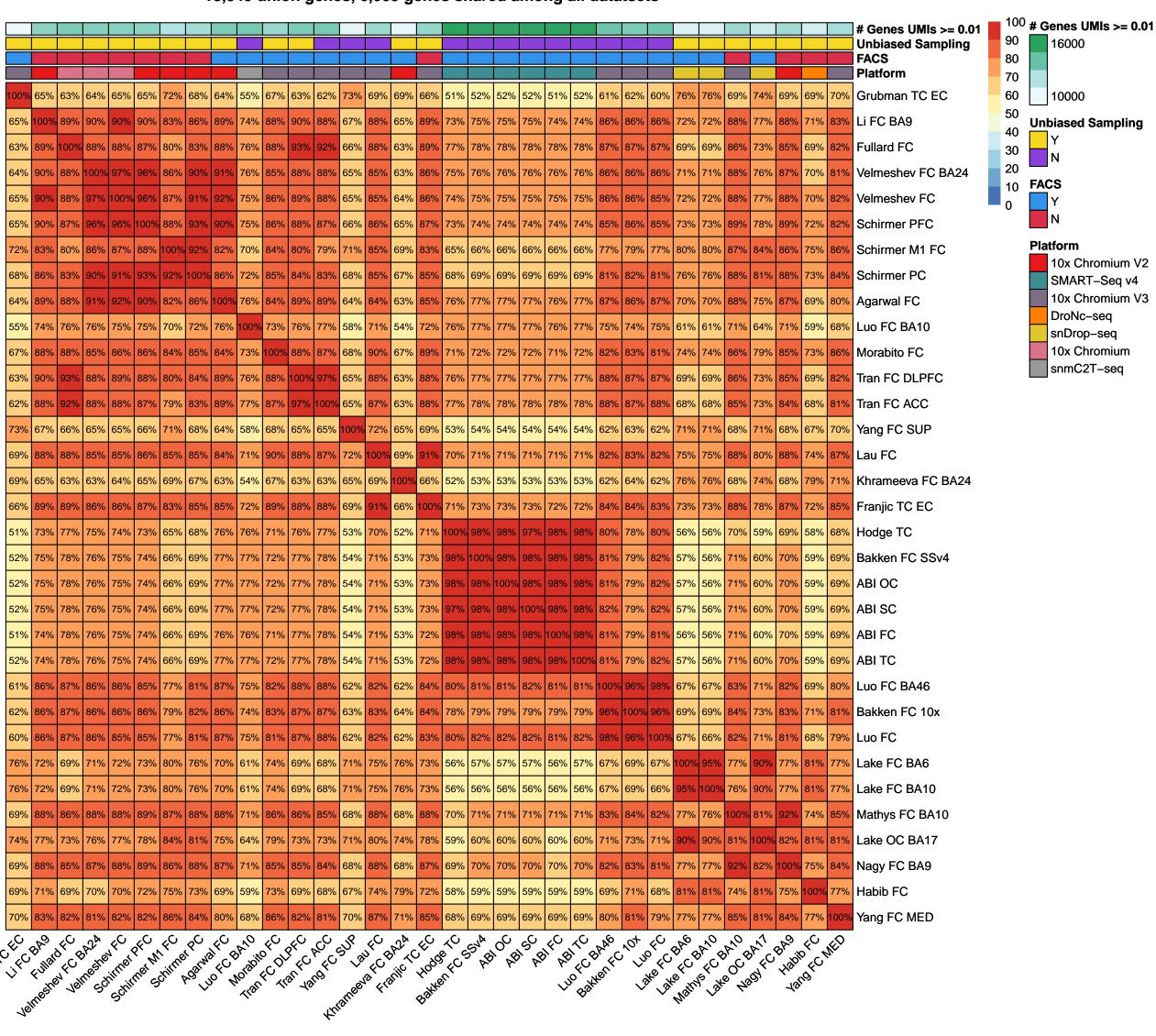
Feature Overlap Percent Restricting to Protein Coding Genes with Mean UMIs > 0 19,723 union genes, 9,147 genes shared among all datatsets



Feature Overlap Percent Restricting to Protein Coding Genes with Mean UMIs >= 0.0005 19,507 union genes, 8,549 genes shared among all datatsets



Feature Overlap Percent Restricting to Protein Coding Genes with Mean UMIs >= 0.01 18,849 union genes, 6,009 genes shared among all datatsets



Feature Overlap Percent Restricting to Protein Coding Genes with Mean UMIs >= 0.1 17,551 union genes, 364 genes shared among all datatsets

15000

5000

FACS

N

Platform

Unbiased Sampling

10x Chromium V2

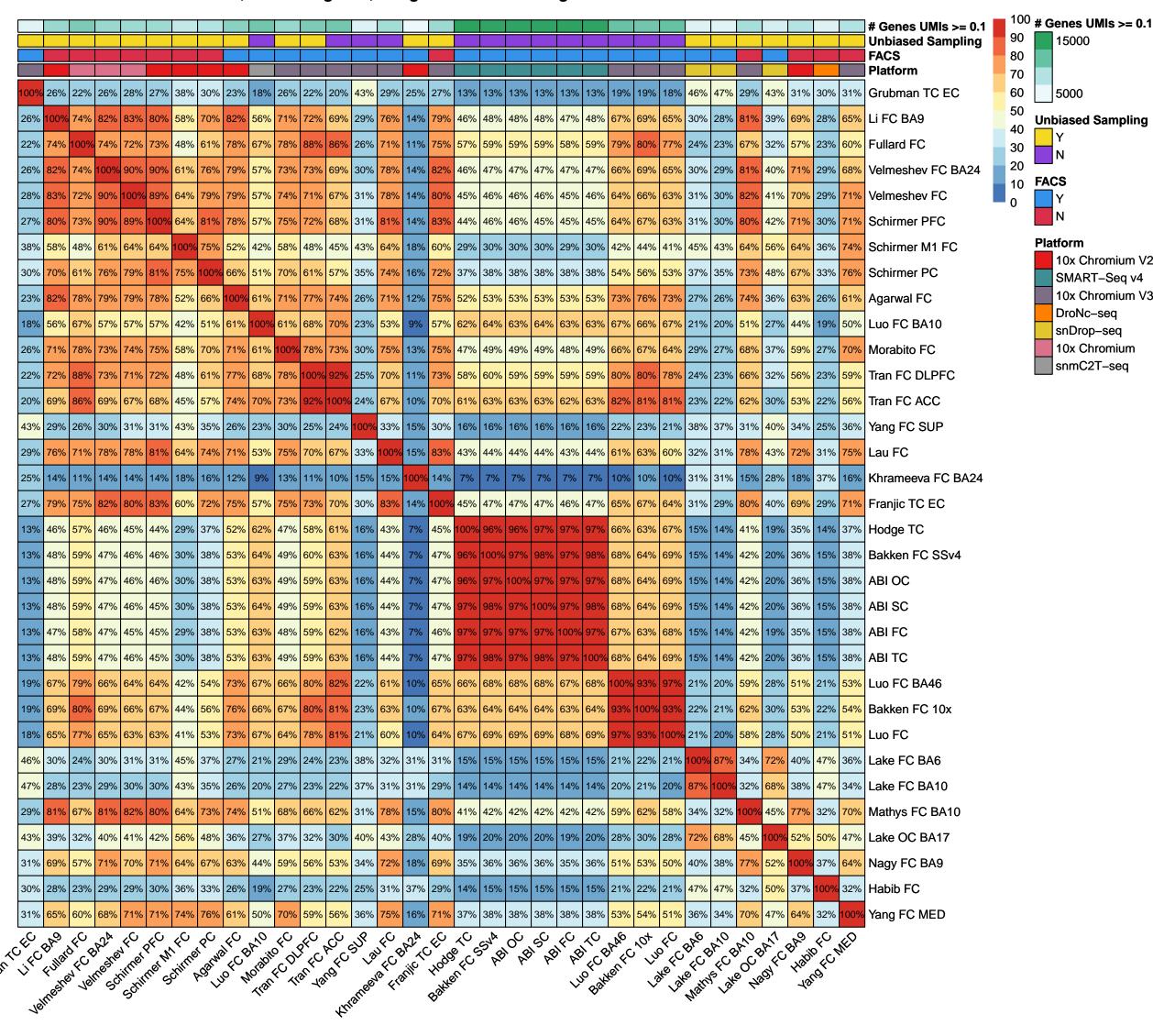
SMART-Seq v4 10x Chromium V3

DroNc-seq

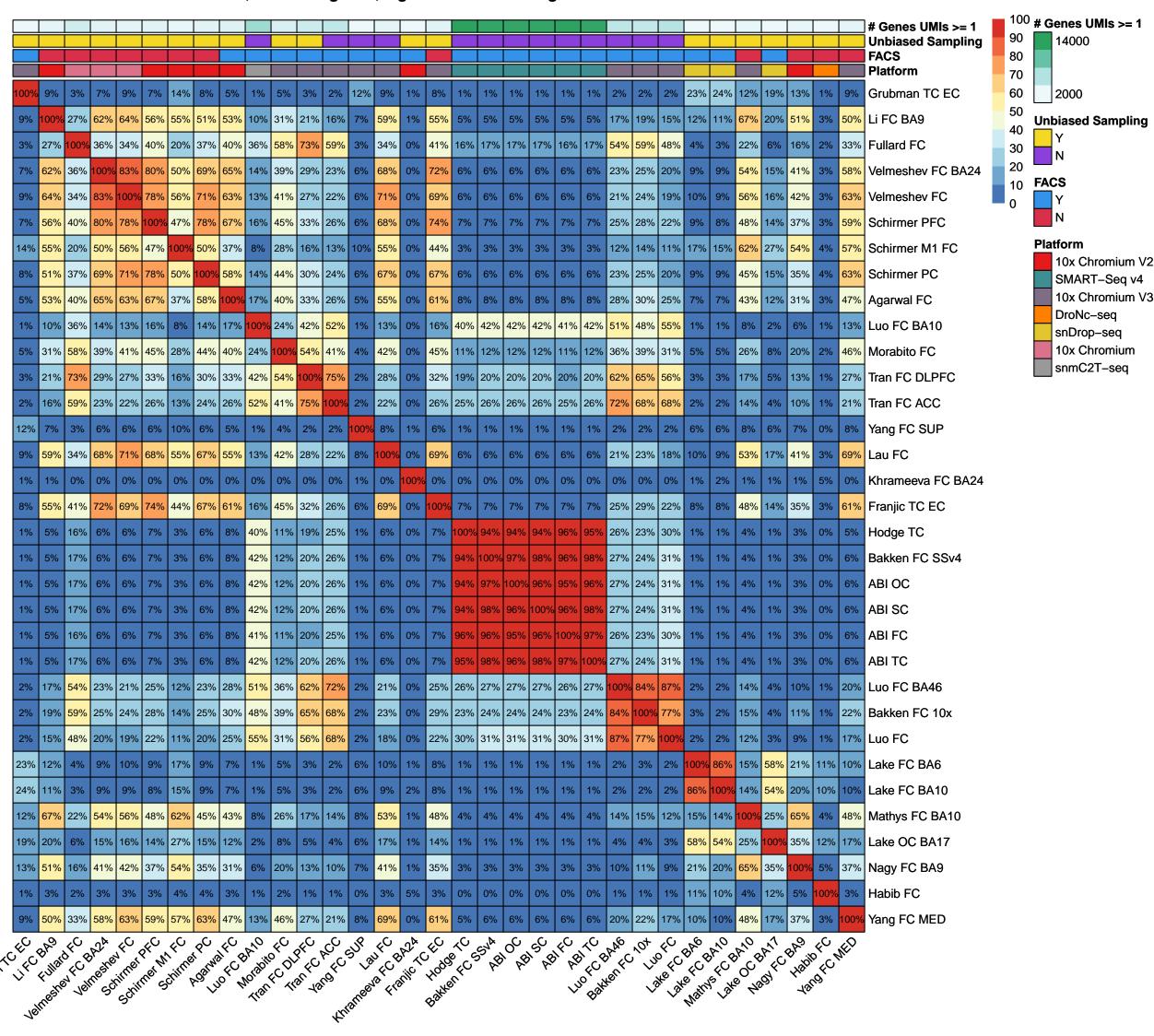
snDrop-seq

10x Chromium

snmC2T-seq



Feature Overlap Percent Restricting to Protein Coding Genes with Mean UMIs >= 1 15,102 union genes, 0 genes shared among all datatsets



Feature Overlap Percent Restricting to Protein Coding Genes with Mean UMIs >= 10 11,120 union genes, 0 genes shared among all datatsets

10000

FACS

Ν

Platform

Unbiased Sampling

10x Chromium V2

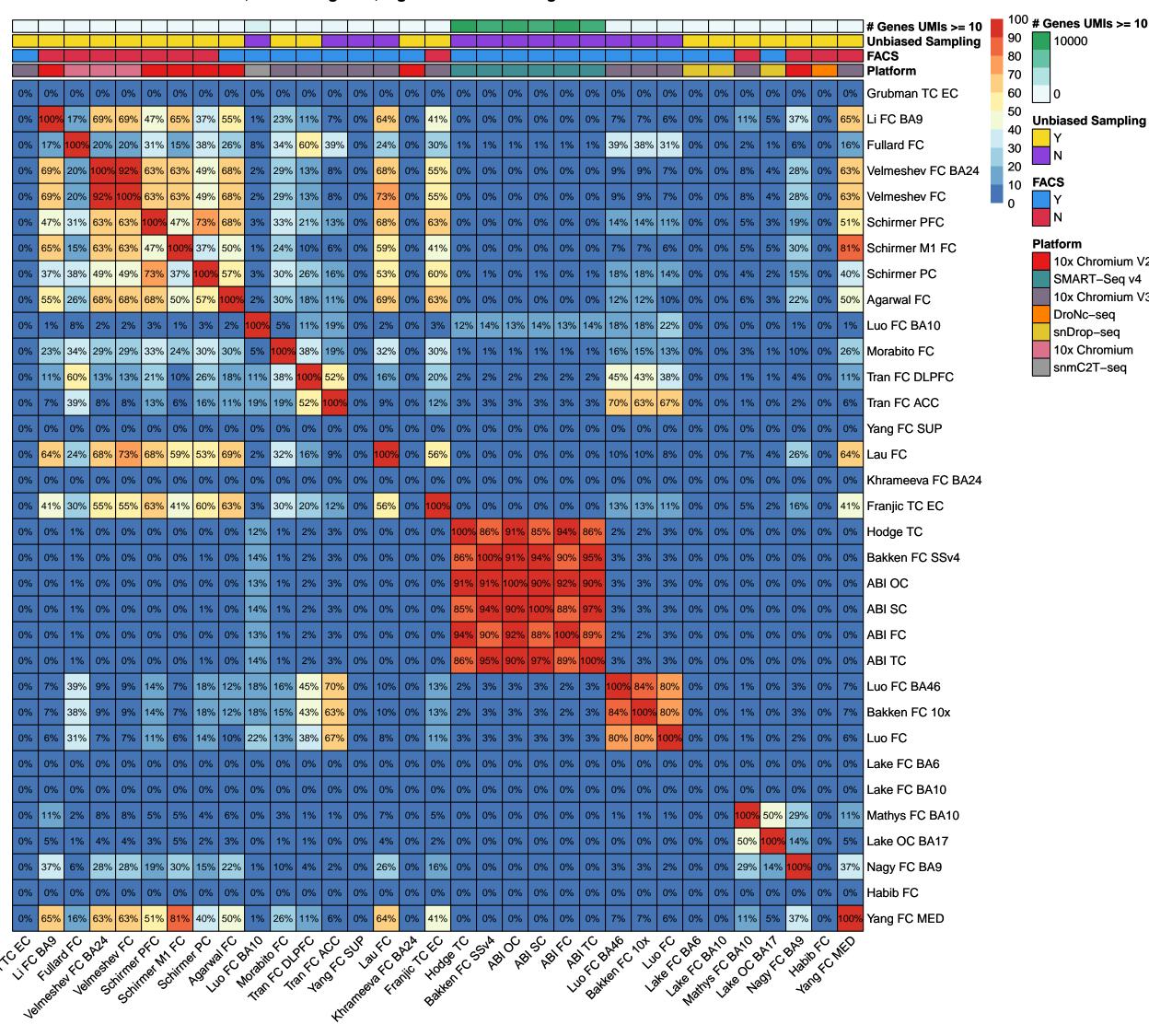
SMART-Seq v4 10x Chromium V3

DroNc-seq

snDrop-seq

10x Chromium

snmC2T-seq



Feature Overlap Percent Restricting to Protein Coding Genes with Mean UMIs >= 500 586 union genes, 0 genes shared among all datatsets

