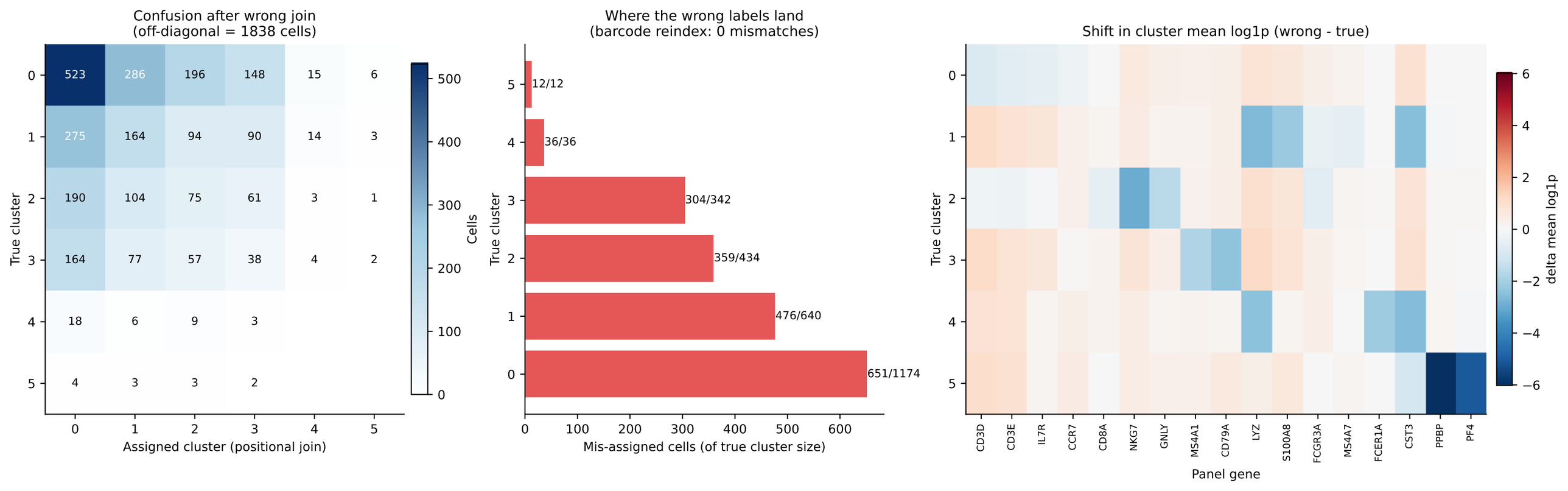
Confusion after wrong join
(off-diagonal = 1838 cells)
True cluster
Assigned cluster (positional join)
Cells
Where the wrong labels land
(barcode reindex: 0 mismatches)
True cluster
Mis-assigned cells (of true cluster size)
12/12
36/36
304/342
359/434
476/640
651/1174
Shift in cluster mean log1p (wrong - true)
True cluster
Panel gene
delta mean log1p
CD3D
CD3E
IL7R
CCR7
CD8A
NKG7
GNLY
MS4A1
CD79A
LYZ
S100A8
FCGR3A
MS4A7
FCER1A
CST3
PPBP
PF4