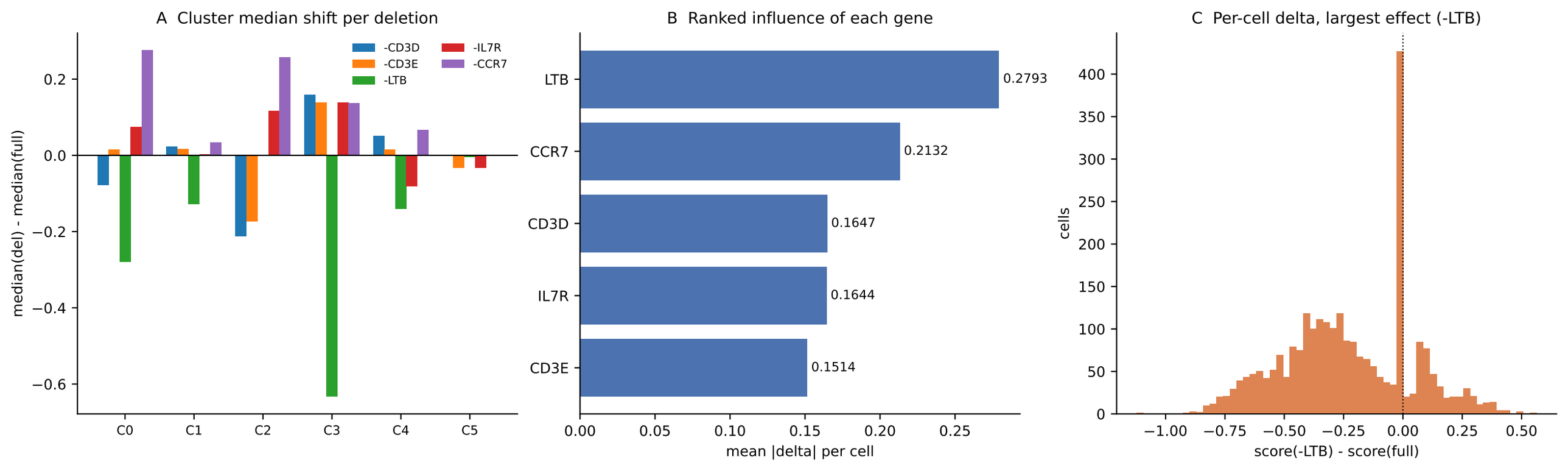
A Cluster median shift per deletion
-CD3D
-CD3E
-LTB
-IL7R
-CCR7
median(del) - median(full)
0.2
0.0
−0.2
−0.4
−0.6
C0
C1
C2
C3
C4
C5
B Ranked influence of each gene
LTB
CCR7
CD3D
IL7R
CD3E
0.2793
0.2132
0.1647
0.1644
0.1514
0.00
0.05
0.10
0.15
0.20
0.25
mean |delta| per cell
C Per-cell delta, largest effect (-LTB)
cells
400
350
300
250
200
150
100
50
0
−1.00
−0.75
−0.50
−0.25
0.00
0.25
0.50
score(-LTB) - score(full)