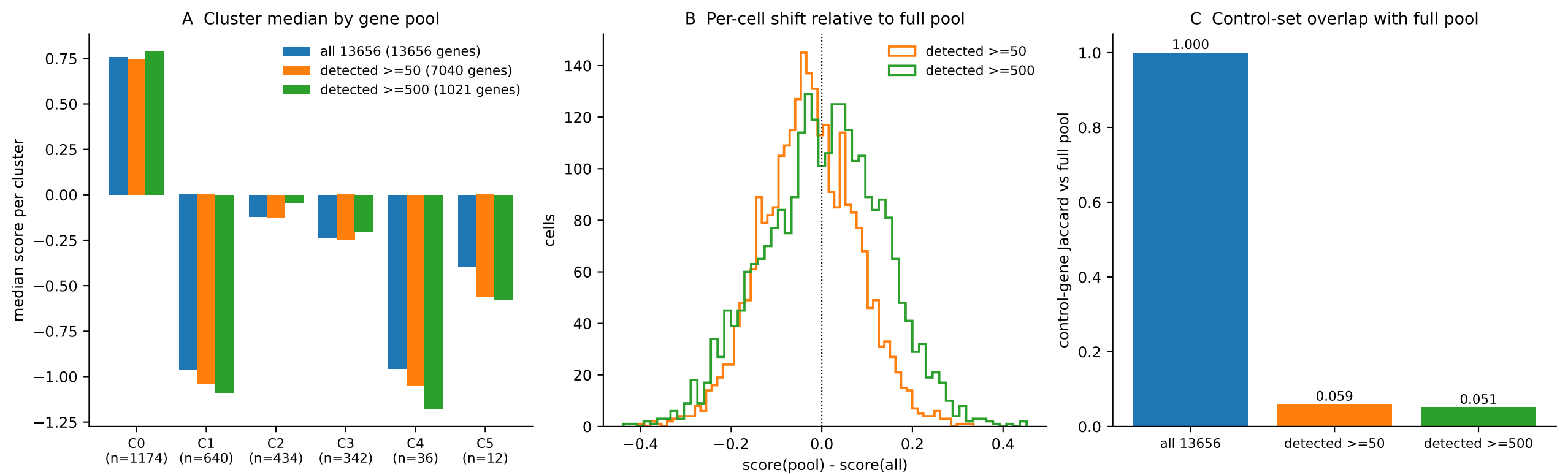
A Cluster median by gene pool
all 13656 (13656 genes)
detected >=50 (7040 genes)
detected >=500 (1021 genes)
median score per cluster
0.75
0.50
0.25
0.00
−0.25
−0.50
−0.75
−1.00
−1.25
C0 (n=1174)
C1 (n=640)
C2 (n=434)
C3 (n=342)
C4 (n=36)
C5 (n=12)
B Per-cell shift relative to full pool
detected >=50
detected >=500
cells
140
120
100
80
60
40
20
0
−0.4
−0.2
0.0
0.2
0.4
score(pool) - score(all)
C Control-set overlap with full pool
control-gene Jaccard vs full pool
1.0
0.8
0.6
0.4
0.2
0.0
1.000
0.059
0.051
all 13656
detected >=50
detected >=500