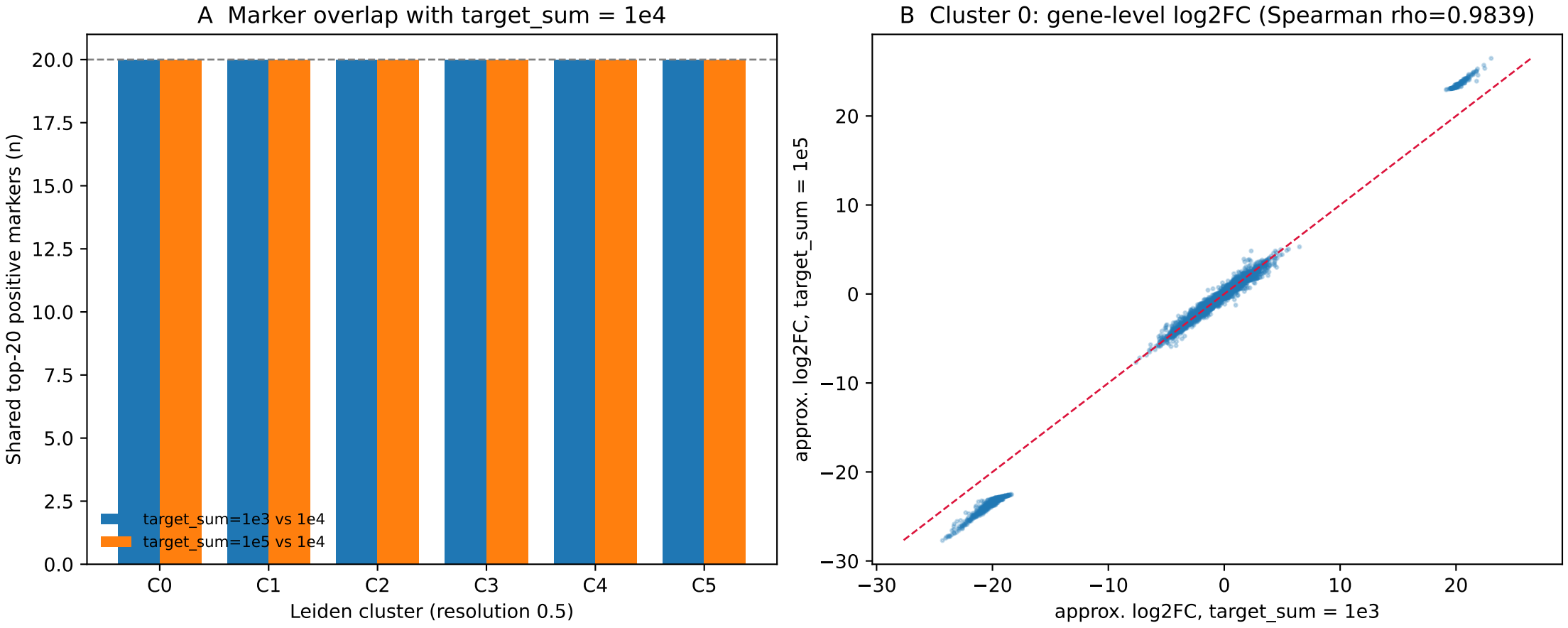

A Marker overlap with target_sum = 1e4
Shared top-20 positive markers (n)
20.0
17.5
15.0
12.5
10.0
7.5
5.0
2.5
0.0
target_sum=1e3 vs 1e4
target_sum=1e5 vs 1e4
C0
C1
C2
C3
C4
C5
Leiden cluster (resolution 0.5)
B Cluster 0: gene-level log2FC (Spearman rho=0.9839)
approx. log2FC, target_sum = 1e5
20
10
0
−10
−20
−30
−30
−20
−10
0
10
20
approx. log2FC, target_sum = 1e3