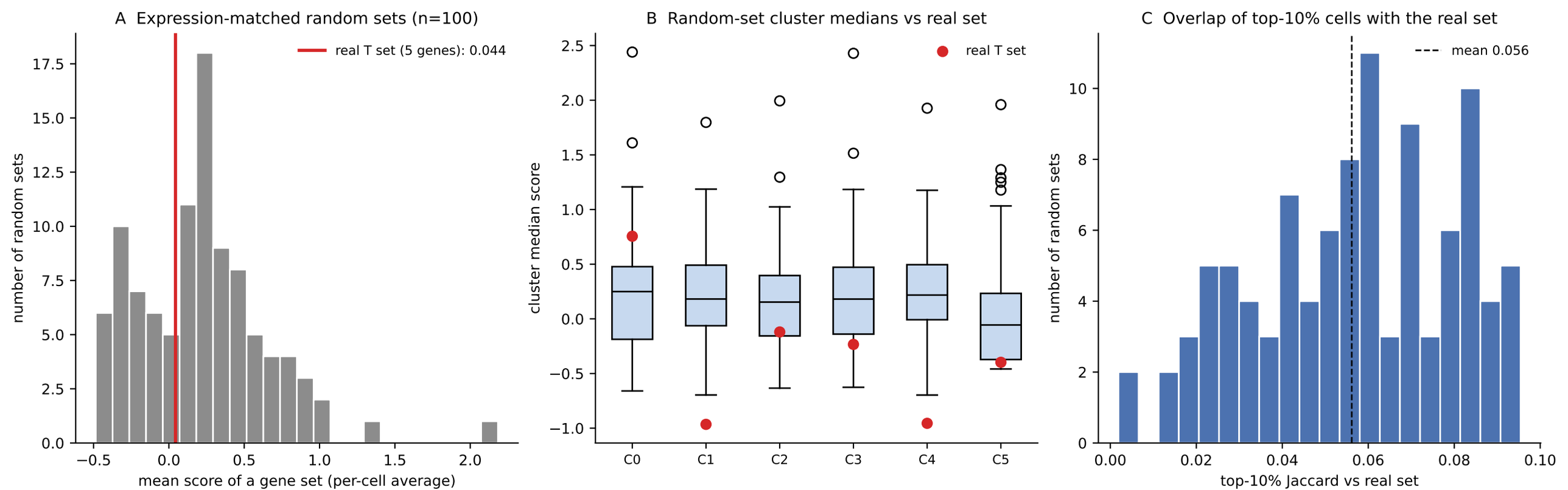
A Expression-matched random sets (n=100)
real T set (5 genes): 0.044
number of random sets
mean score of a gene set (per-cell average)
B Random-set cluster medians vs real set
real T set
cluster median score
C0
C1
C2
C3
C4
C5
C Overlap of top-10% cells with the real set
mean 0.056
number of random sets
top-10% Jaccard vs real set