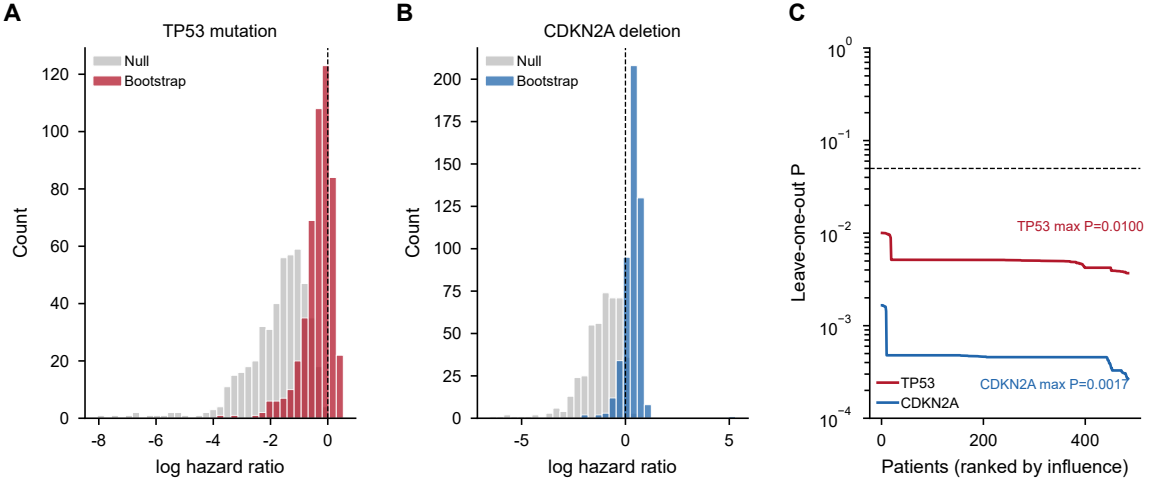

A
TP53 mutation
Null
Bootstrap
Count
log hazard ratio
B
CDKN2A deletion
Null
Bootstrap
Count
log hazard ratio
C
Leave-one-out P
TP53 max P=0.0100
TP53
CDKN2A
CDKN2A max P=0.0017
Patients (ranked by influence)