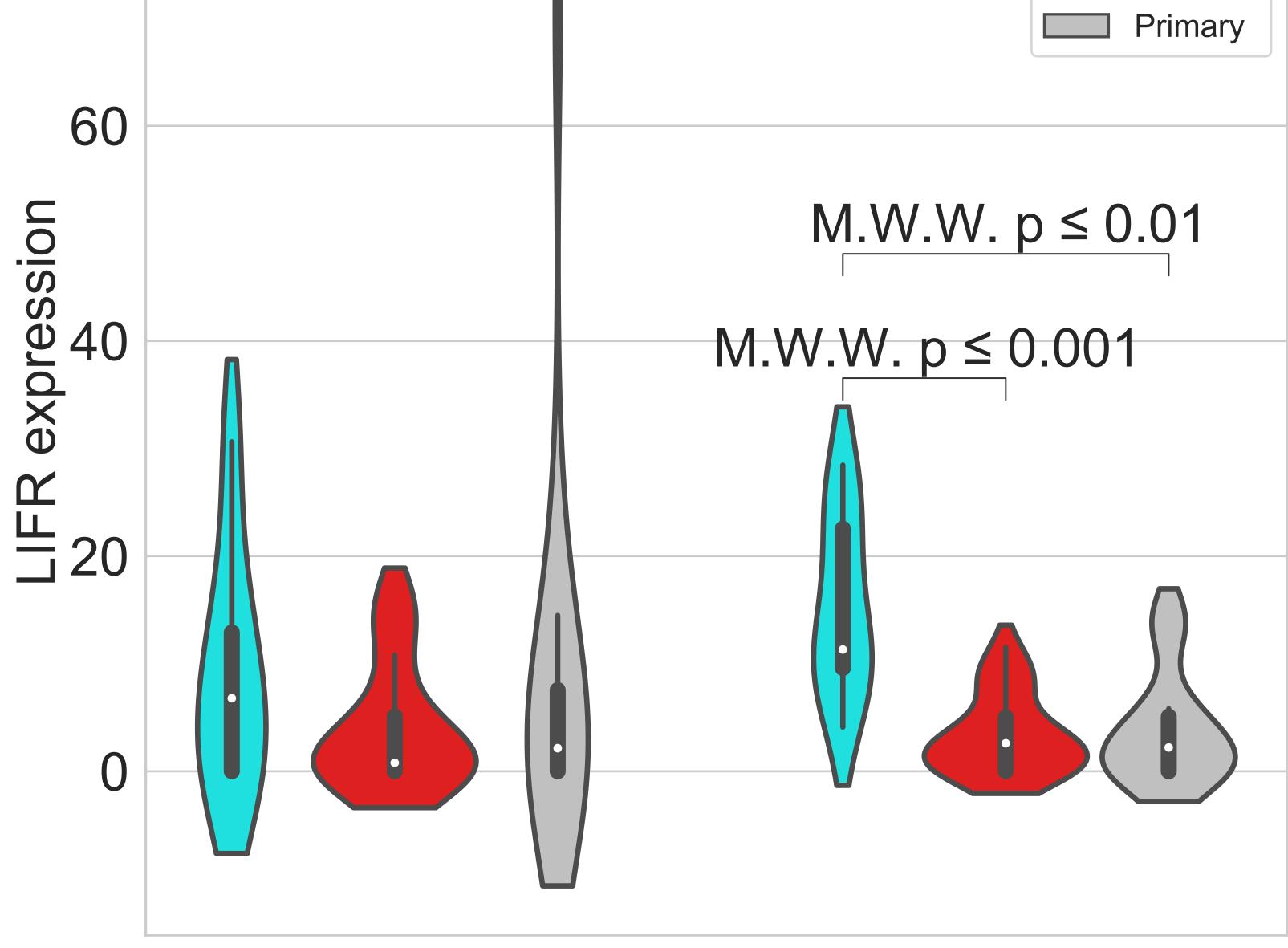
LIFR: Kruskal-Wallis p=0.012 Stage Normal CIS+AIS Primary M.W.W. p ≤ 0.01 M.W.W. $p \le 0.001$

80



Higher Lower Mutation Shared Count per TMB