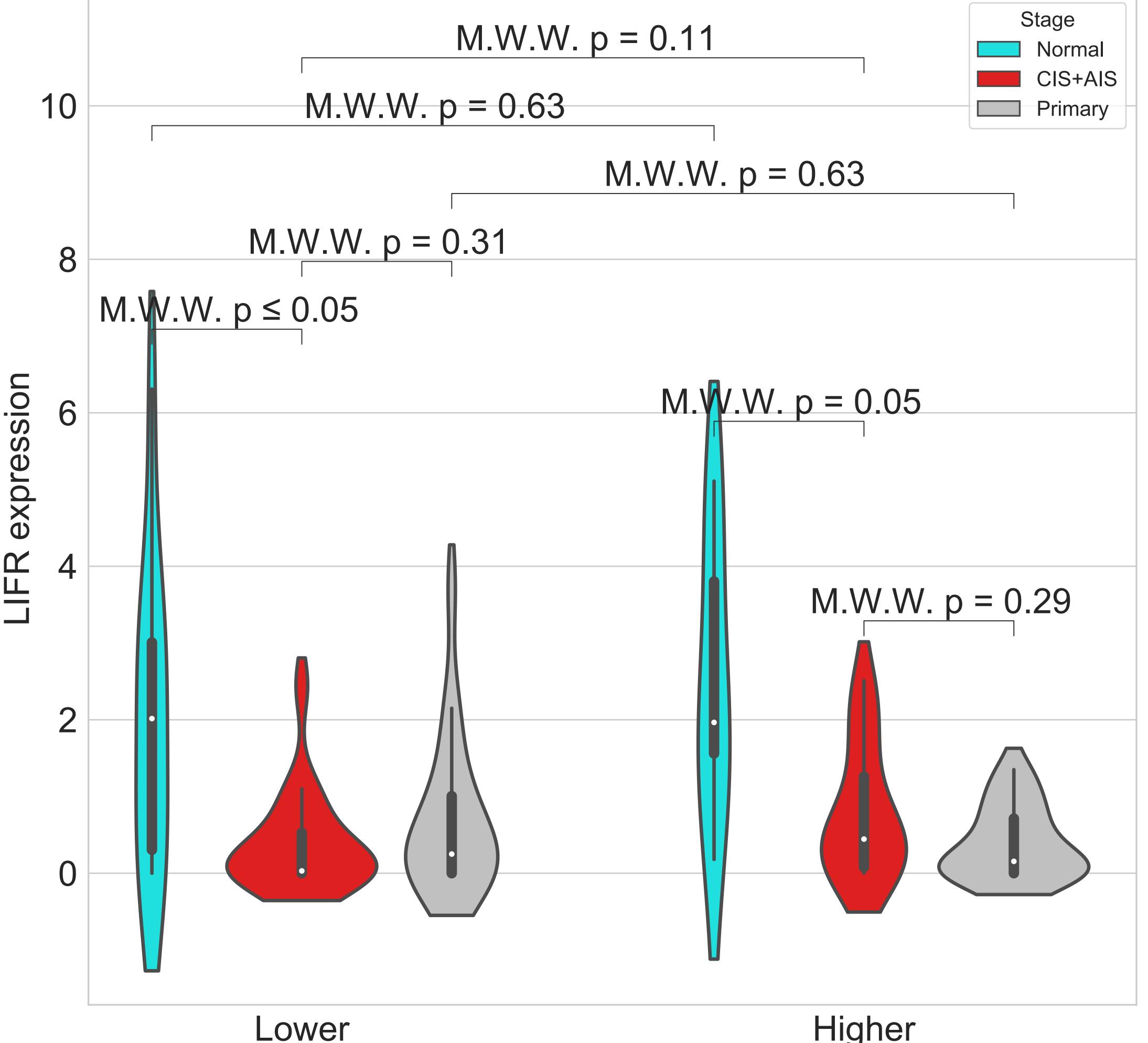
LIFR: Kruskal-Wallis p=0.008

M.W.W. p = 0.11



er
Mutation Shared Count per TMB