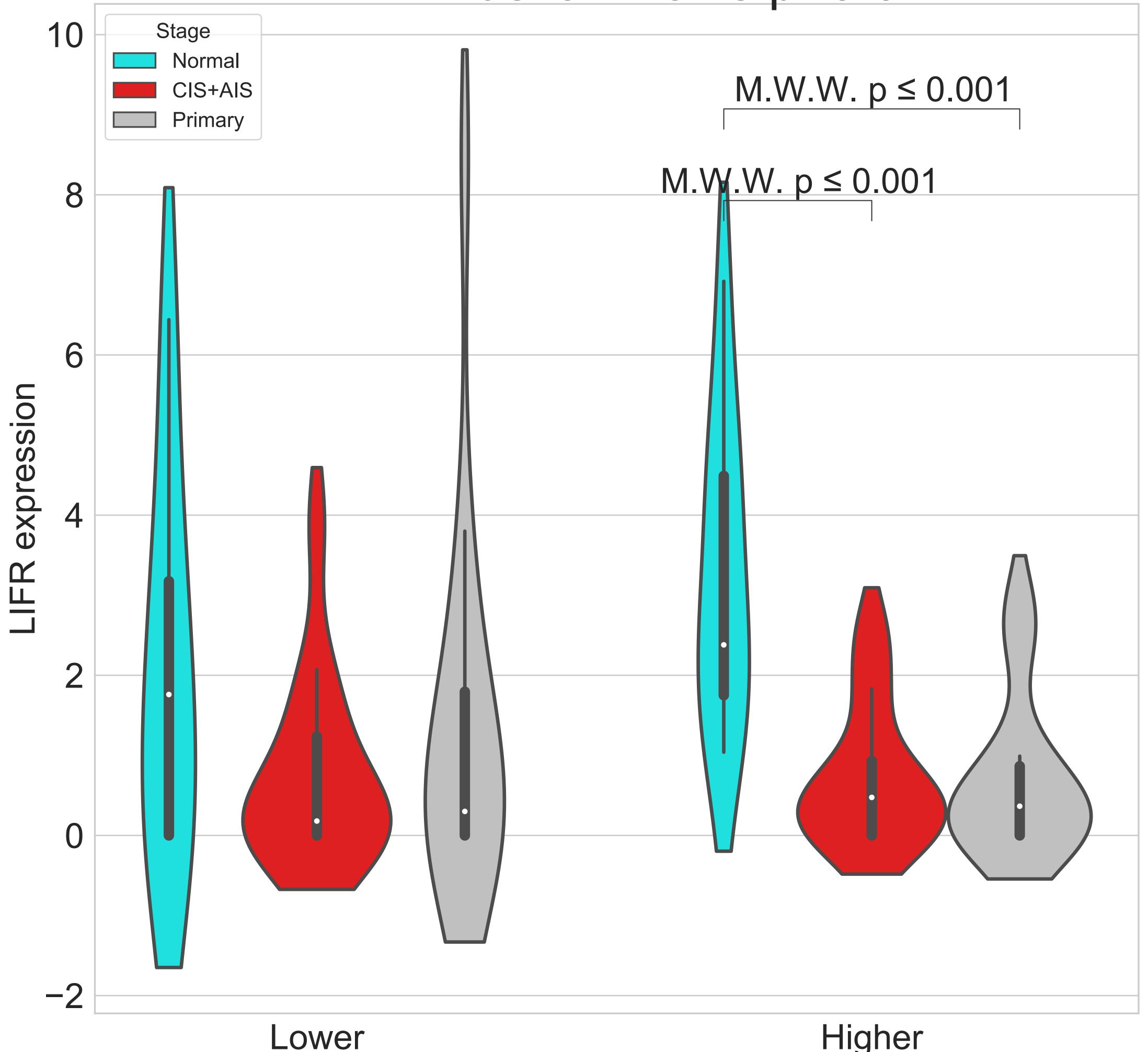
LIFR: Kruskal-Wallis p=0.011



ver
Mutation Shared Count per TMB