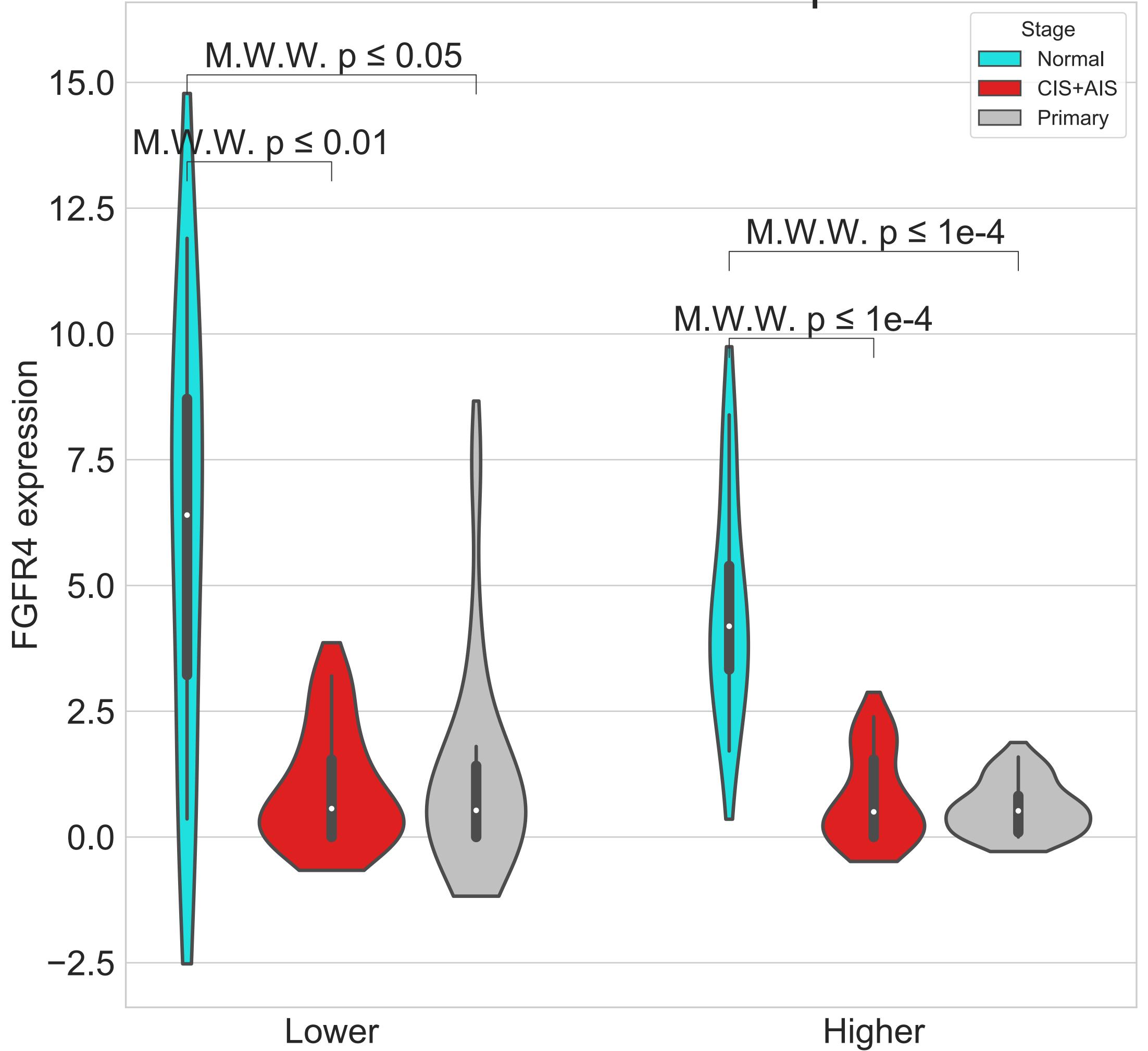
FGFR4: Kruskal-Wallis p=0.000



Mutation Shared Count per TMB