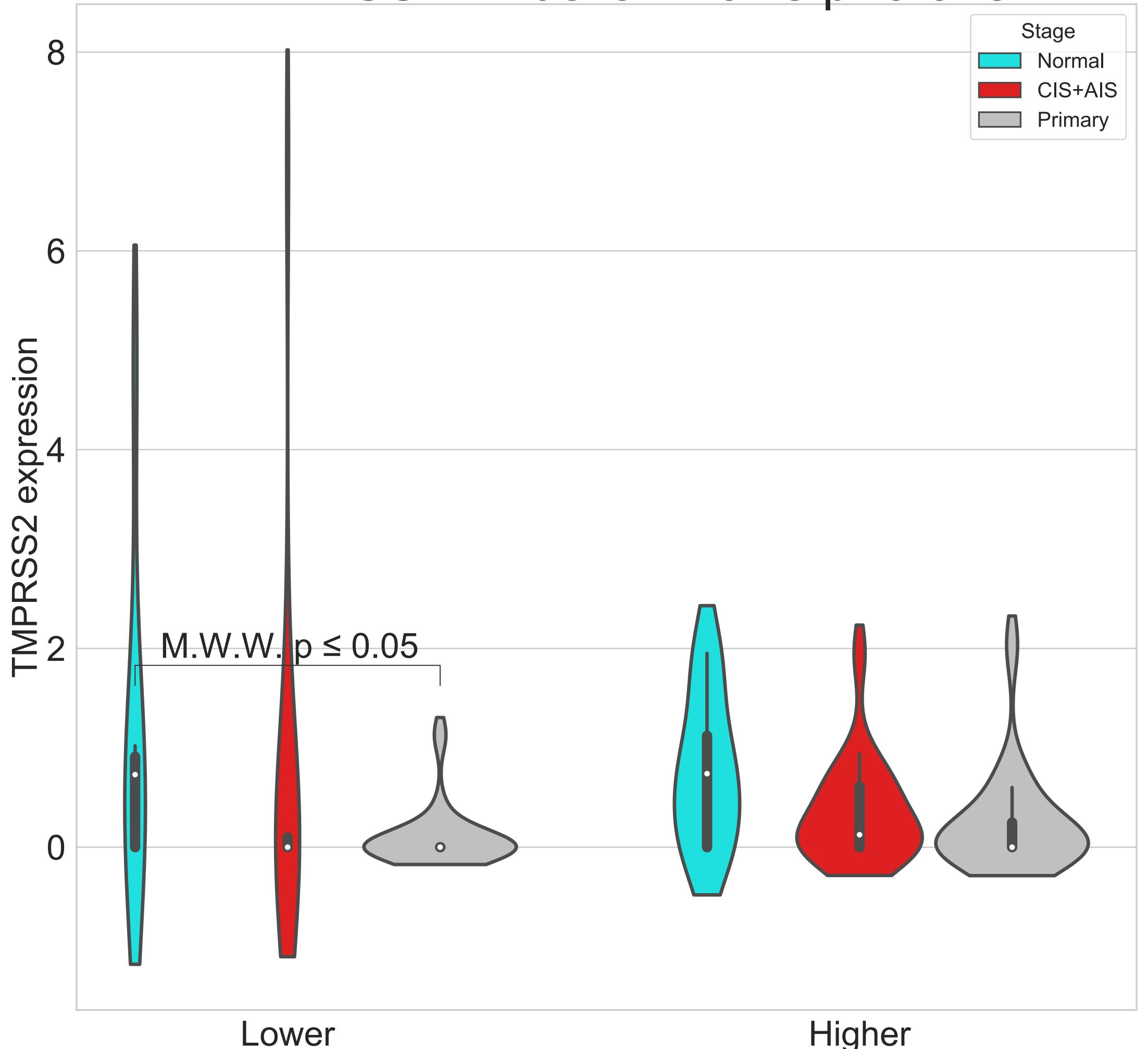
TMPRSS2: Kruskal-Wallis p=0.045



Higher Mutation Shared Count per TMB