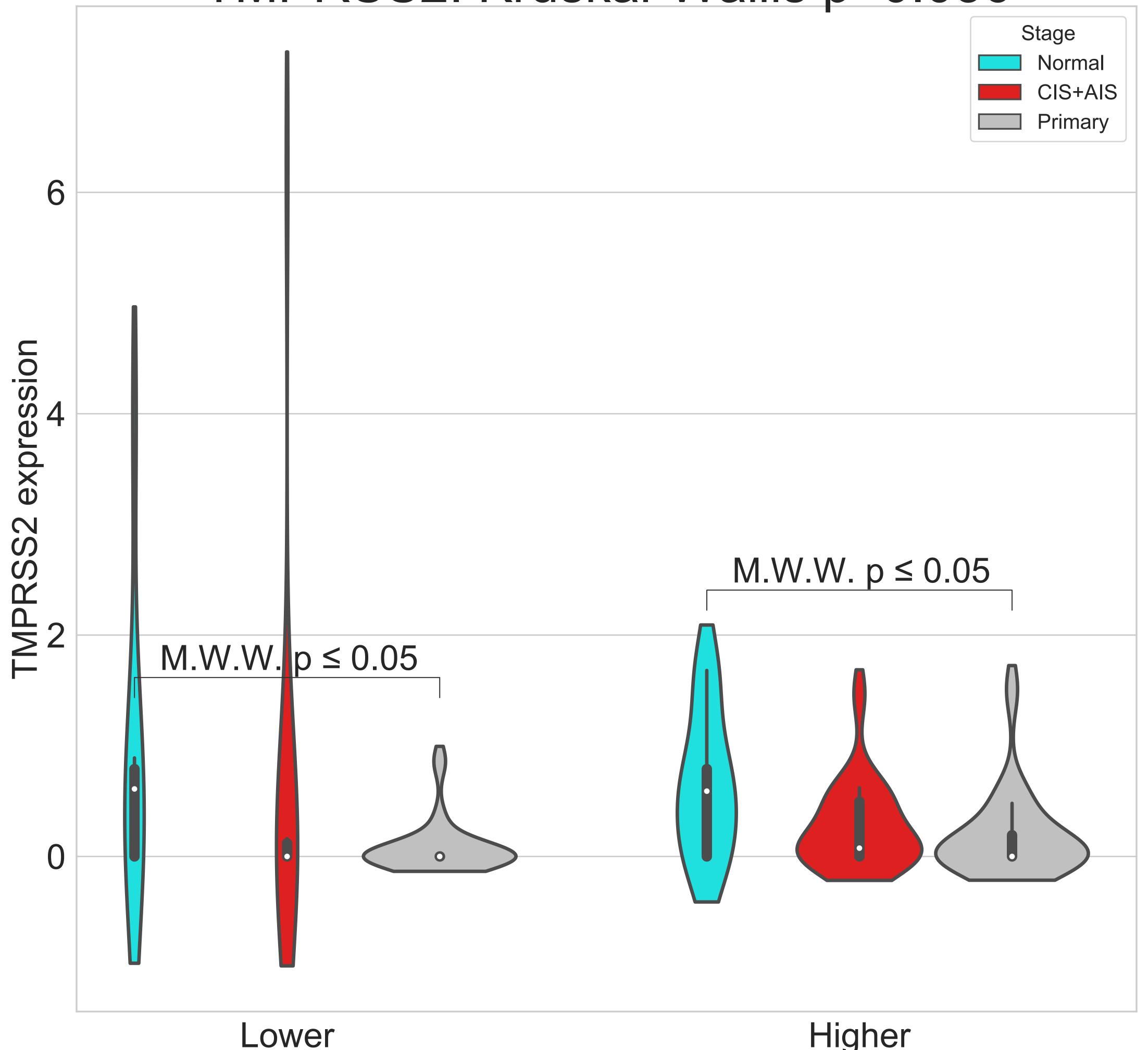
TMPRSS2: Kruskal-Wallis p=0.036



Higher Mutation Shared Count per TMB