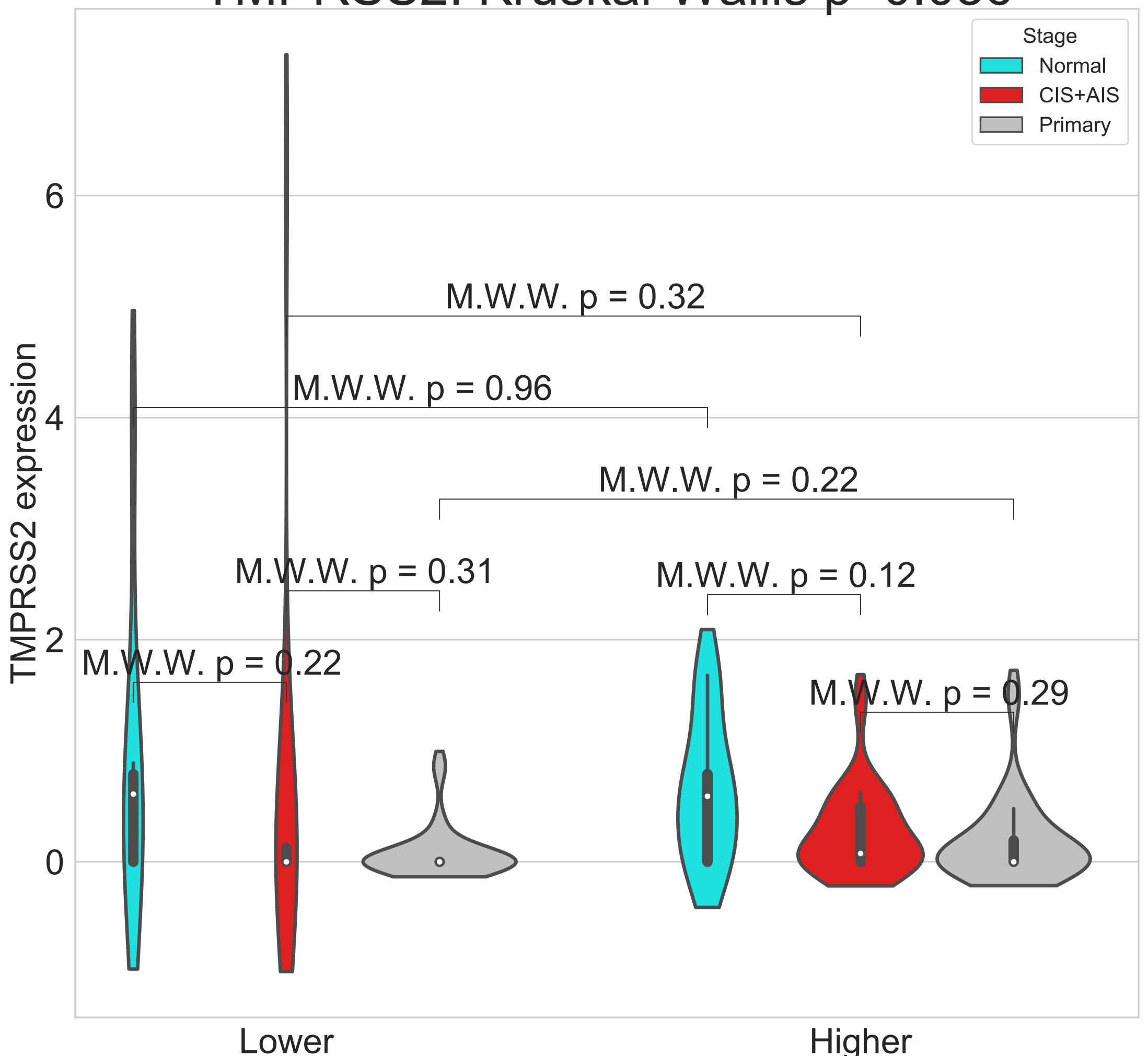
TMPRSS2: Kruskal-Wallis p=0.036



r Higher Mutation Shared Count per TMB