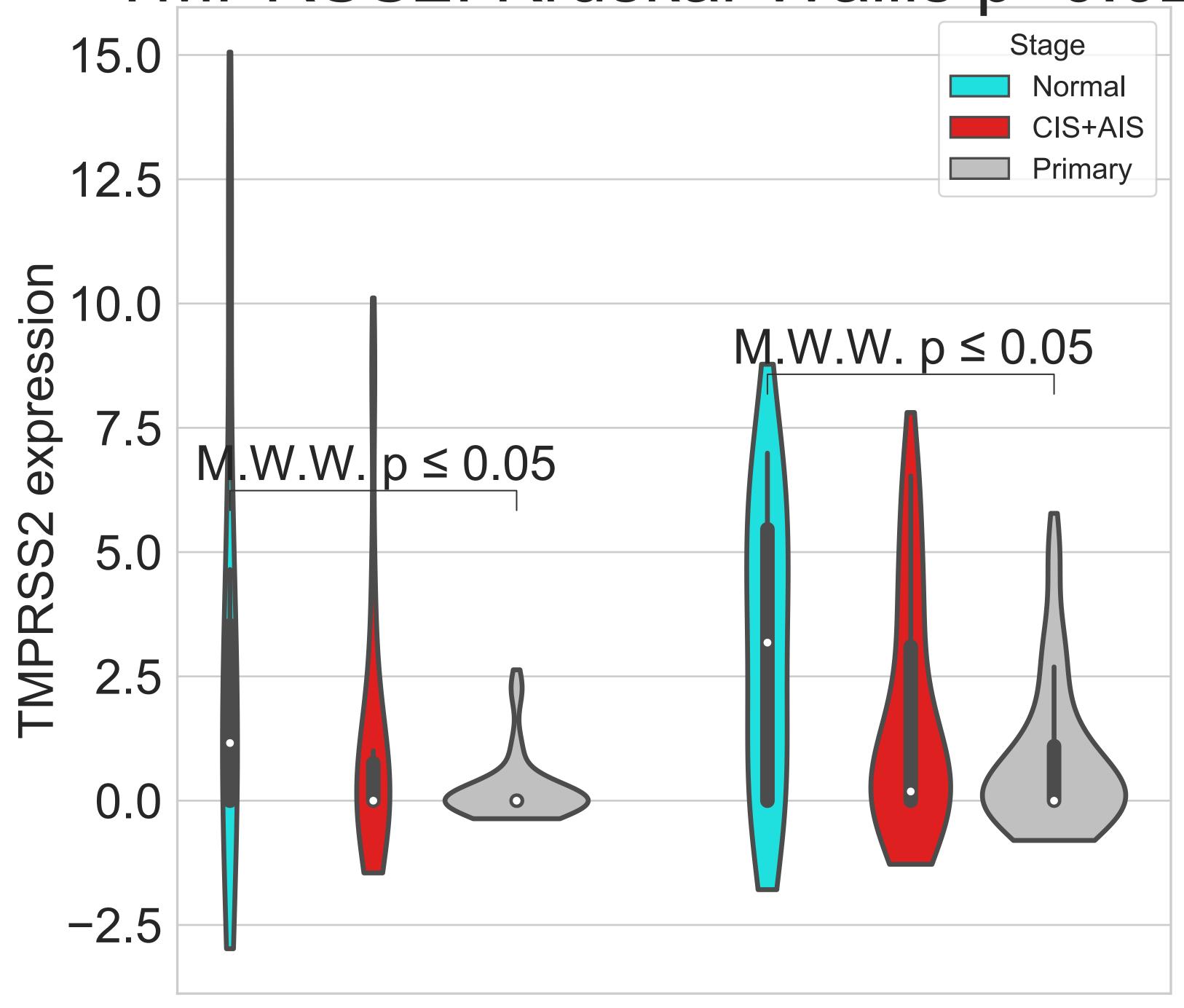
TMPRSS2: Kruskal-Wallis p=0.026



Lower Higher Mutation Shared Count per TMB