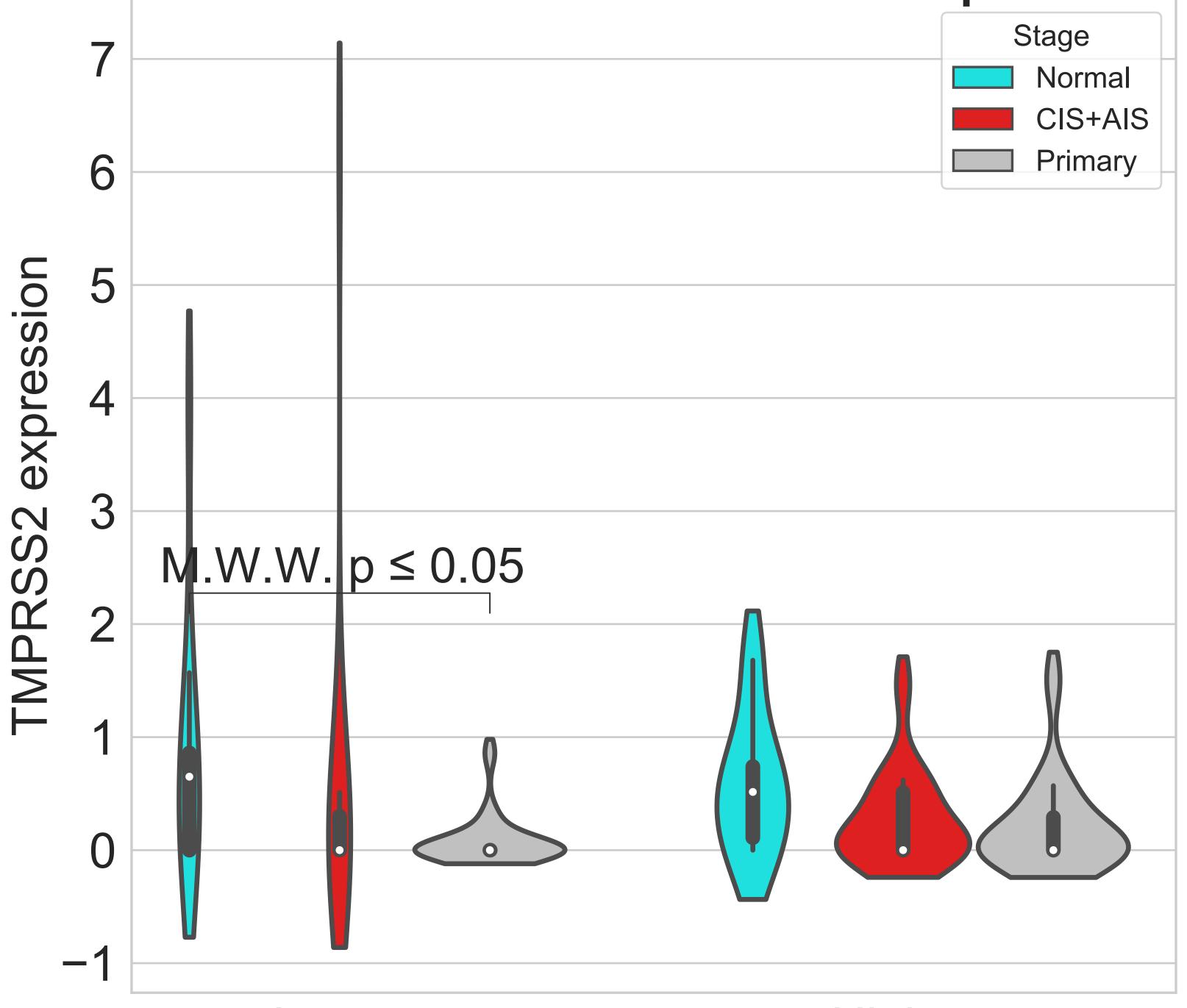
TMPRSS2: Kruskal-Wallis p=0.047



Lower Higher Mutation Shared Count (SYN) per TMB