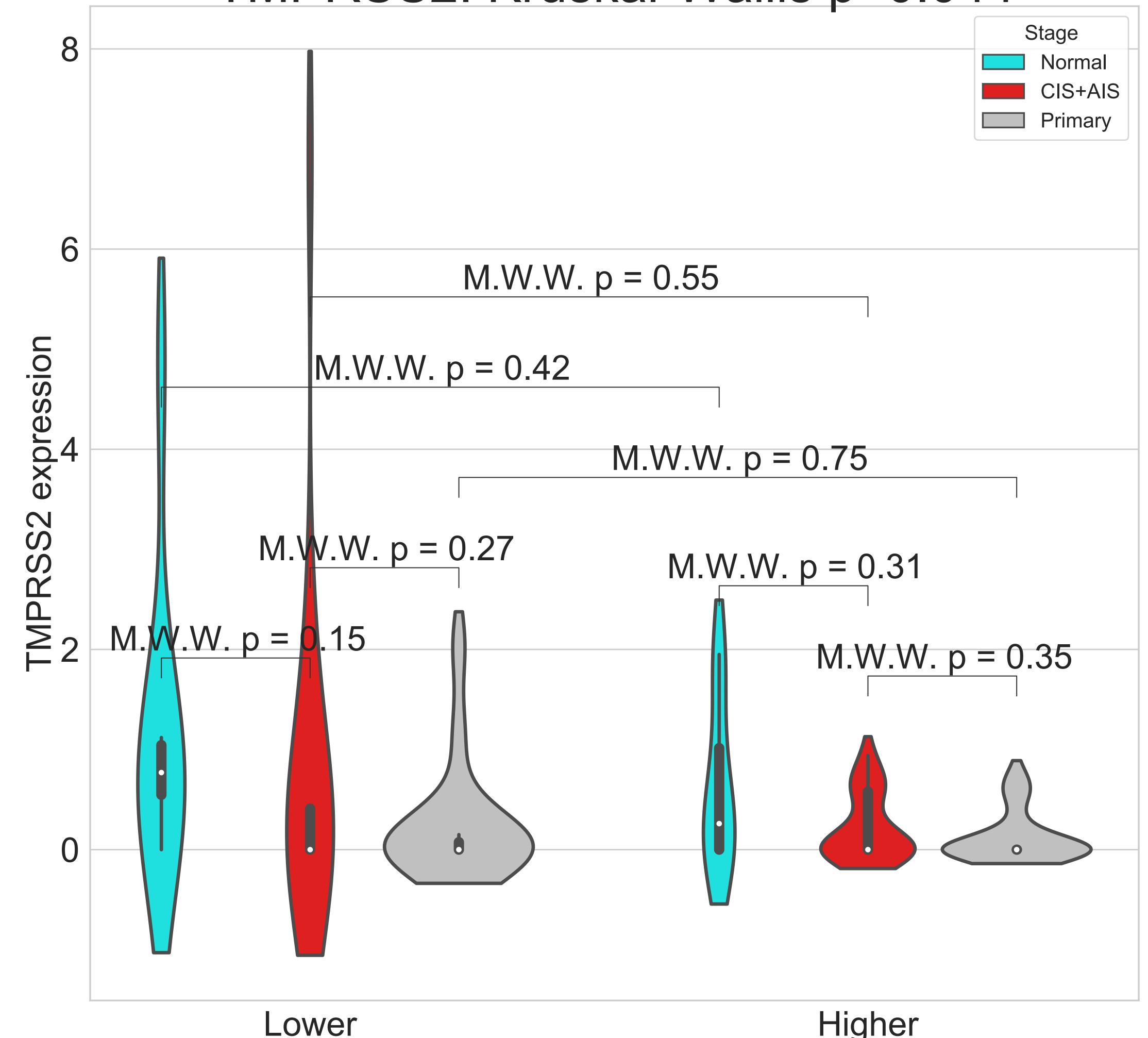
TMPRSS2: Kruskal-Wallis p=0.044



Higher Mutation Shared Count (SYN)