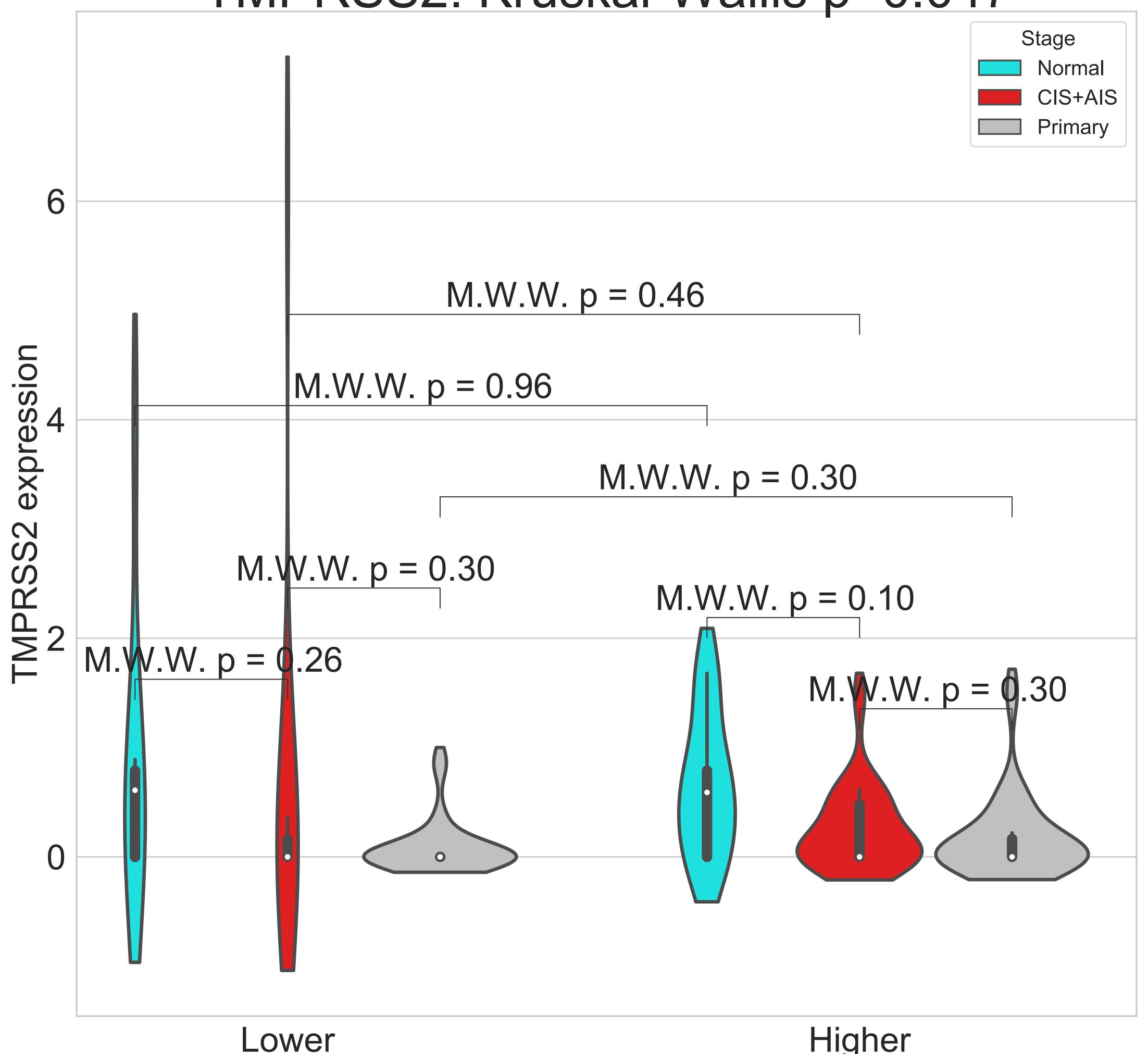
TMPRSS2: Kruskal-Wallis p=0.047



Higher Mutation Shared Proportion (SYN)