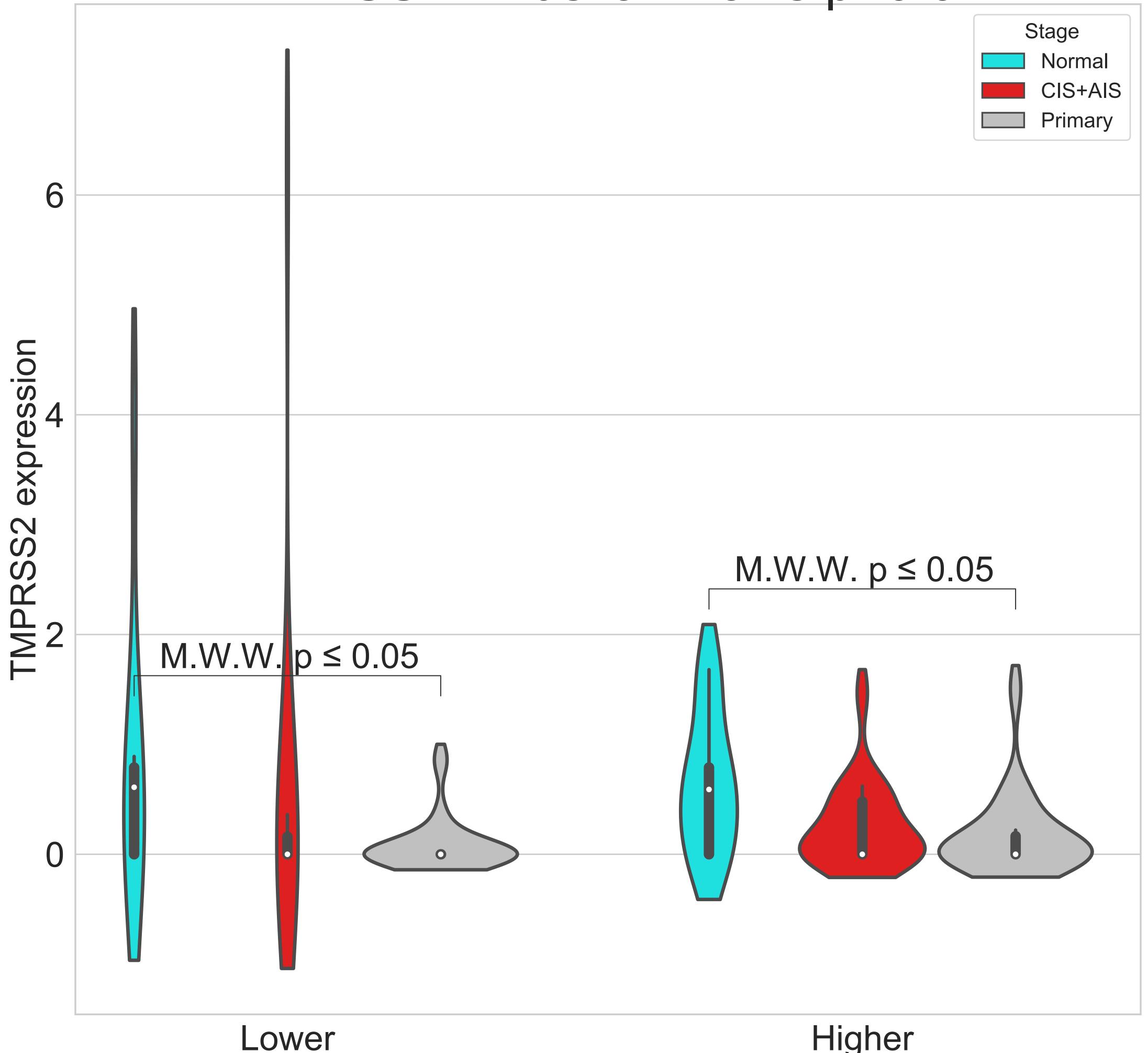
TMPRSS2: Kruskal-Wallis p=0.047



er
Mutation Shared Proportion (SYN)