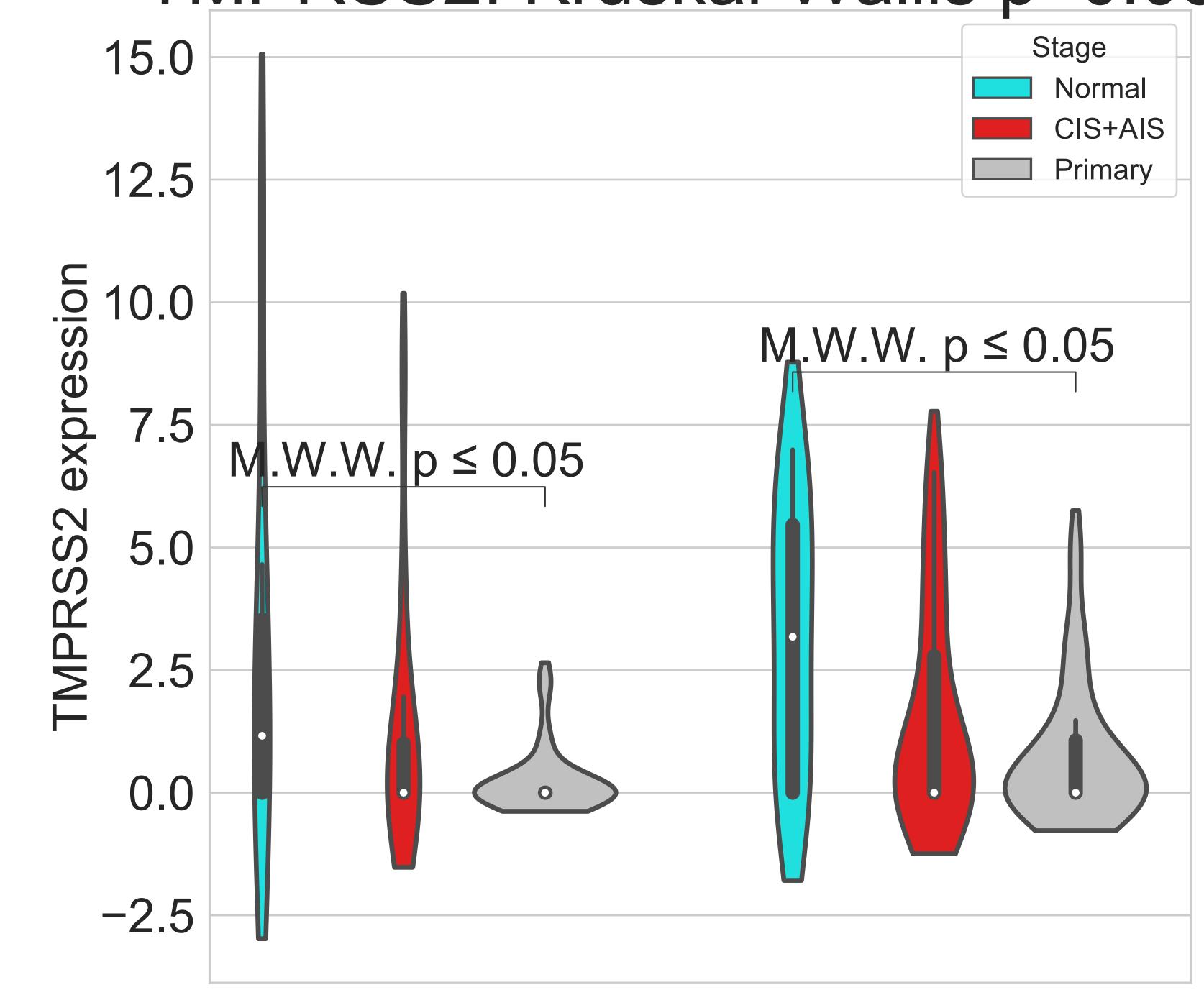
TMPRSS2: Kruskal-Wallis p=0.035



Lower
Mutation Shared Proportion (SYN)