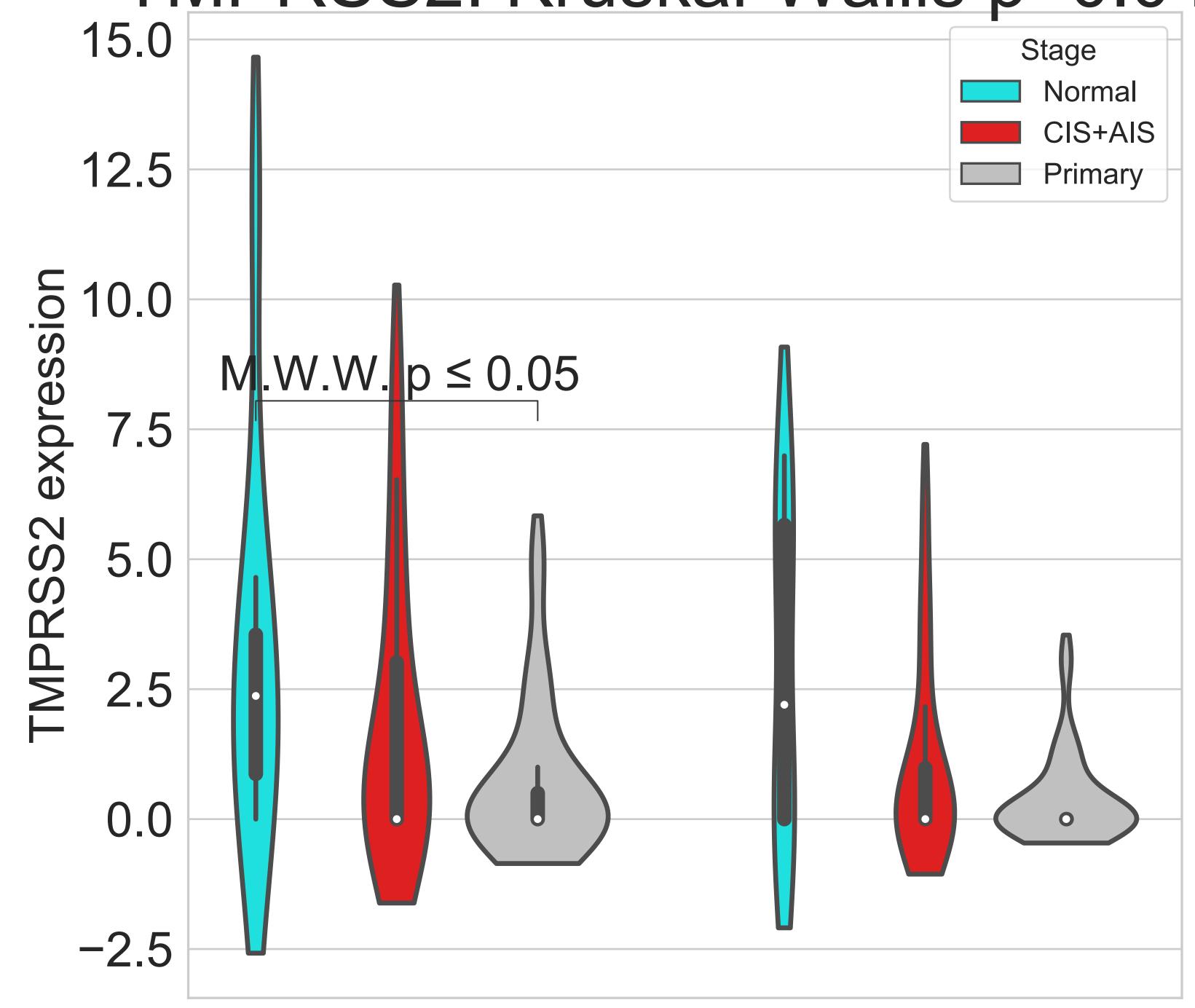
TMPRSS2: Kruskal-Wallis p=0.048



Lower
Mutation Shared Count (SYN)