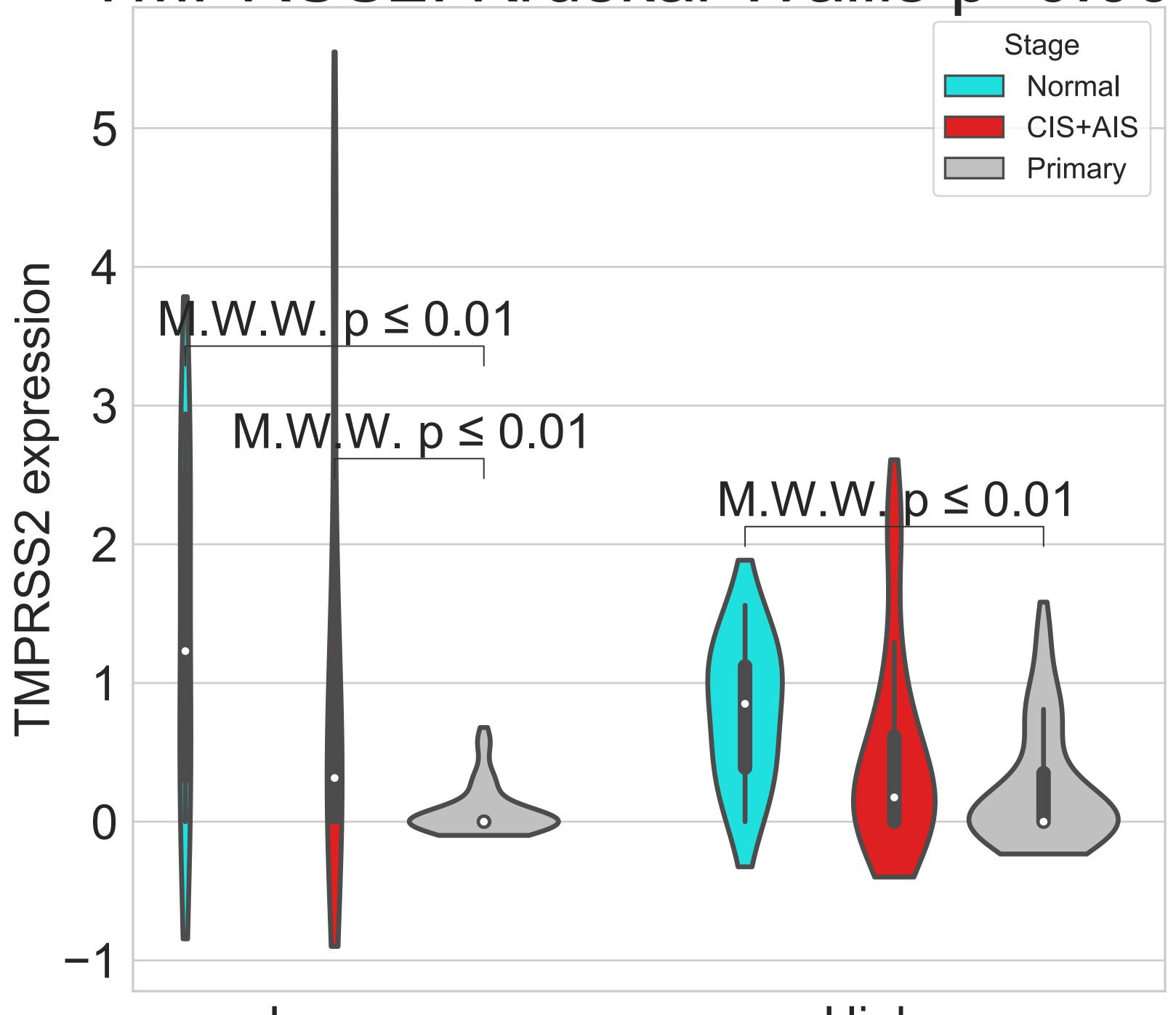
TMPRSS2: Kruskal-Wallis p=0.002



Lower
Higher
Mutation Shared Proportion