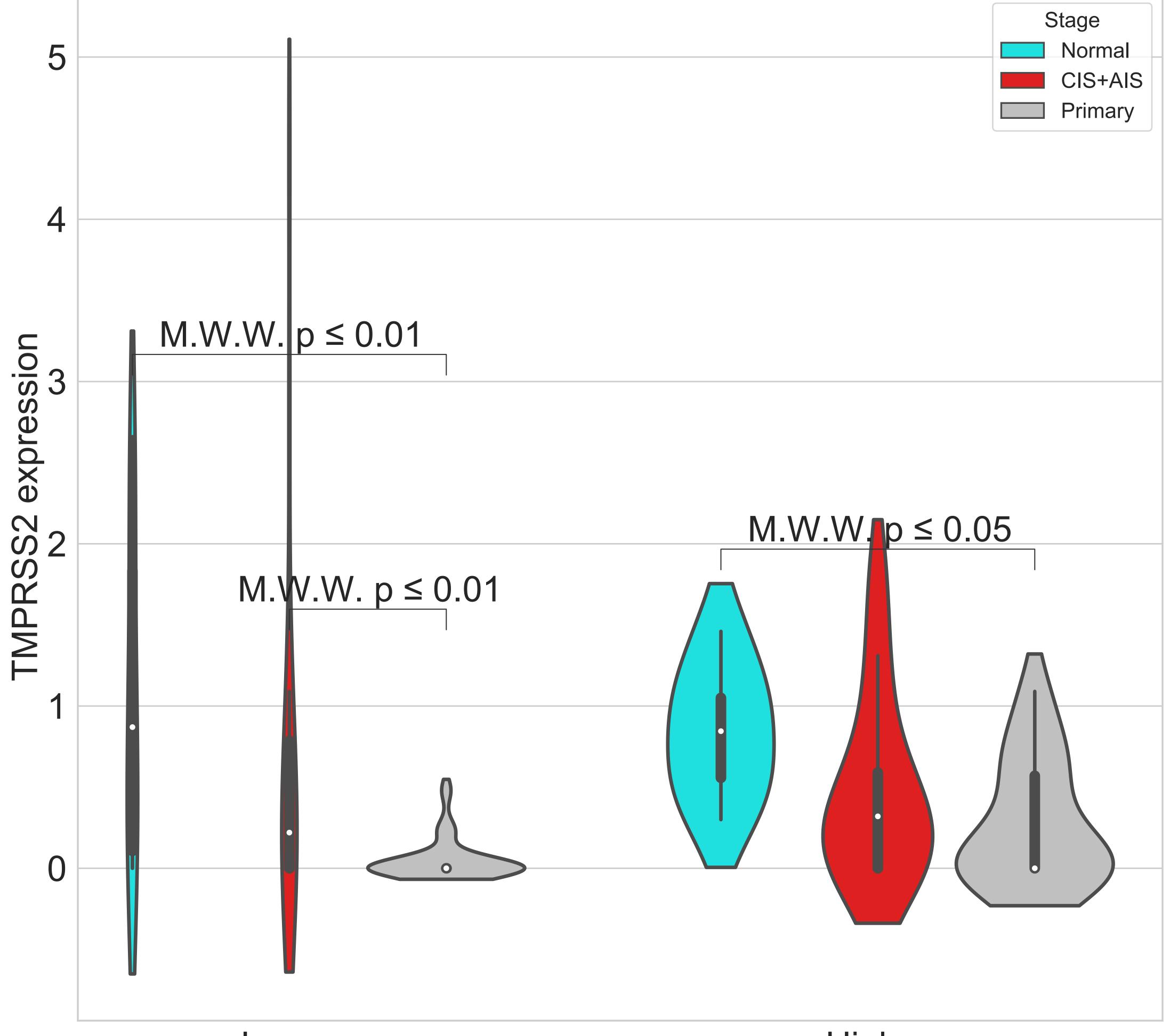
TMPRSS2: Kruskal-Wallis p=0.001



Higher Lower Mutation Shared Proportion (Union)