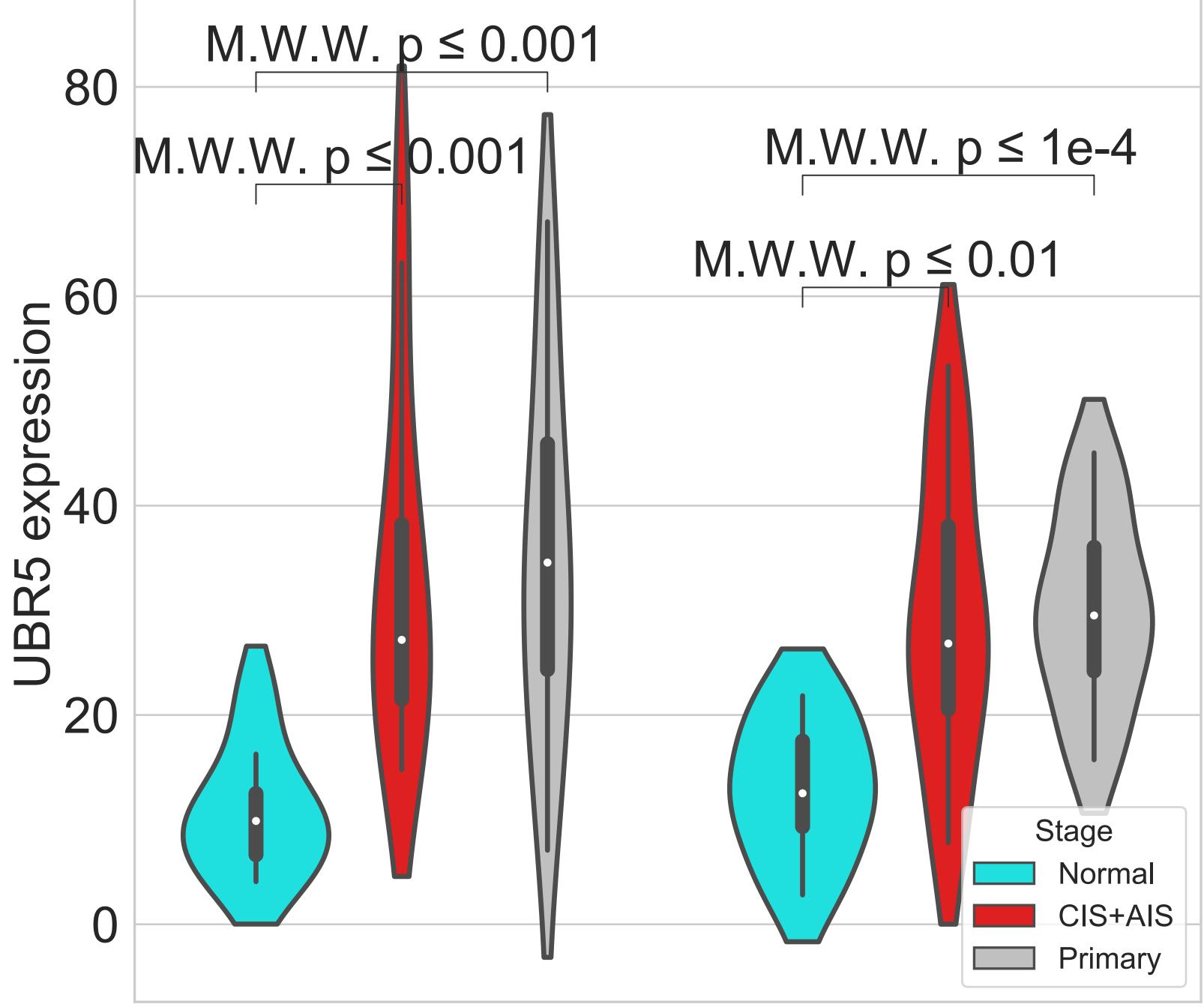
UBR5: Kruskal-Wallis p=0.000 MWW p≤0.001



Lower Higher Mutation Shared Proportion (Union & SYN)