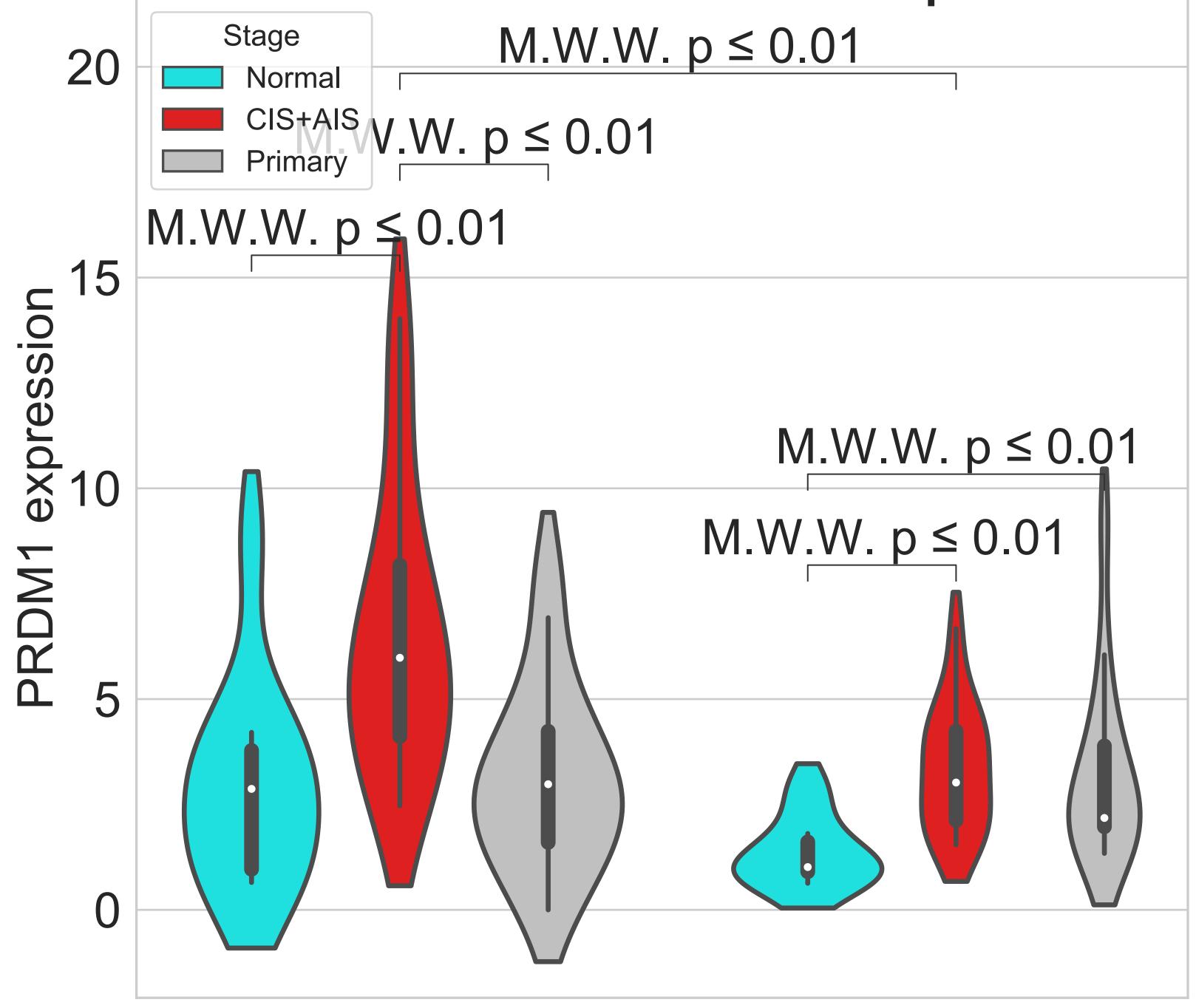
PRDM1: Kruskal-Wallis p=0.000



Lower Higher Mutation Shared Proportion (SYN)