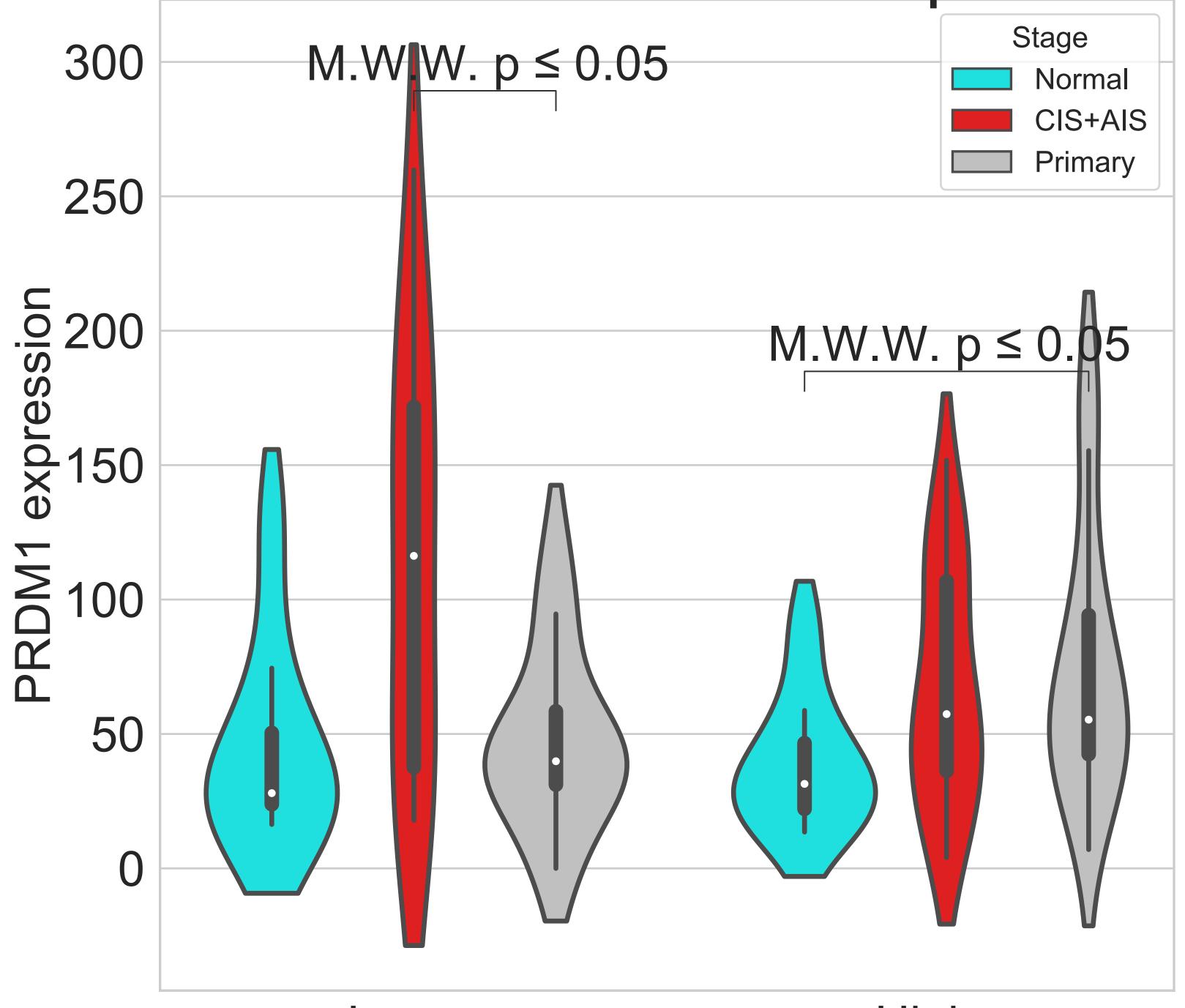
PRDM1: Kruskal-Wallis p=0.038



Lower Higher Mutation Shared Proportion (Union & SYN)