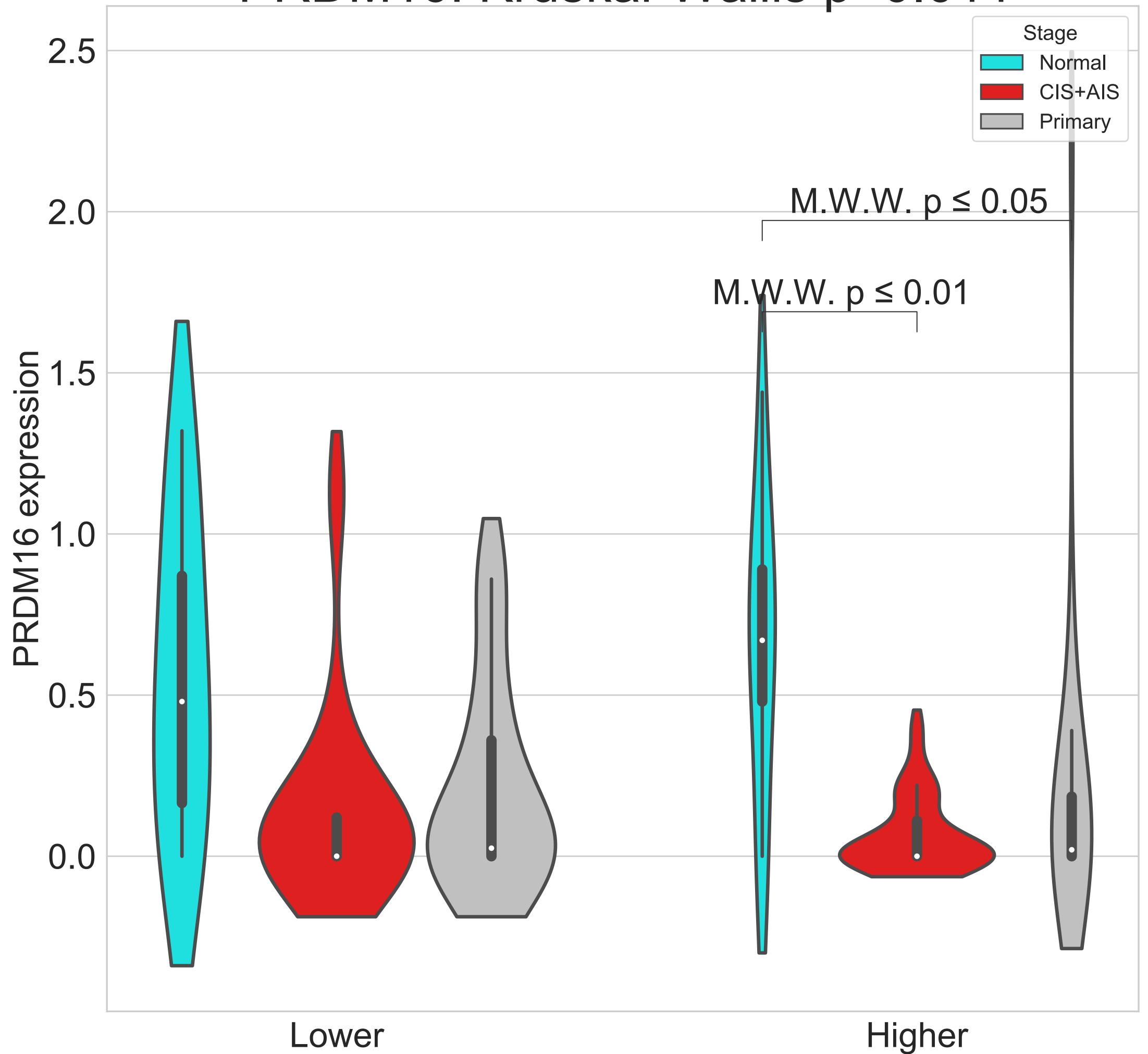
PRDM16: Kruskal-Wallis p=0.011



Mutation Shared Proportion (SYN)