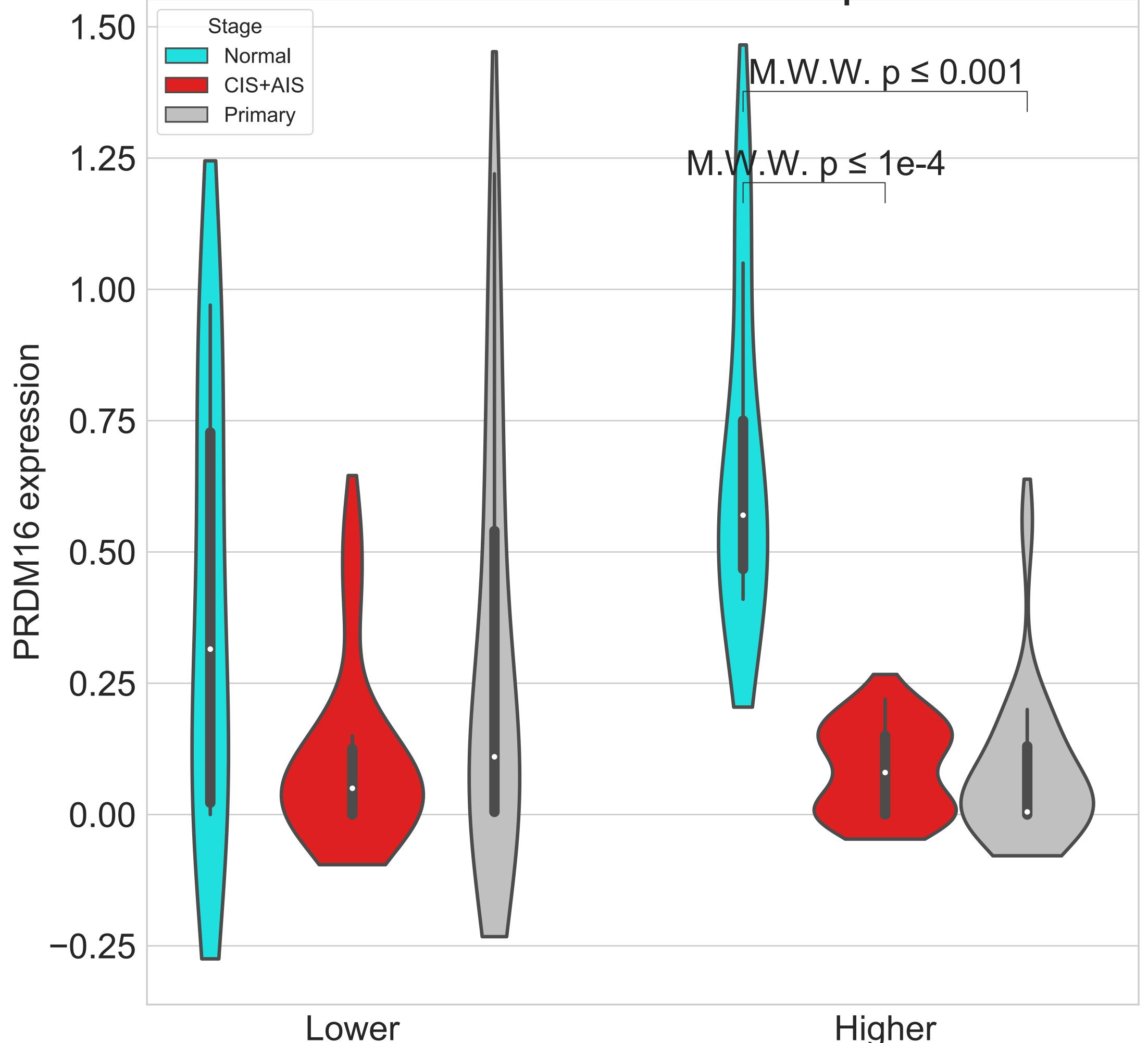
PRDM16: Kruskal-Wallis p=0.001



Higher Mutation Shared Count (SYN)