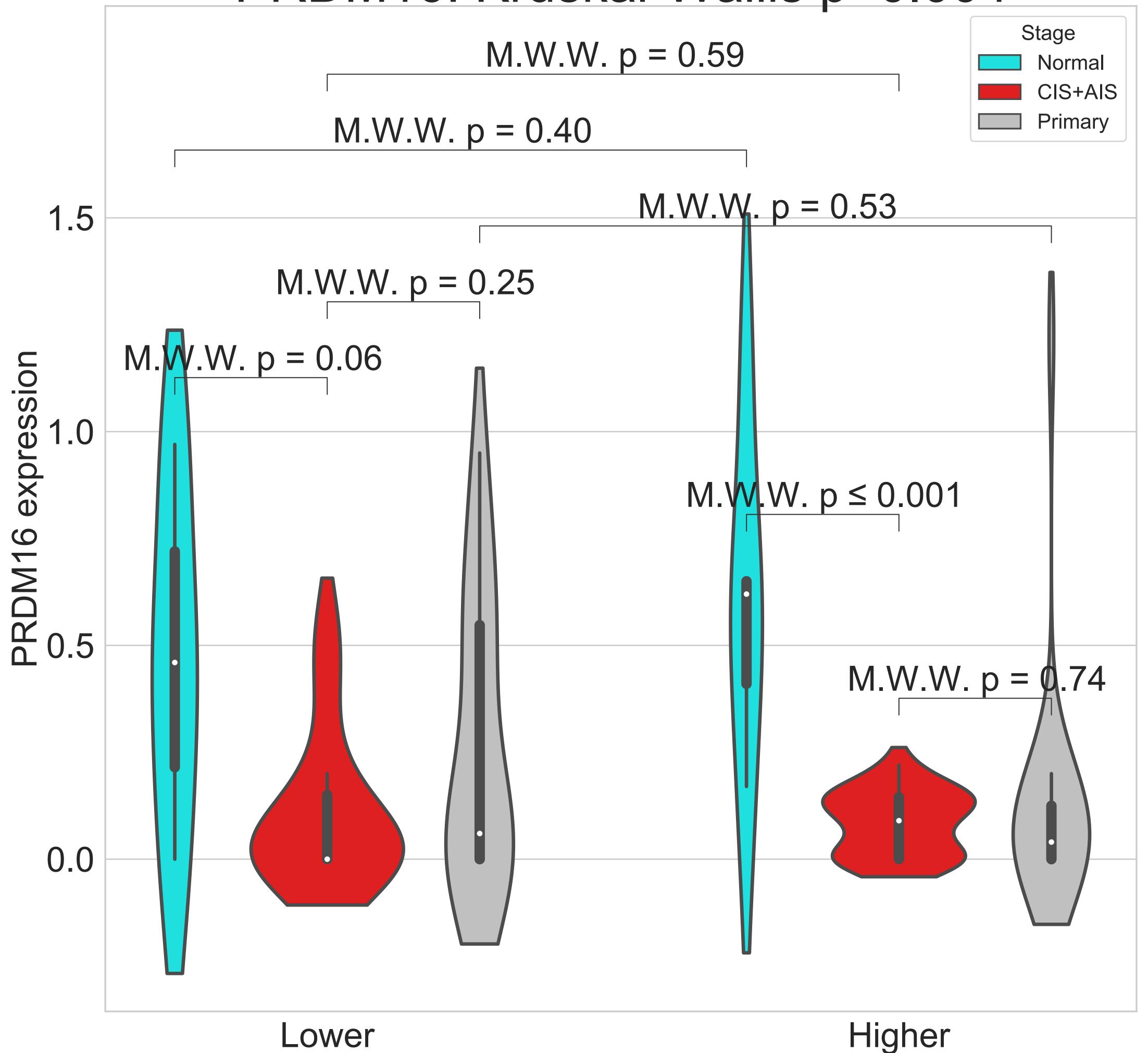
PRDM16: Kruskal-Wallis p=0.004



Mutation Shared Count (SYN) per TMB