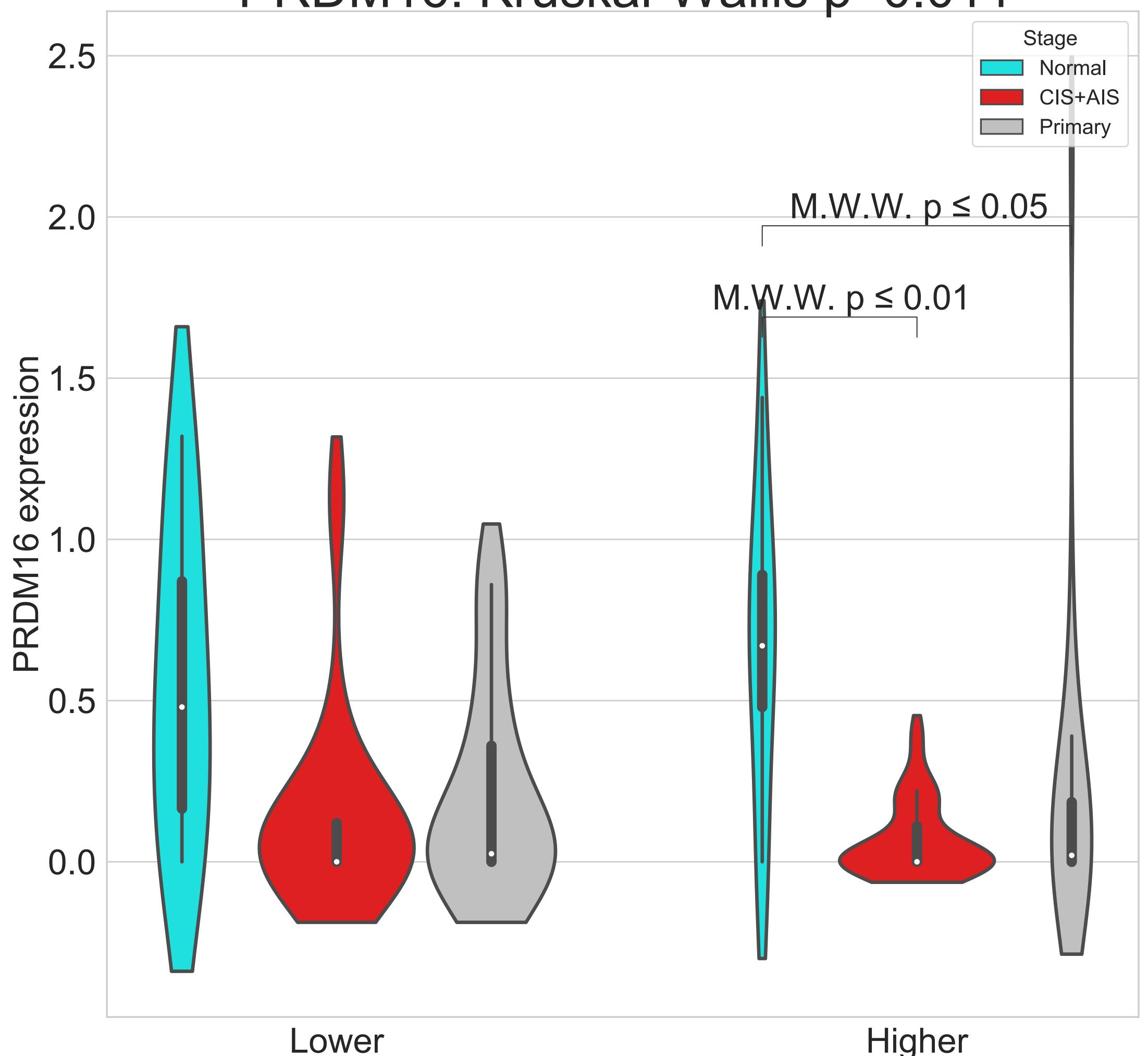
PRDM16: Kruskal-Wallis p=0.011



Lower
Mutation Shared Count (SYN) per TMB