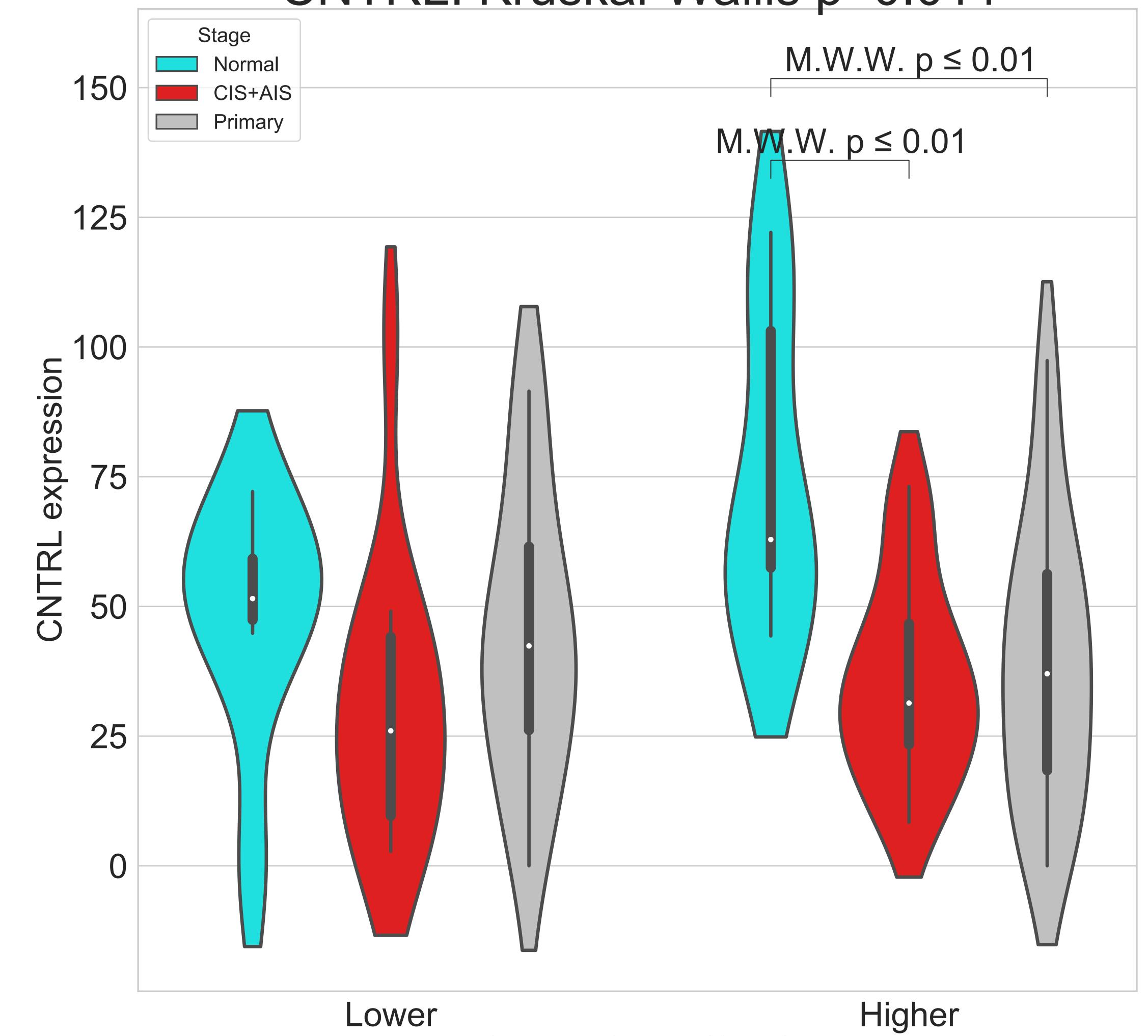
CNTRL: Kruskal-Wallis p=0.011



Mutation Shared Count (SYN) per TMB