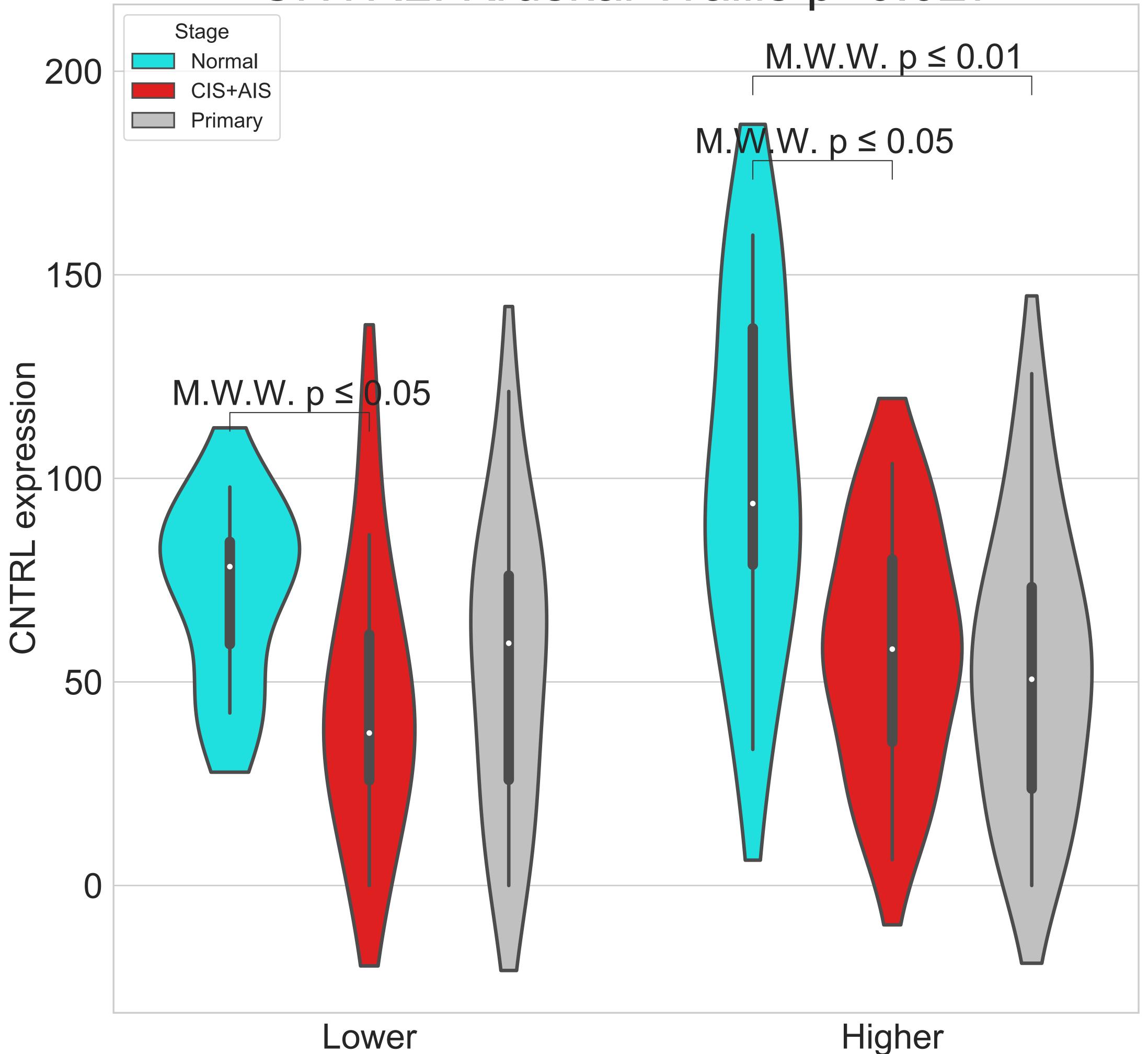
CNTRL: Kruskal-Wallis p=0.027



Lower
Mutation Shared Count (SYN) per TMB