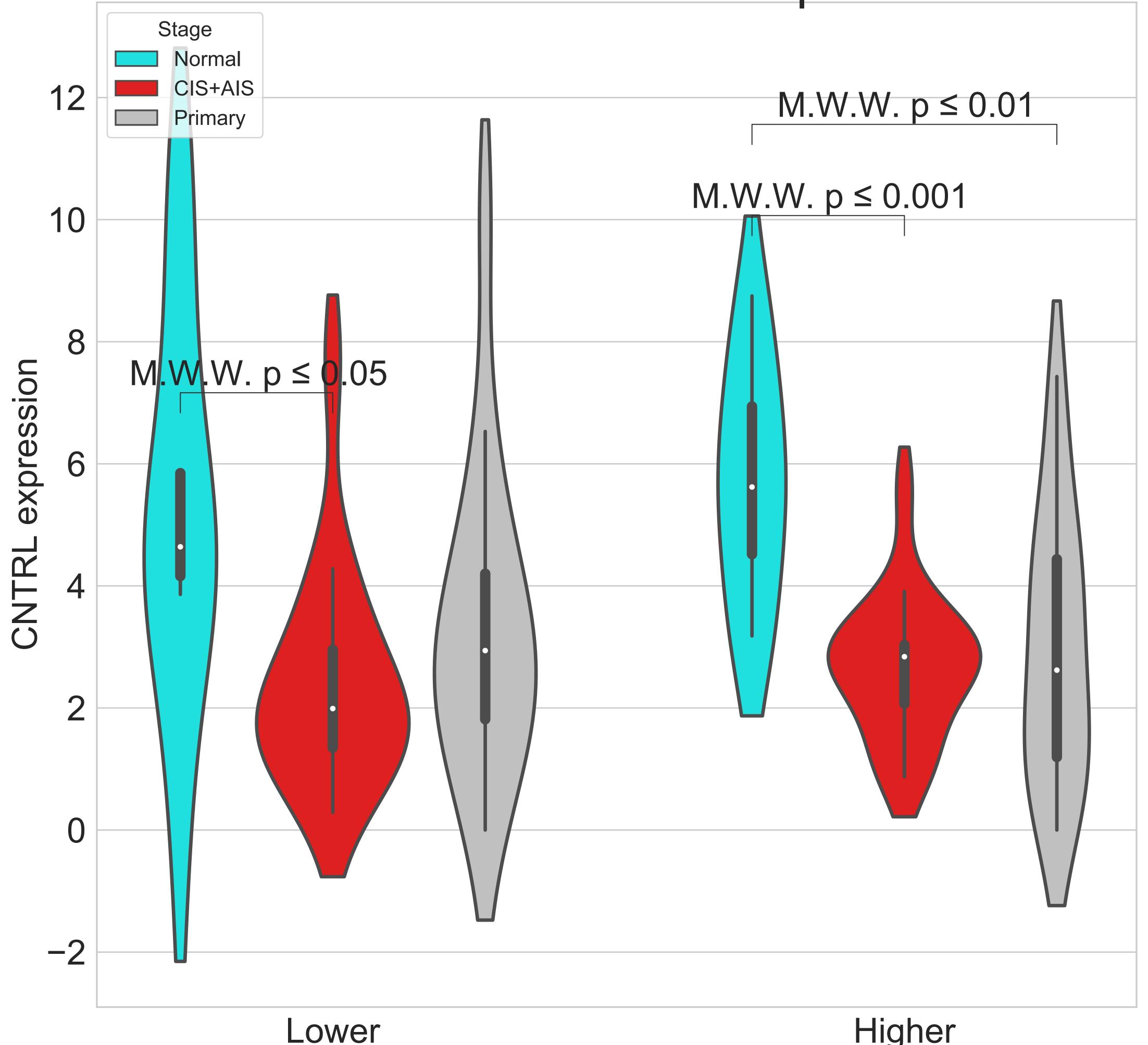
CNTRL: Kruskal-Wallis p=0.004



Higher Mutation Shared Count (SYN)