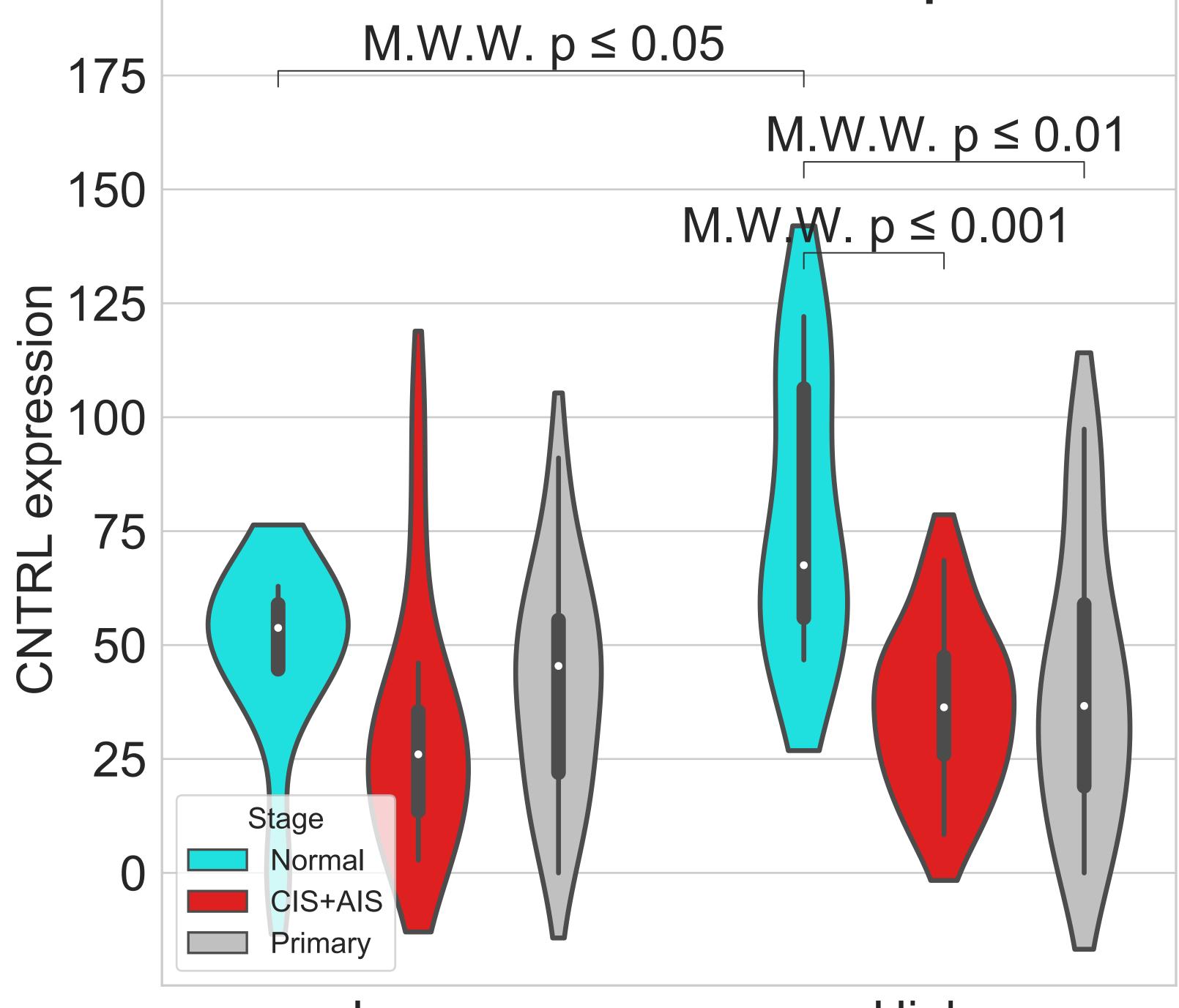
CNTRL: Kruskal-Wallis p=0.007



Lower
Mutation Shared Count (SYN)