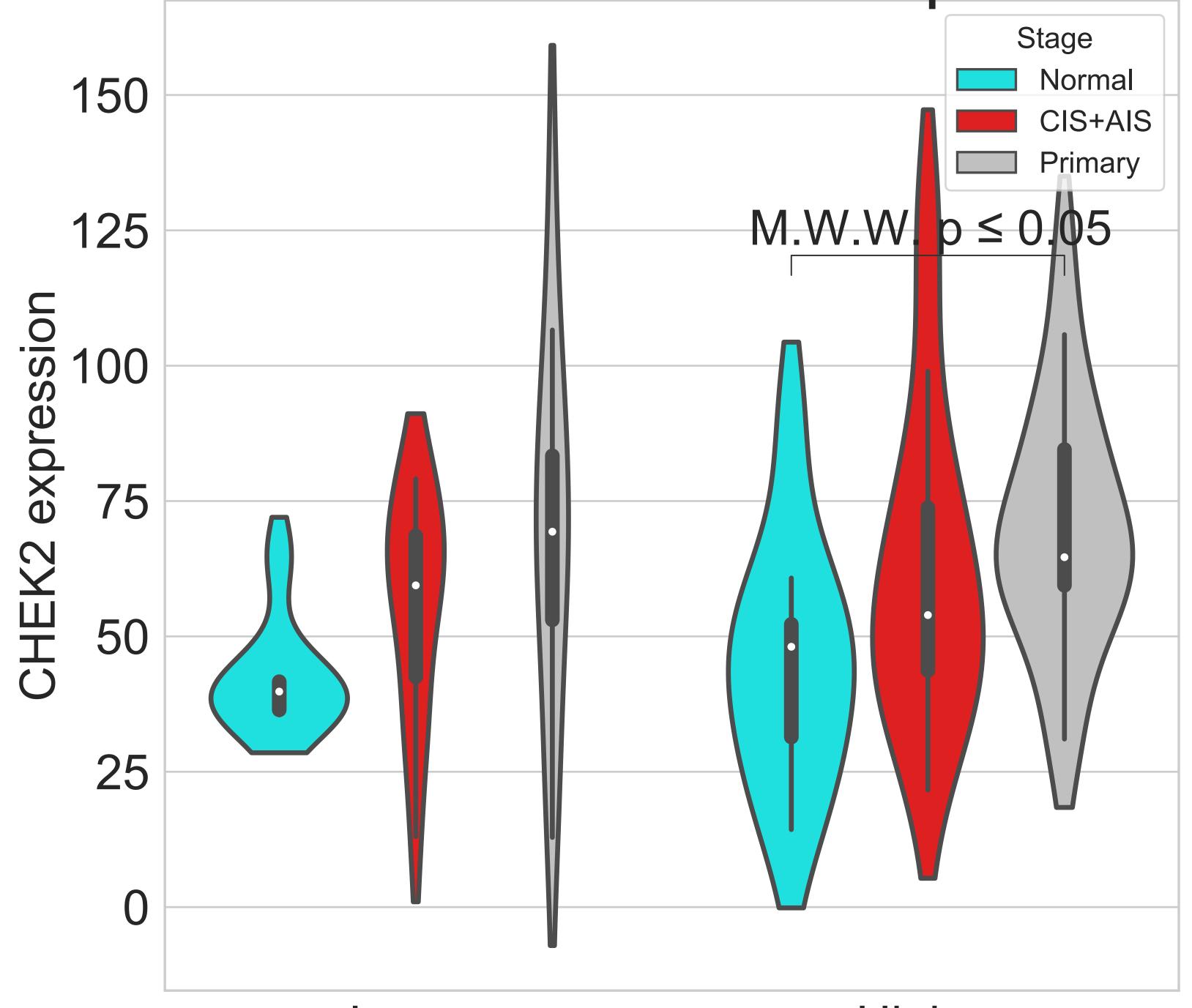
CHEK2: Kruskal-Wallis p=0.033



Lower Higher Mutation Shared Count (SYN) per TMB