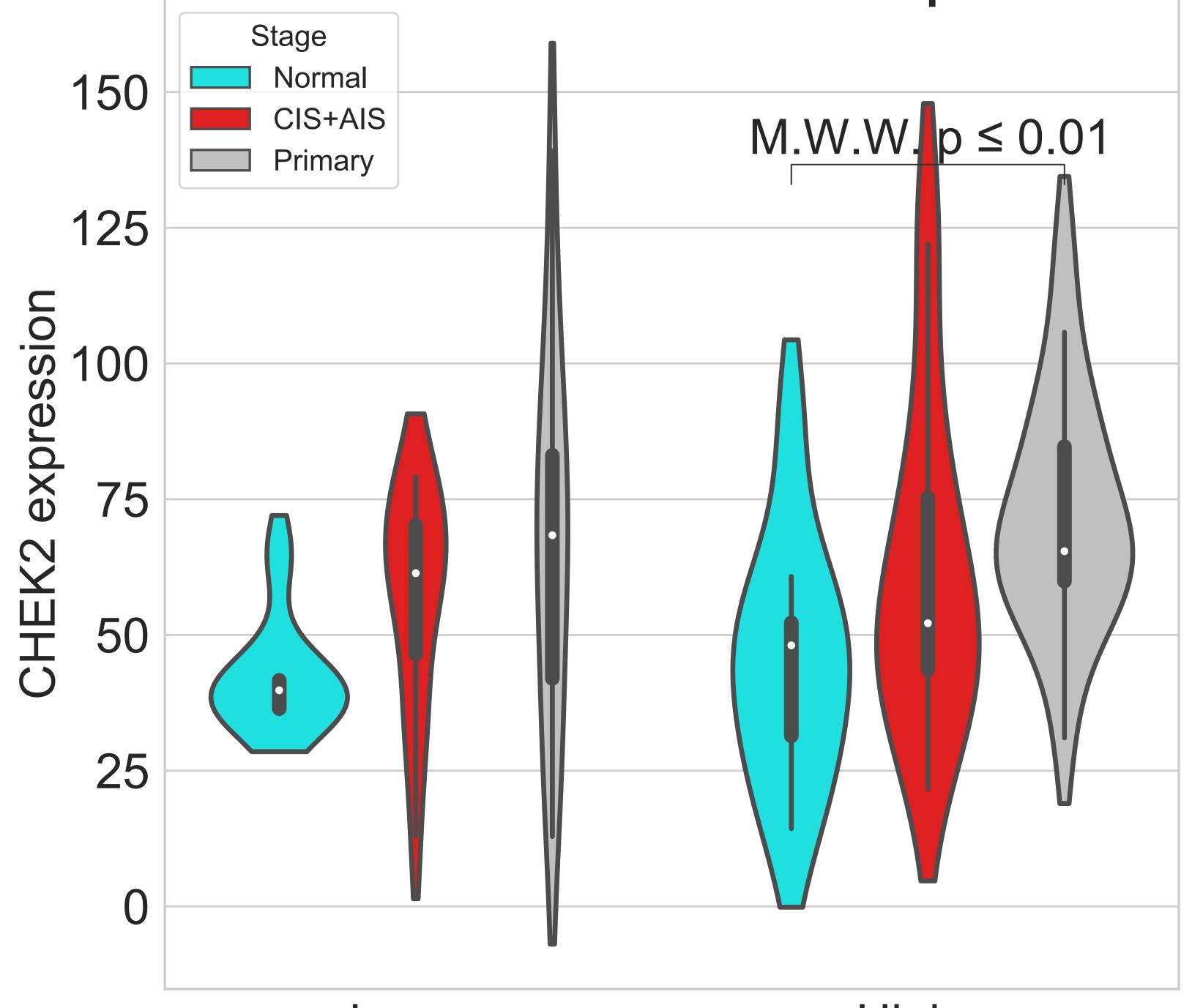
CHEK2: Kruskal-Wallis p=0.027



Lower Higher Mutation Shared Count per TMB