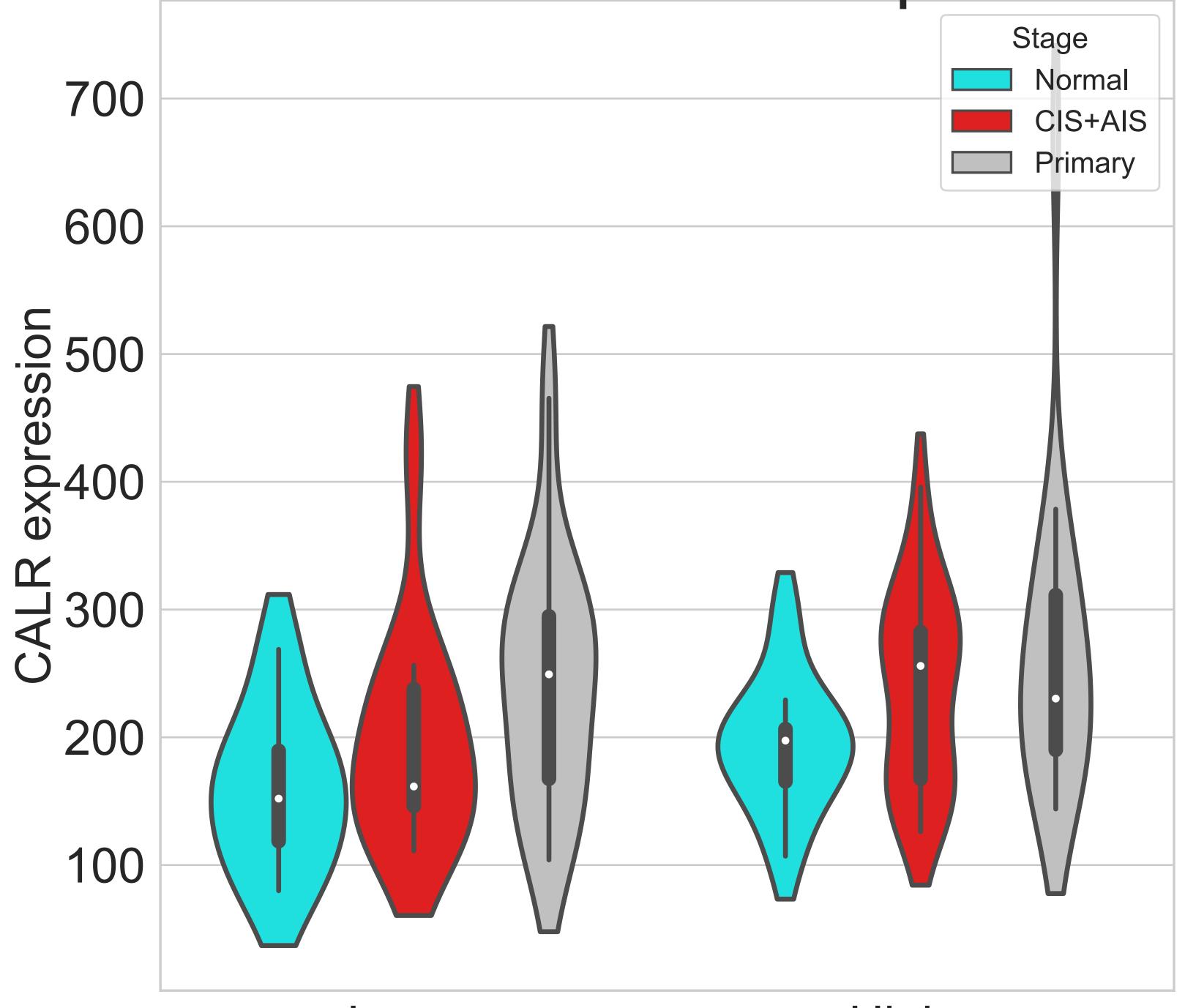
CALR: Kruskal-Wallis p=0.035



Lower Higher Mutation Shared Count (SYN) per TMB