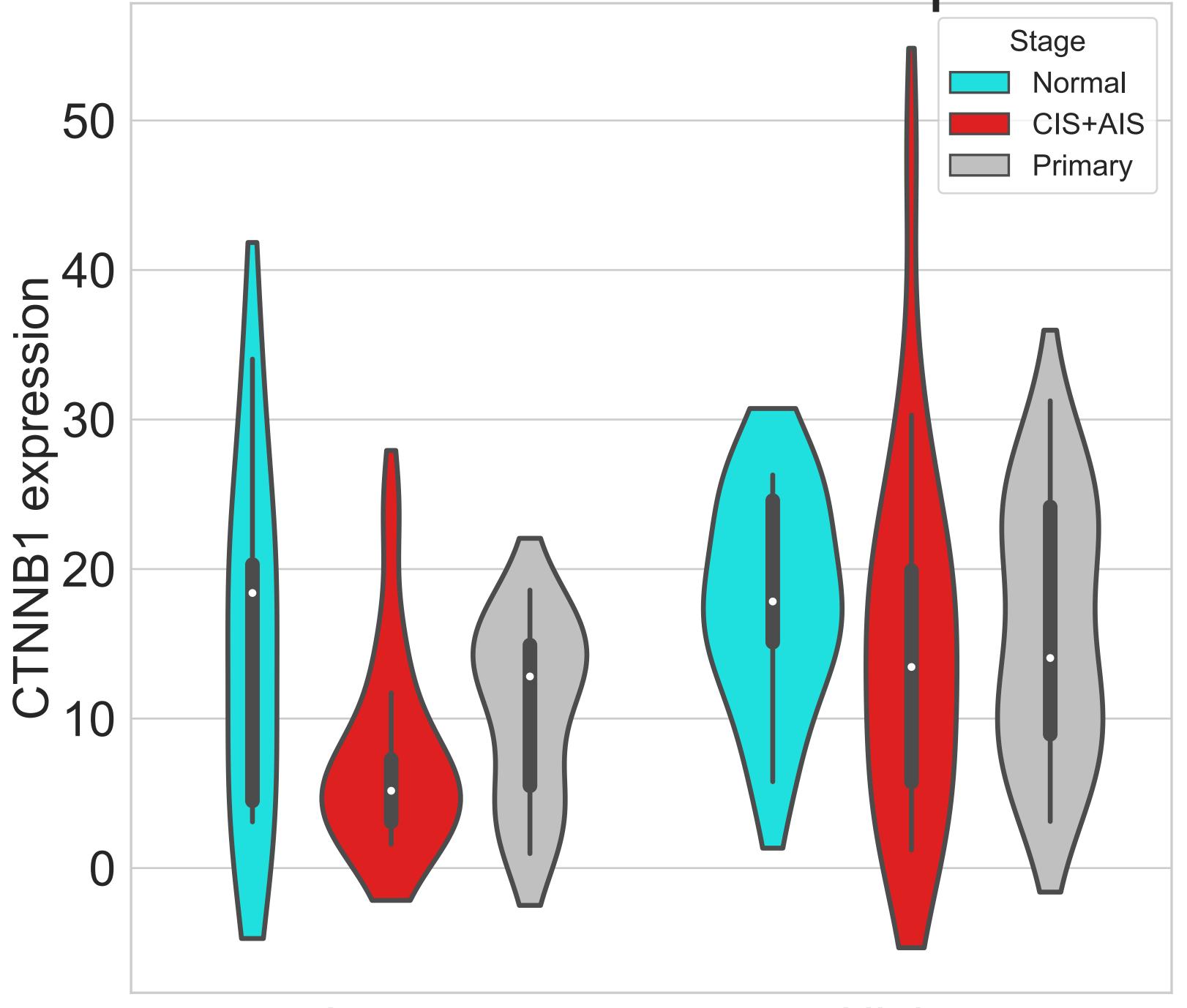
CTNNB1: Kruskal-Wallis p=0.019



Lower Higher Mutation Shared Count (SYN) per TMB