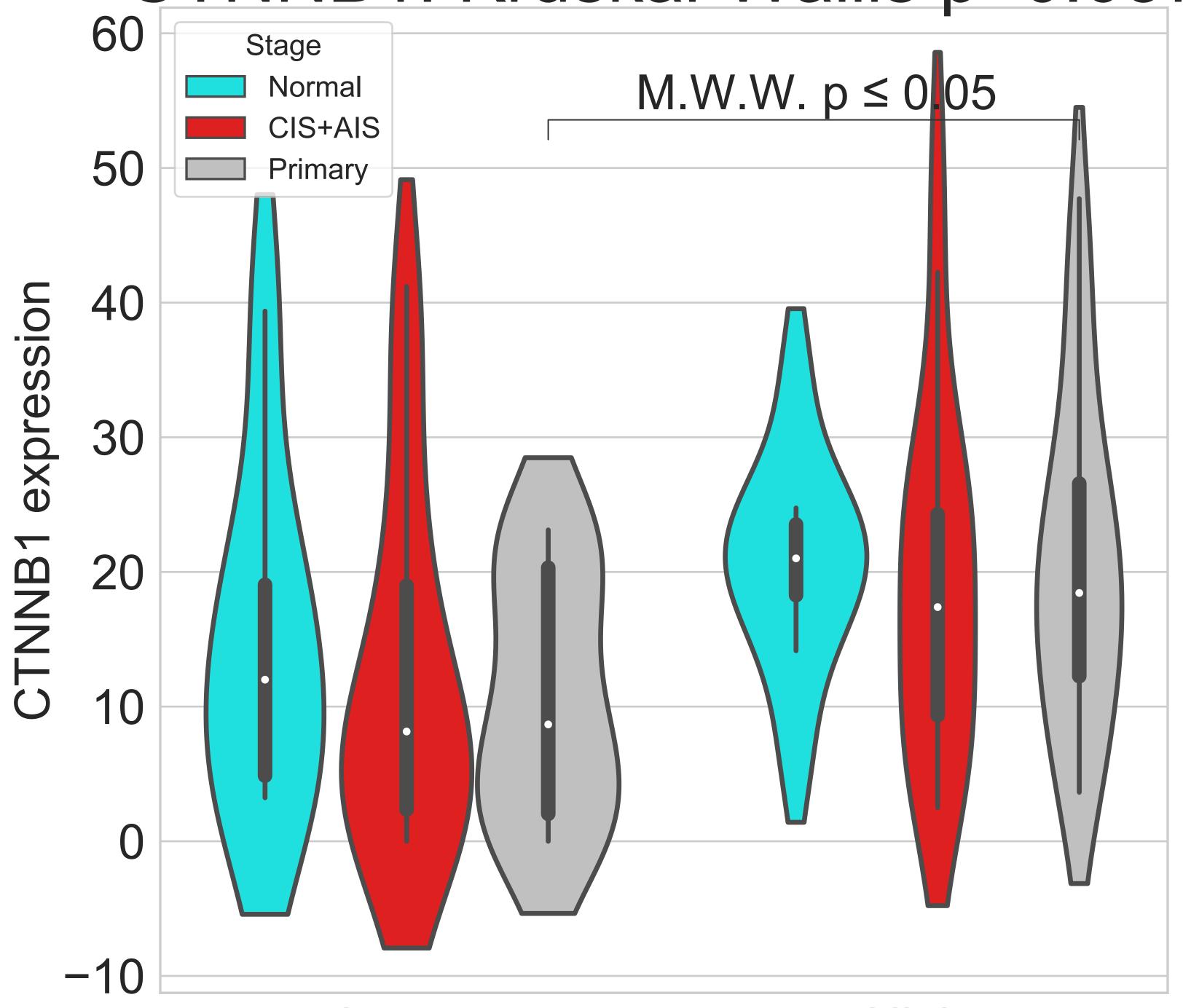
CTNNB1: Kruskal-Wallis p=0.037



Lower Higher Mutation Shared Count (SYN) per TMB