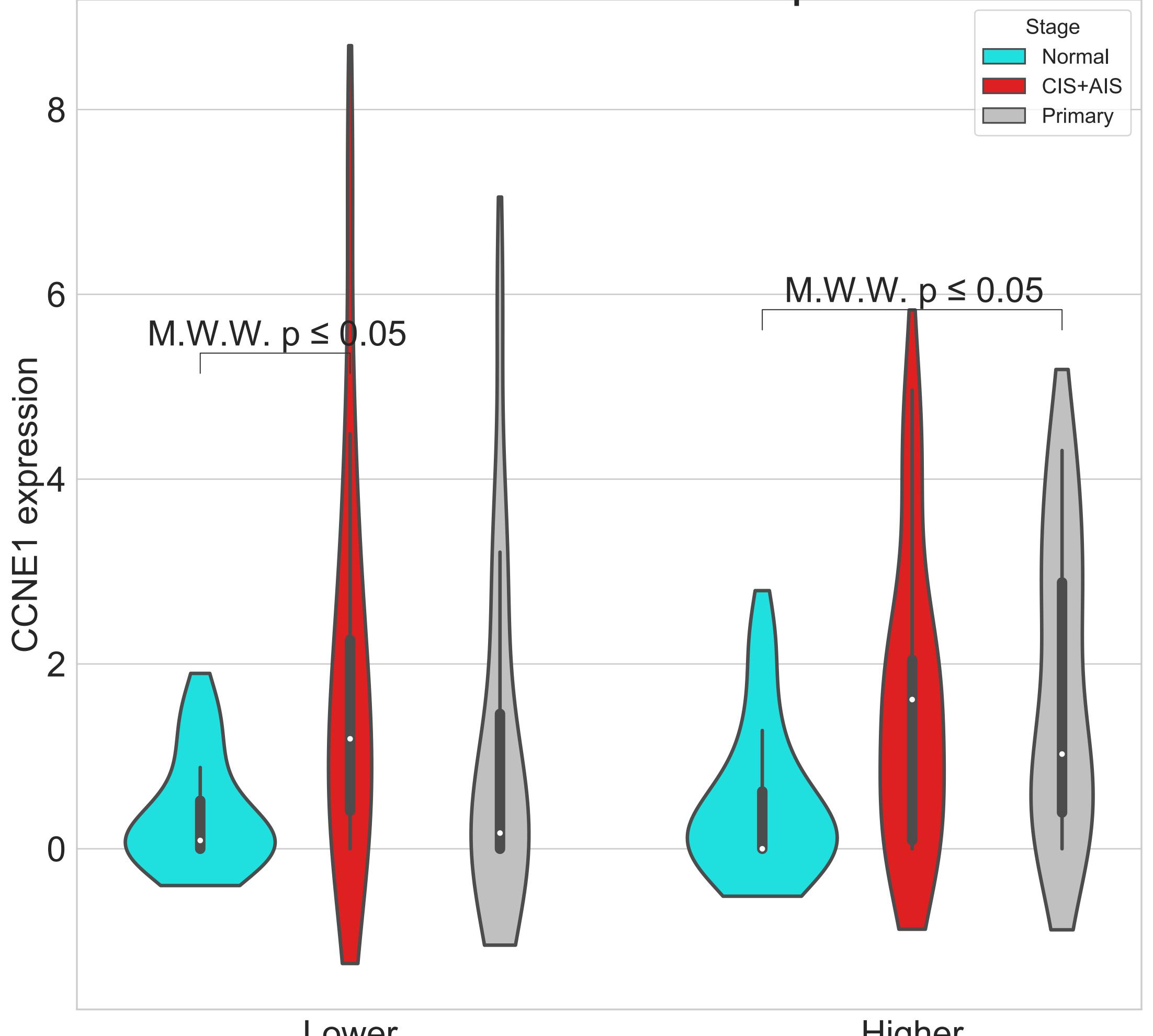
CCNE1: Kruskal-Wallis p=0.044



Higher Lower Mutation Shared Count per TMB