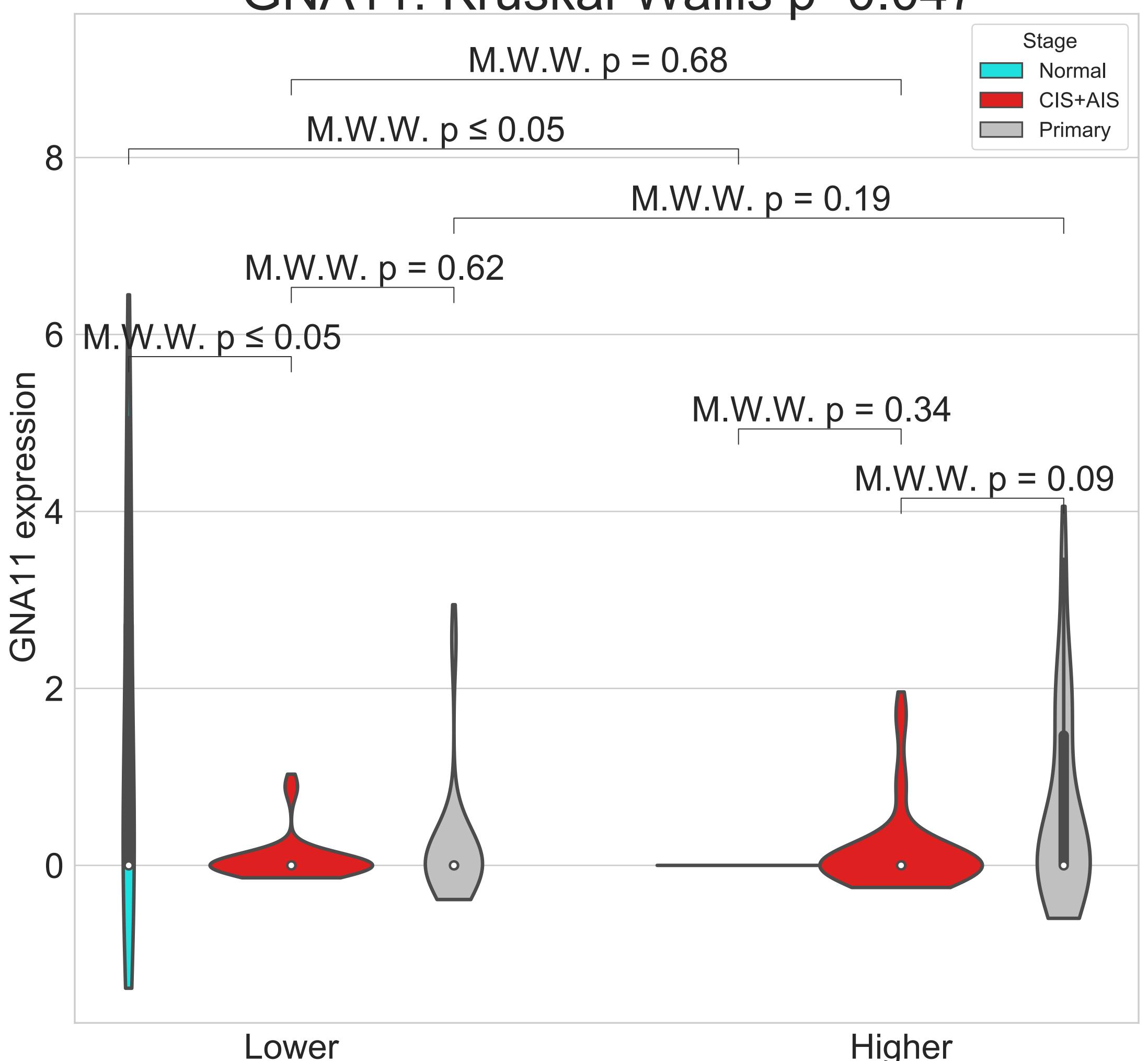
GNA11: Kruskal-Wallis p=0.047



Higher

Mutation Shared Count per TMB