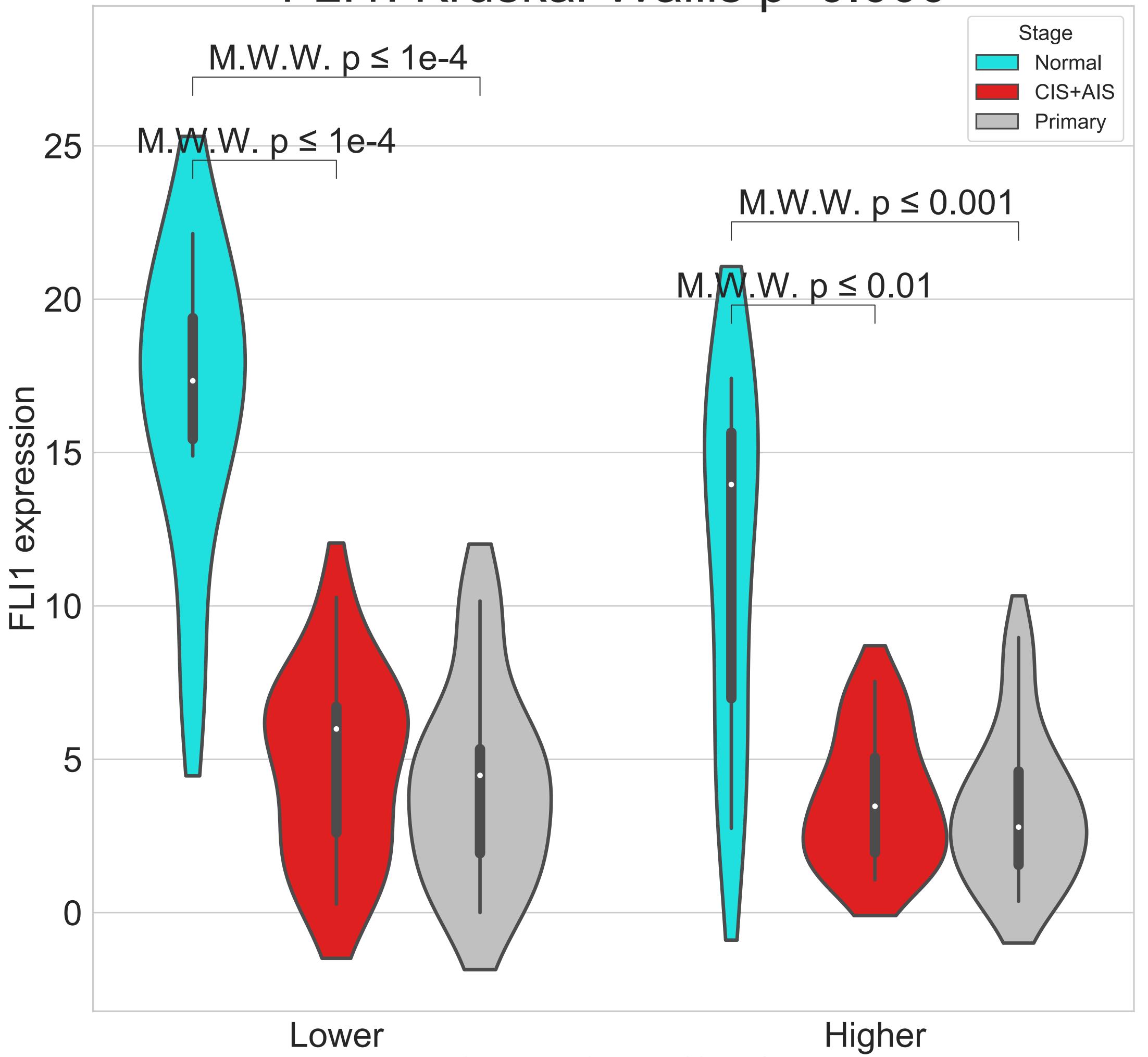
FLI1: Kruskal-Wallis p=0.000



Mutation Shared Count (SYN) per TMB