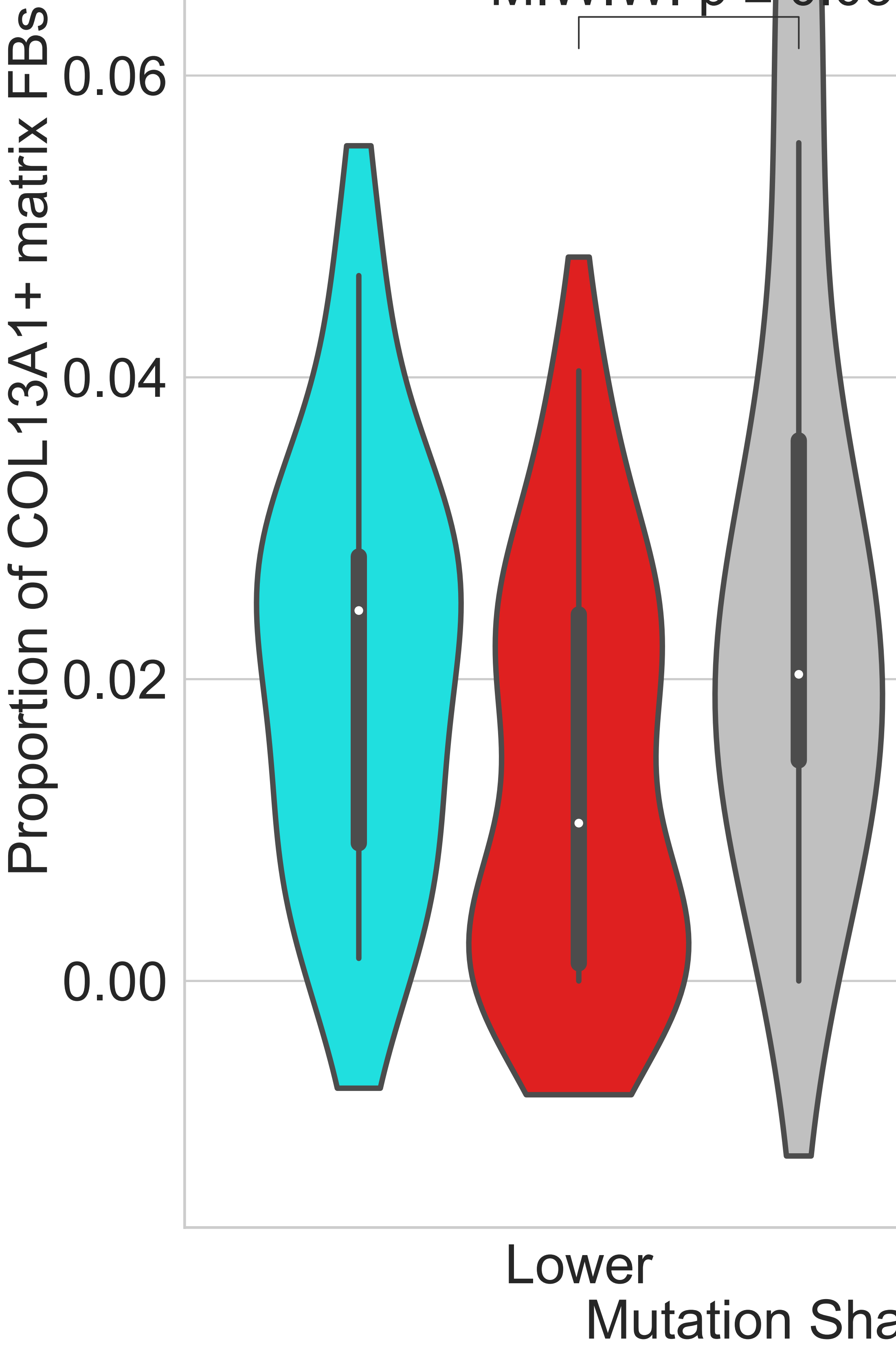


Kruskal-Wallis p=0.046



Higher

Mutation Shared Count per TMB