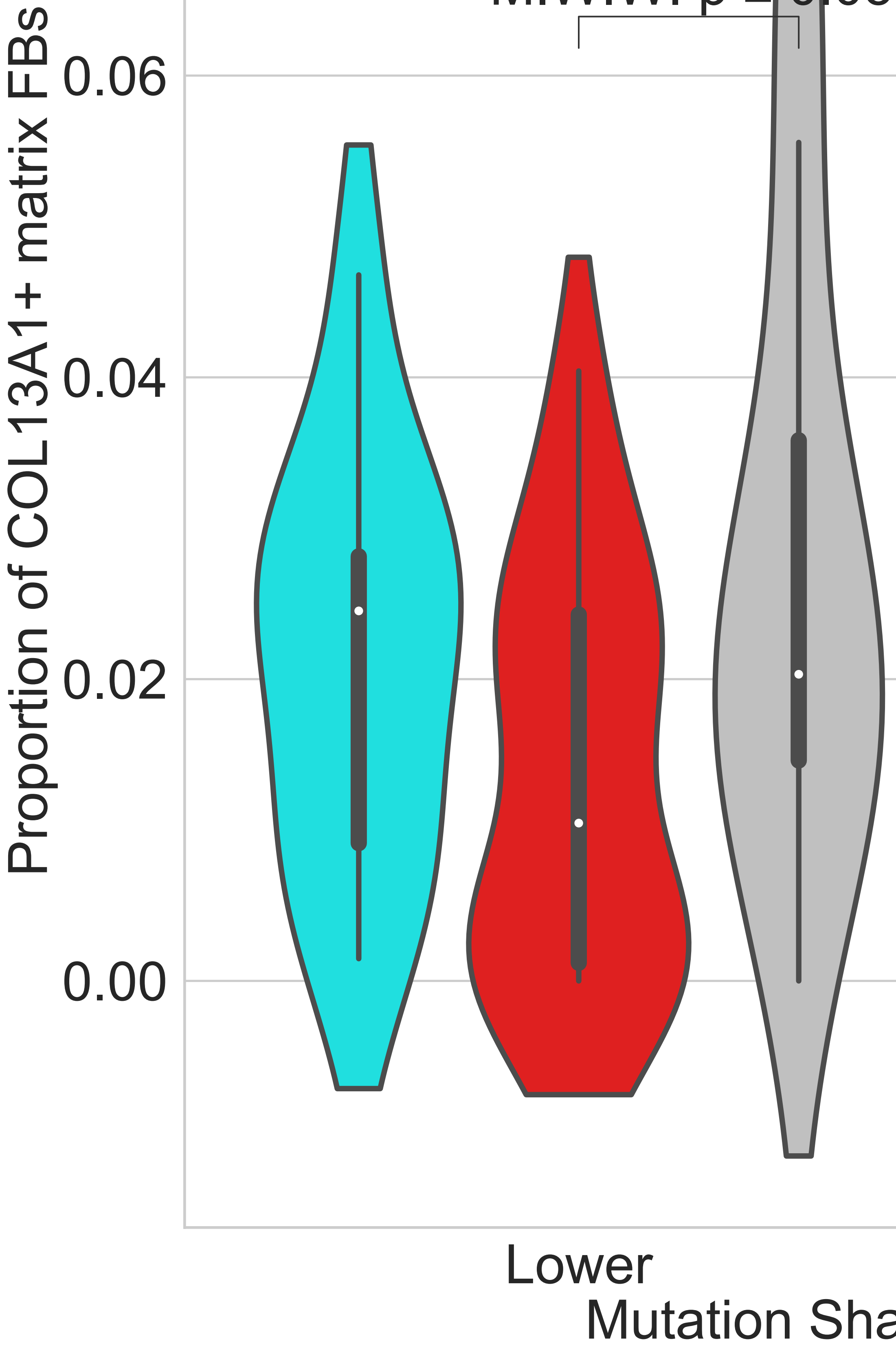


Kruskal-Wallis p=0.047



Higher Mutation Shared Count per TMB