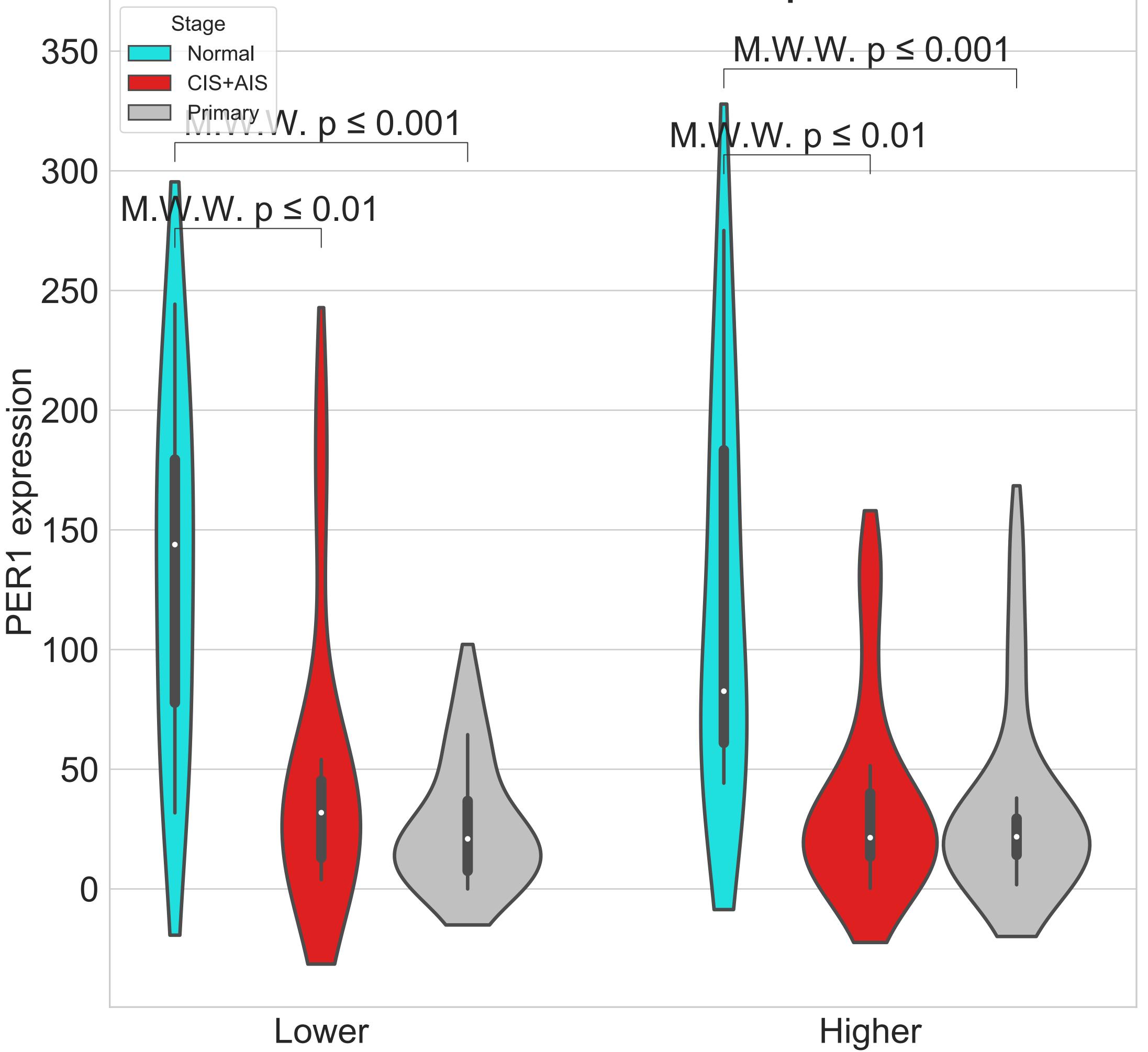
PER1: Kruskal-Wallis p=0.000



Mutation Shared Count per TMB