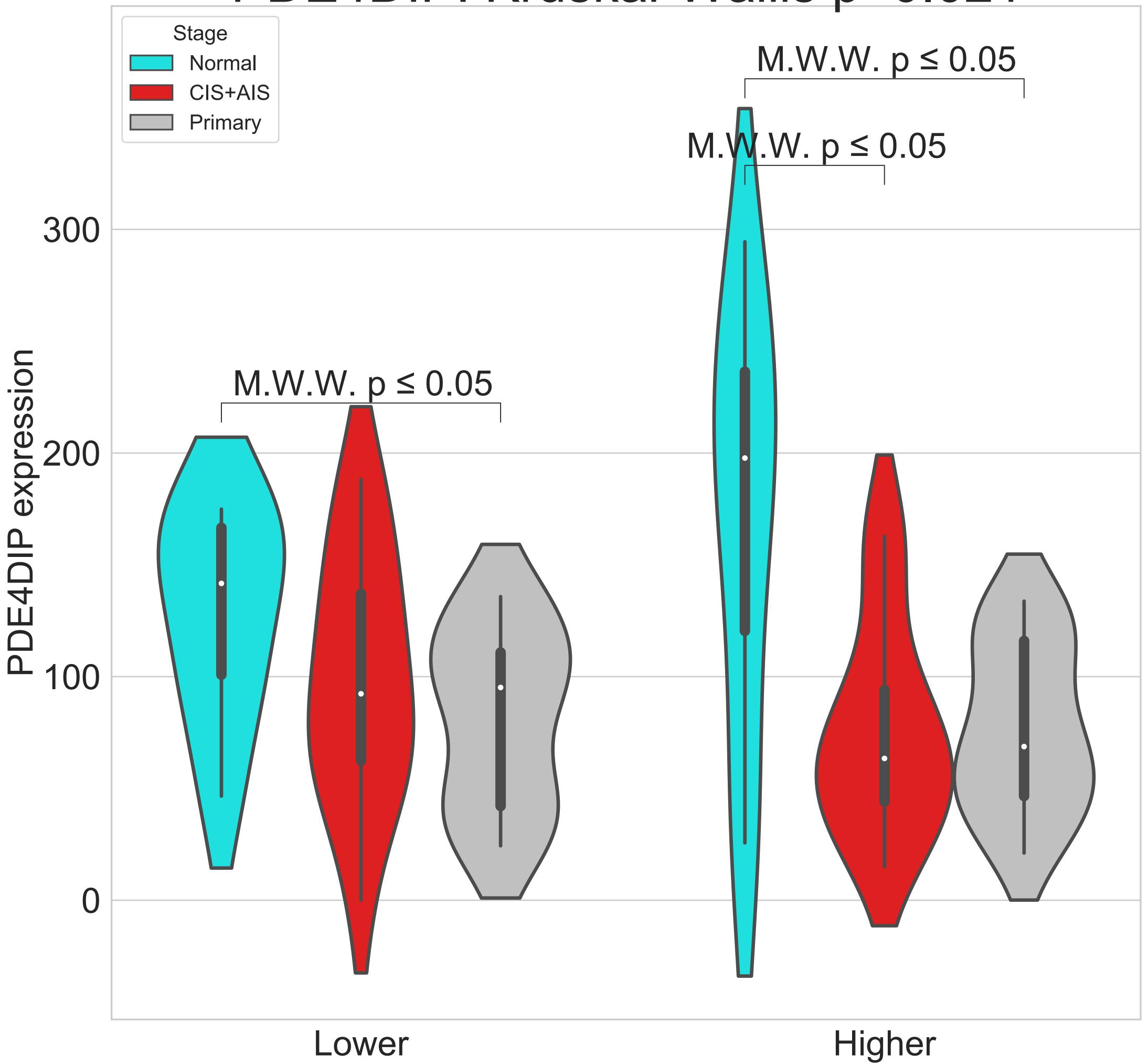
PDE4DIP: Kruskal-Wallis p=0.021



Mutation Shared Count per TMB