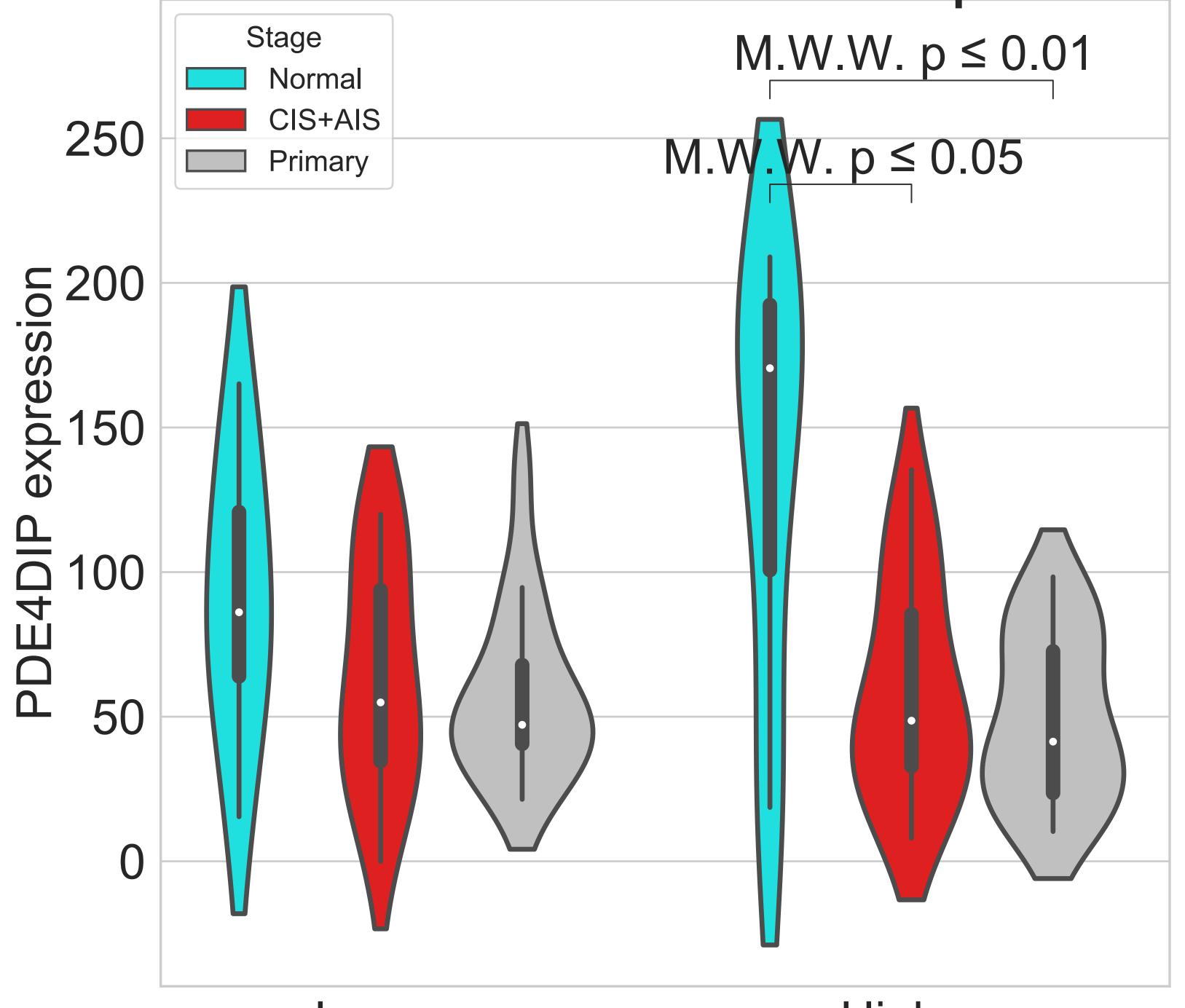
PDE4DIP: Kruskal-Wallis p=0.037



Lower Higher Mutation Shared Count per TMB