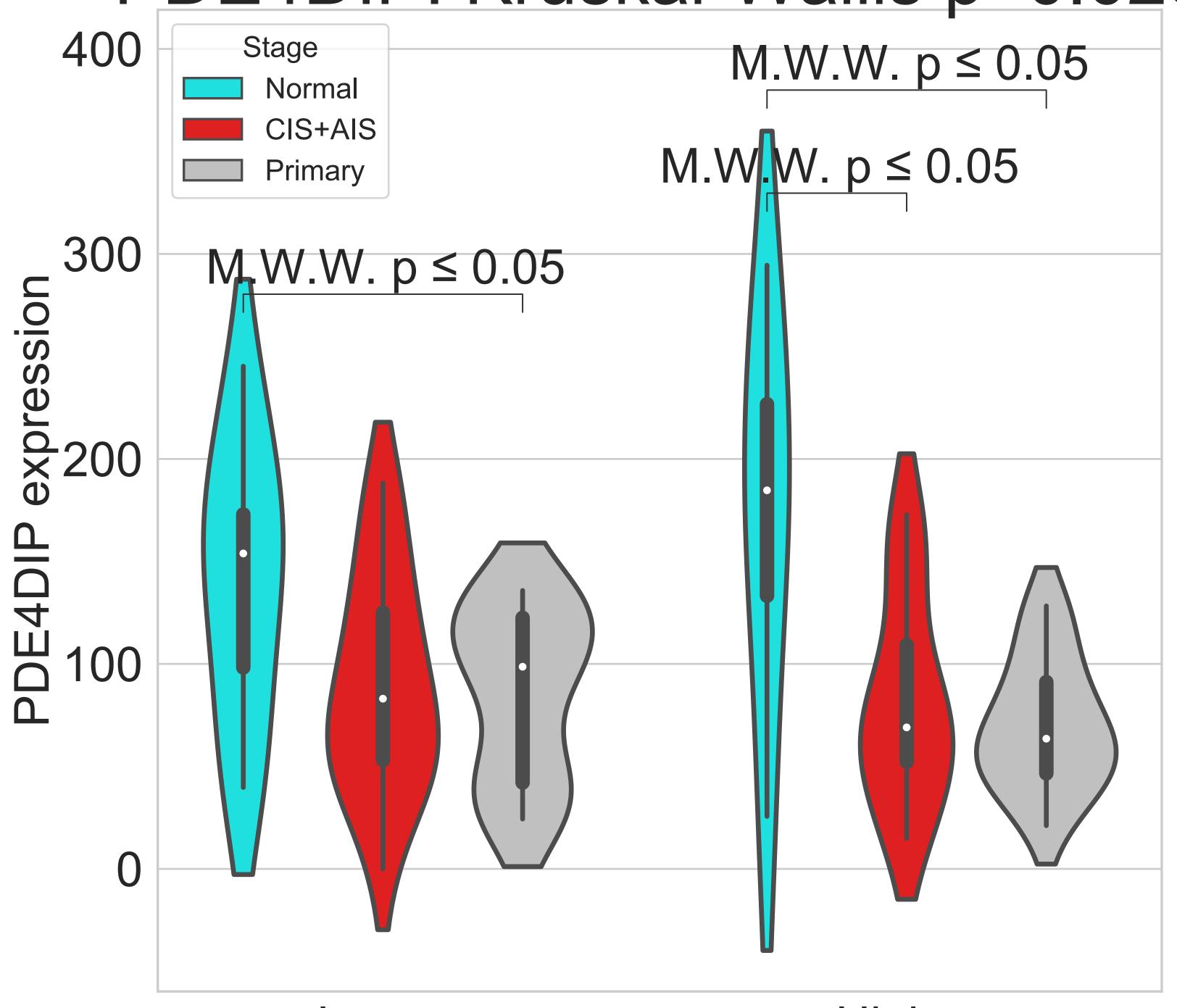
PDE4DIP: Kruskal-Wallis p=0.028



Lower
Mutation Shared Count per TMB