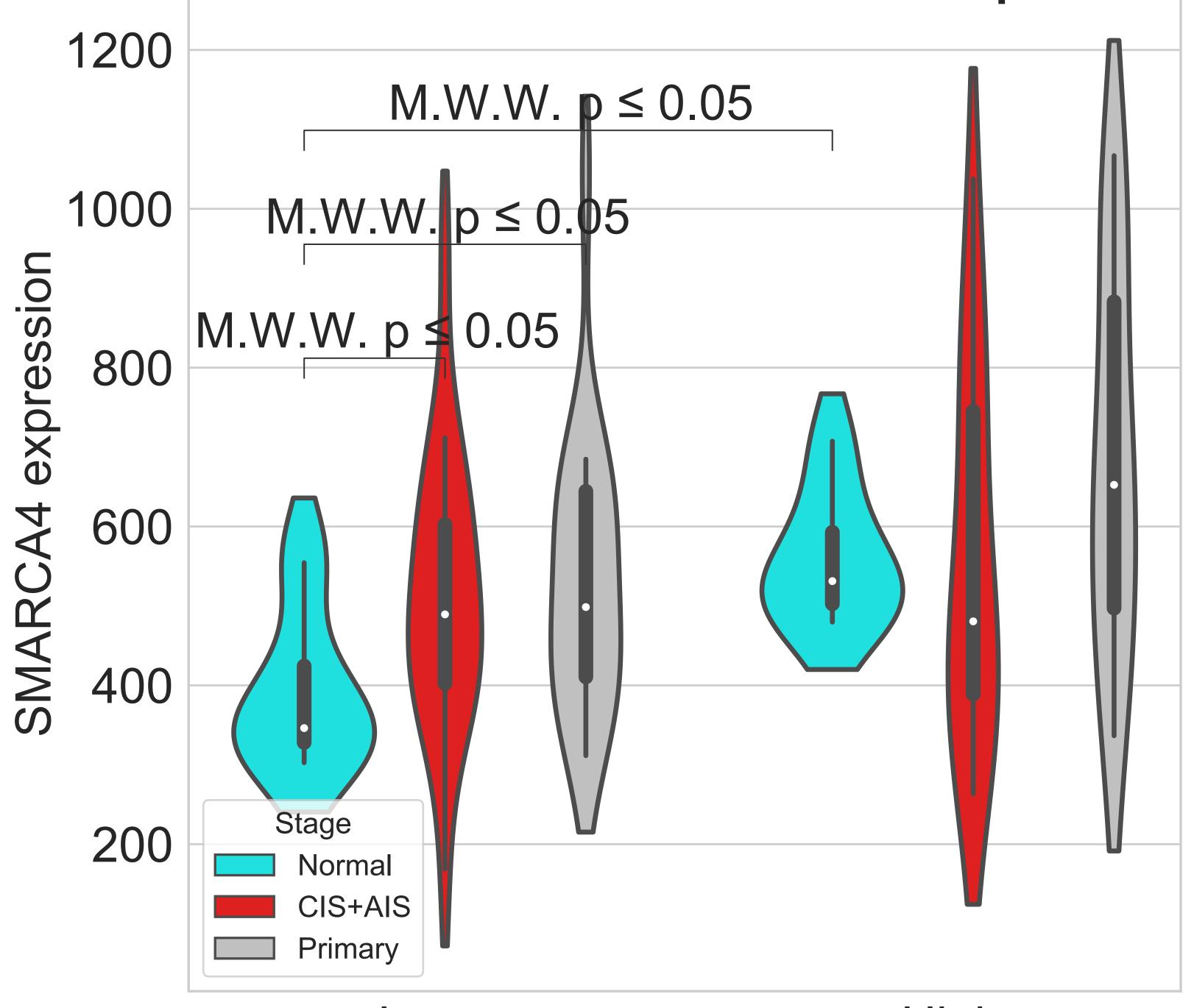
SMARCA4: Kruskal-Wallis p=0.022



Lower Higher Mutation Shared Count per TMB