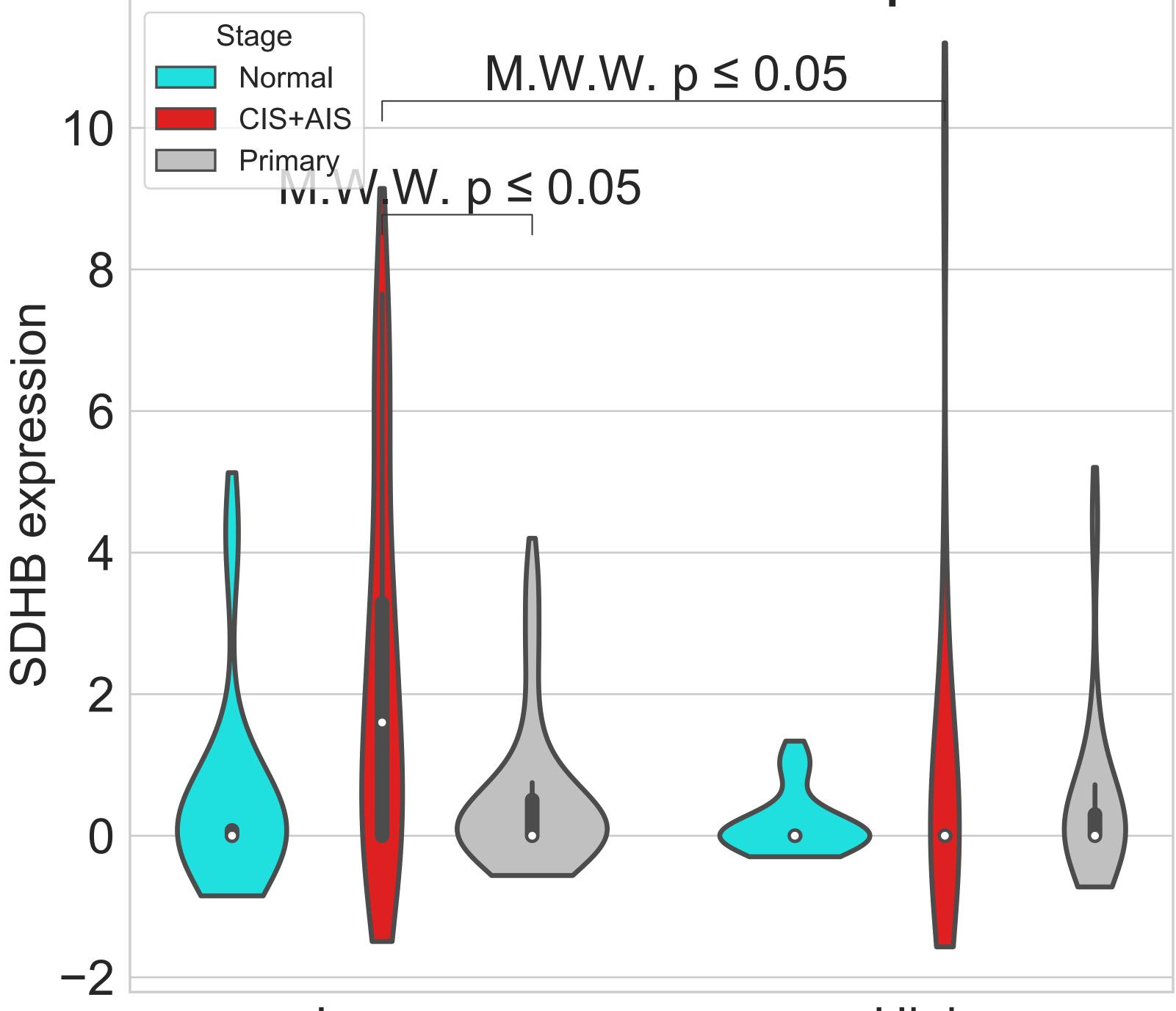
SDHB: Kruskal-Wallis p=0.044



Lower Higher Mutation Shared Count per TMB