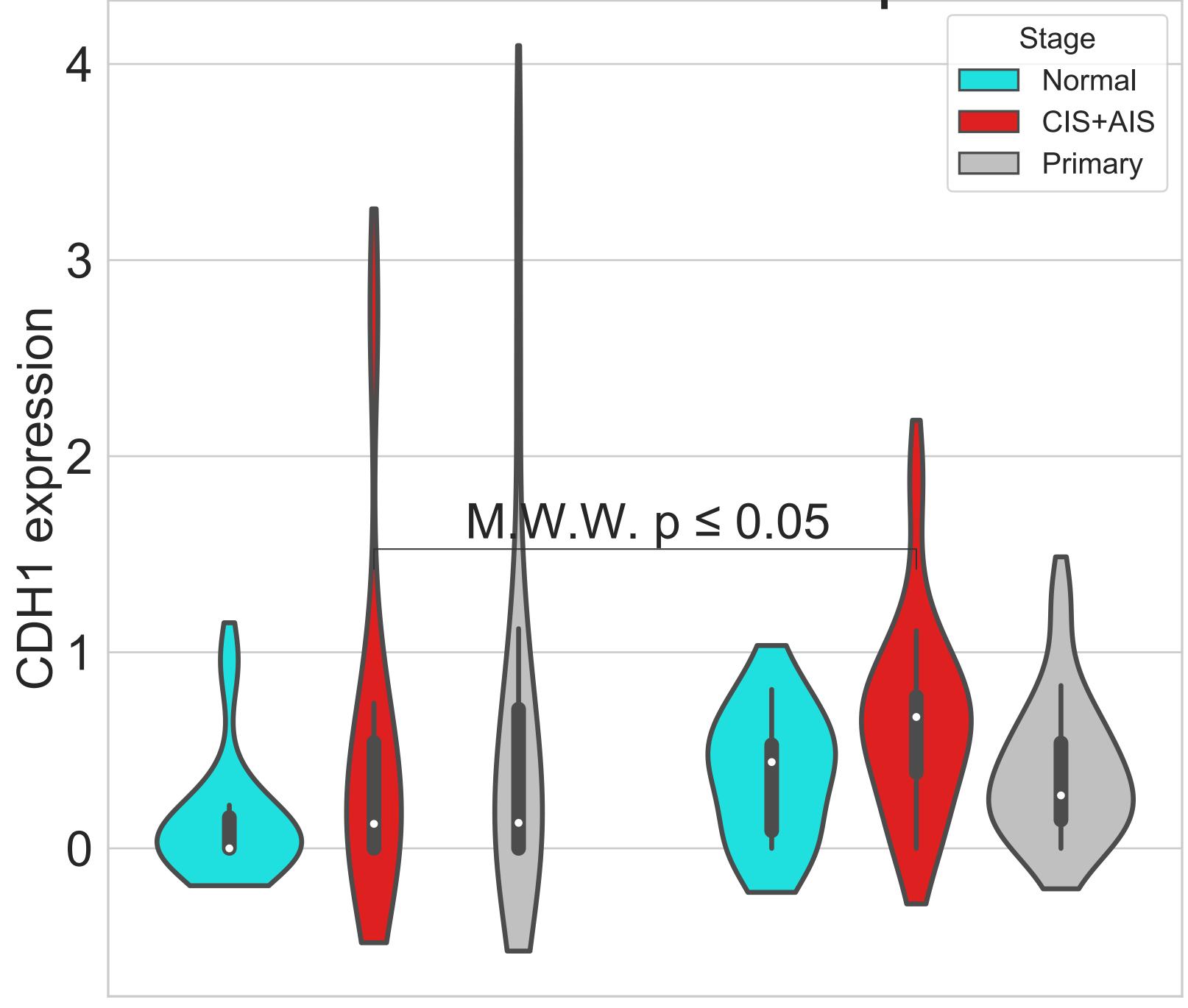
CDH1: Kruskal-Wallis p=0.042



Lower Higher Mutation Shared Count per TMB