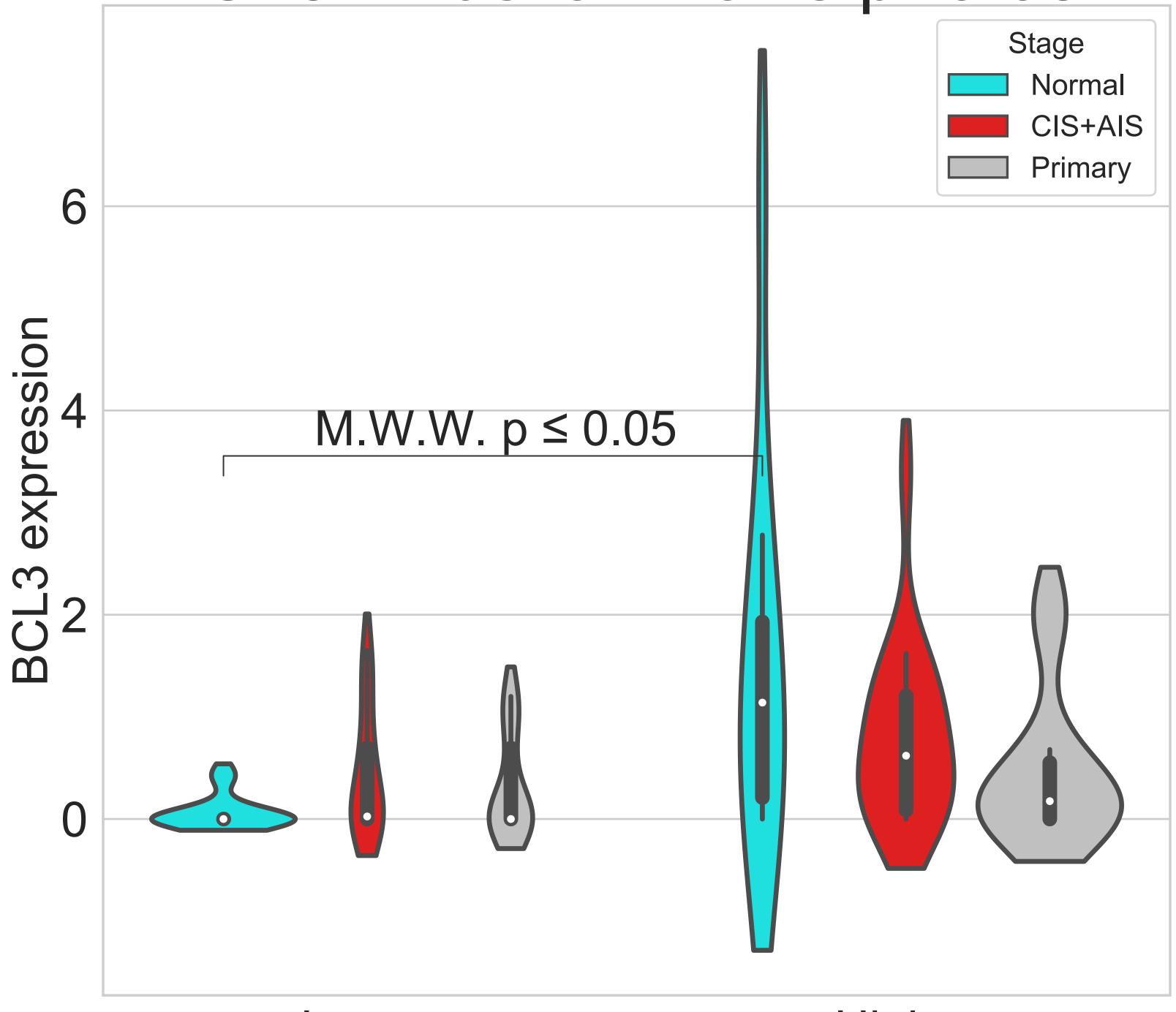
BCL3: Kruskal-Wallis p=0.032



Lower Higher Mutation Shared Count per TMB