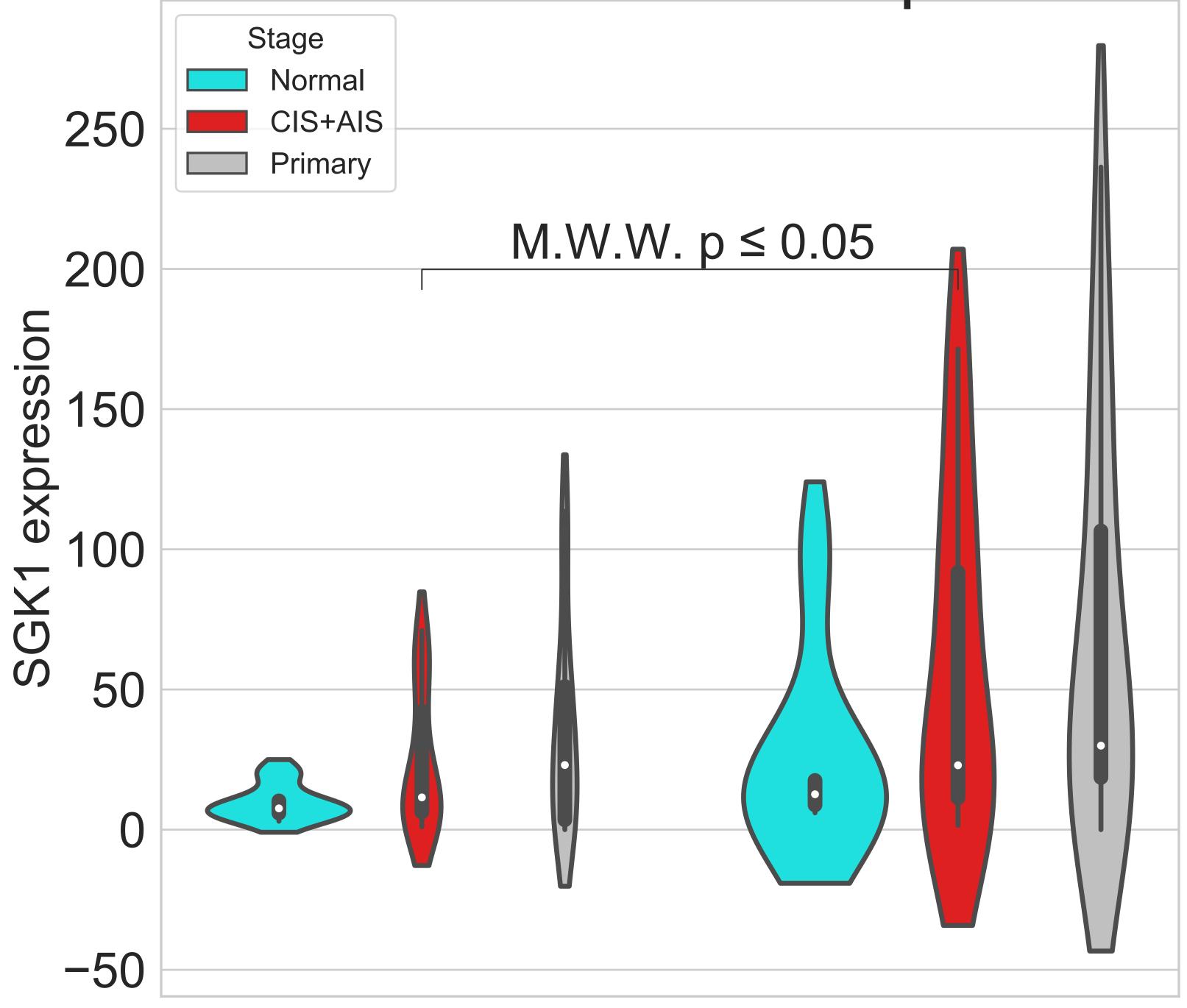
SGK1: Kruskal-Wallis p=0.045



Lower Higher Mutation Shared Count per TMB