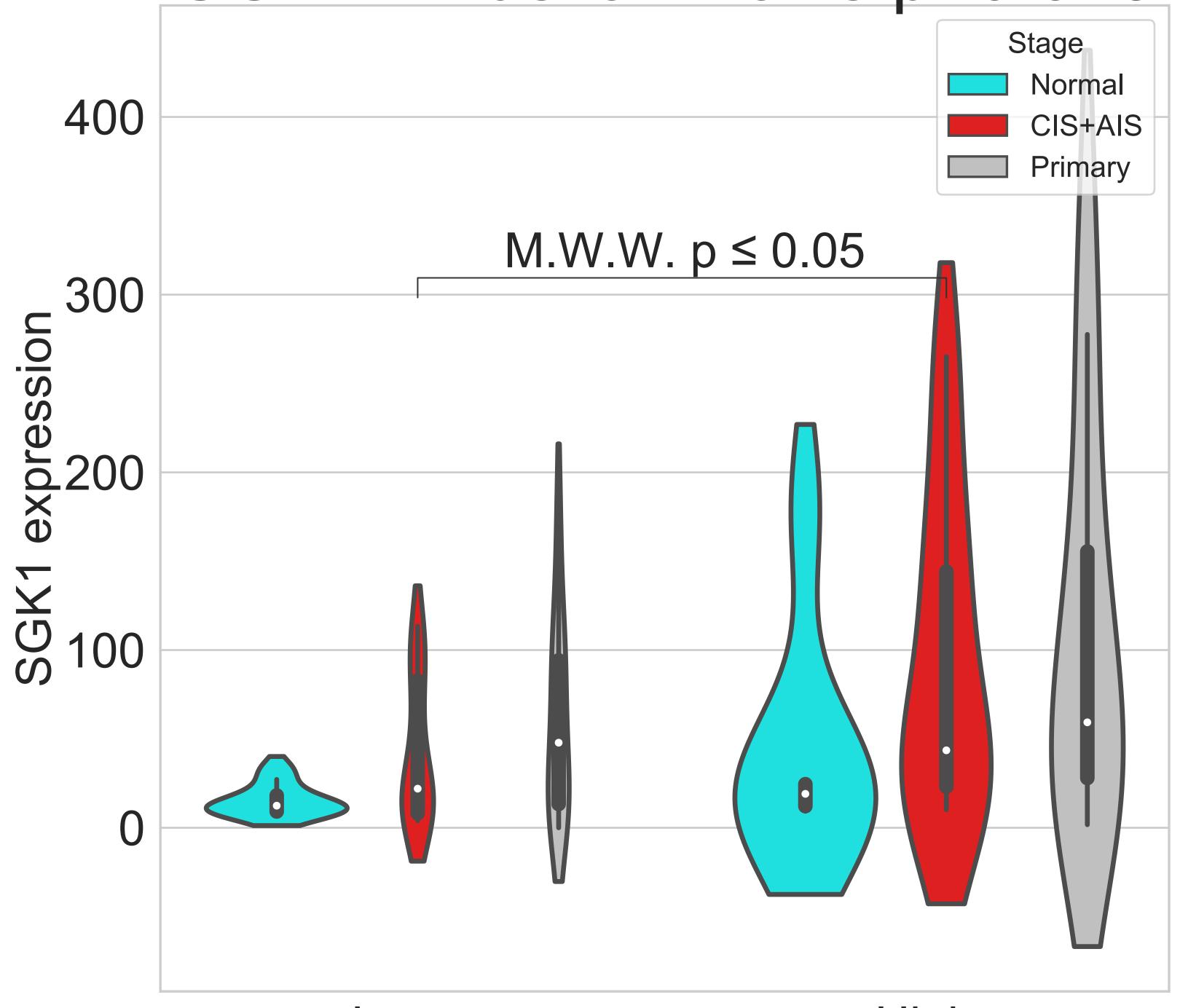
SGK1: Kruskal-Wallis p=0.023



Lower Higher Mutation Shared Count per TMB