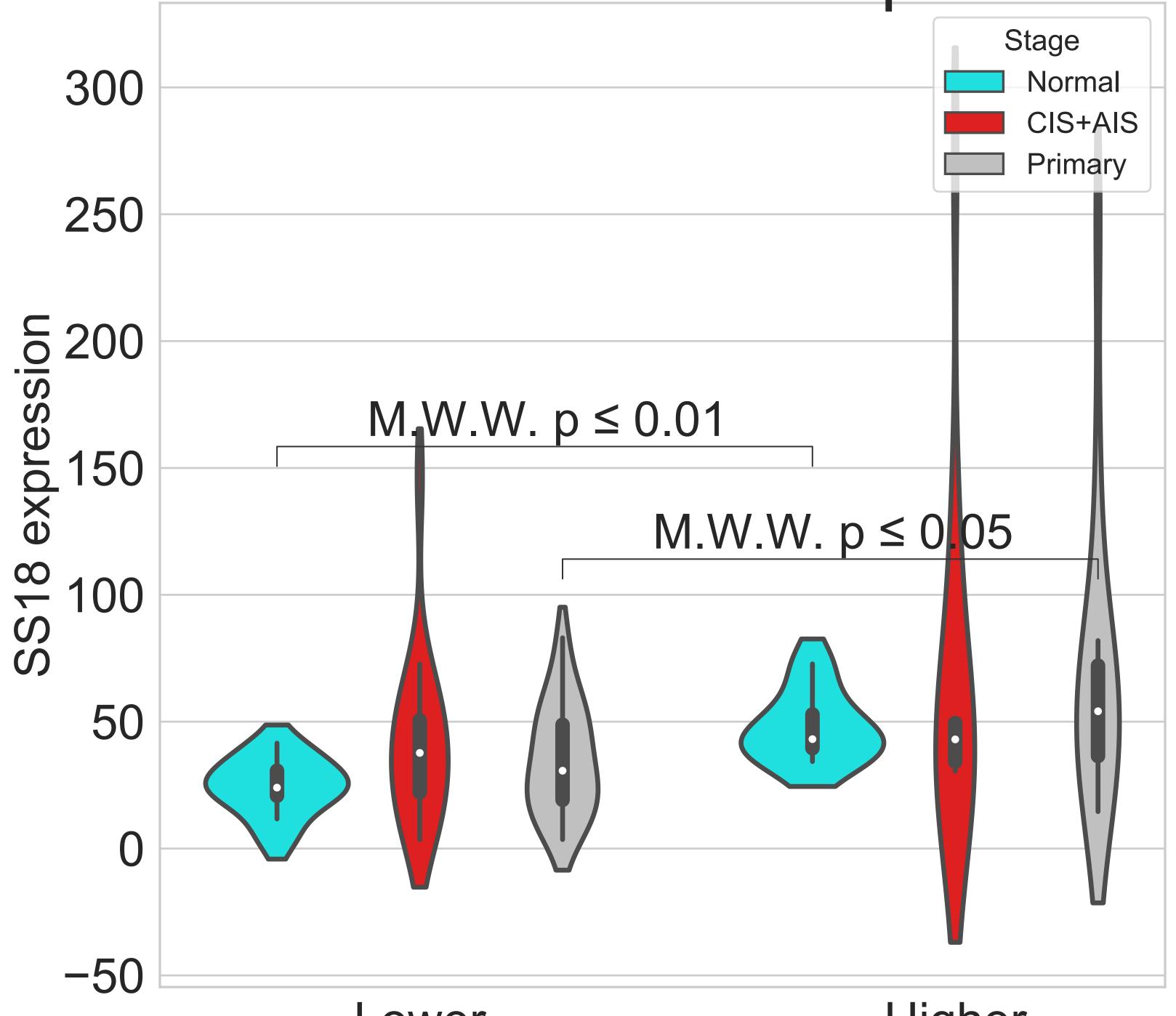
SS18: Kruskal-Wallis p=0.024



Lower Higher Mutation Shared Count per TMB