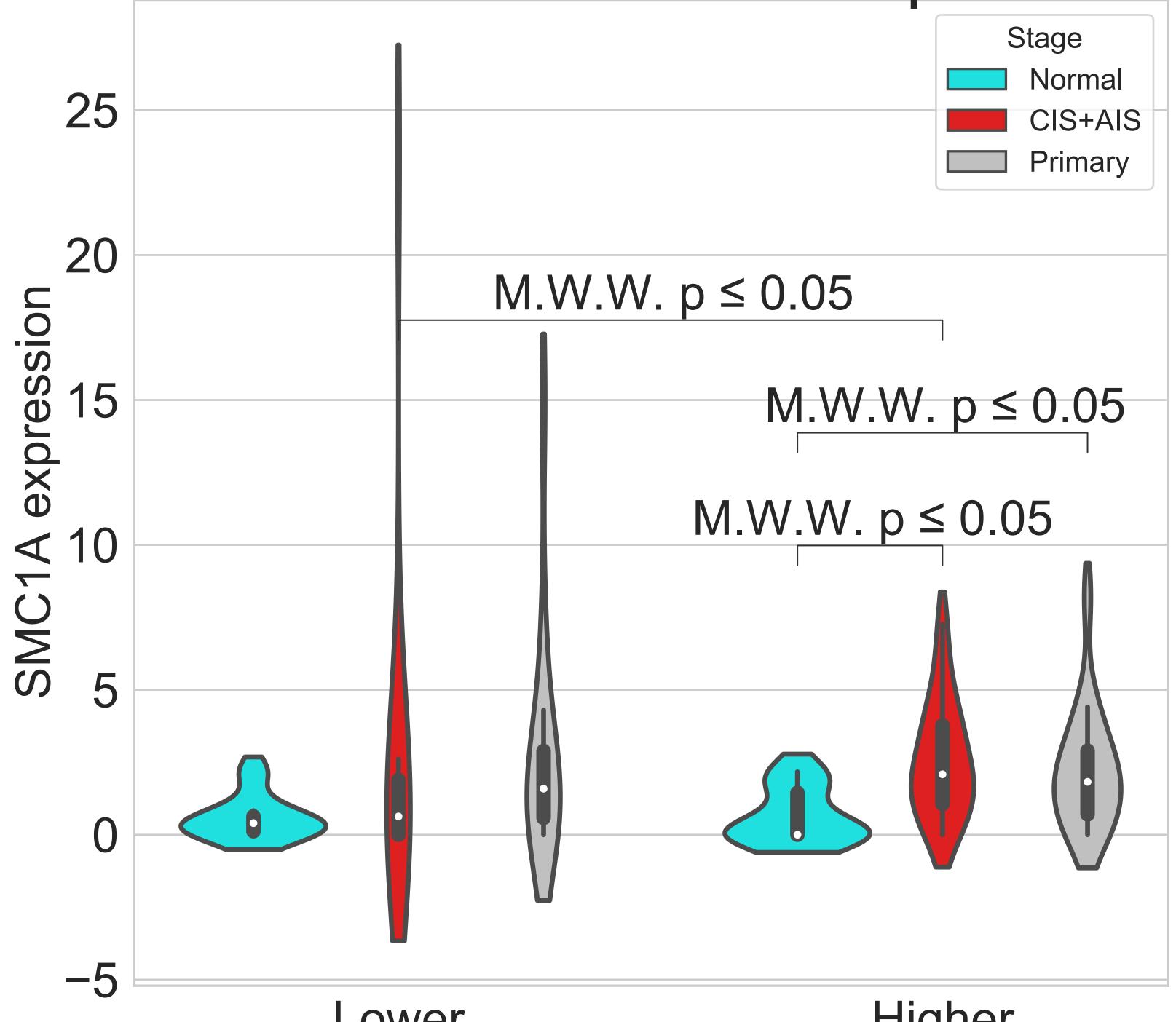
SMC1A: Kruskal-Wallis p=0.022



Lower
Mutation Shared Count per TMB