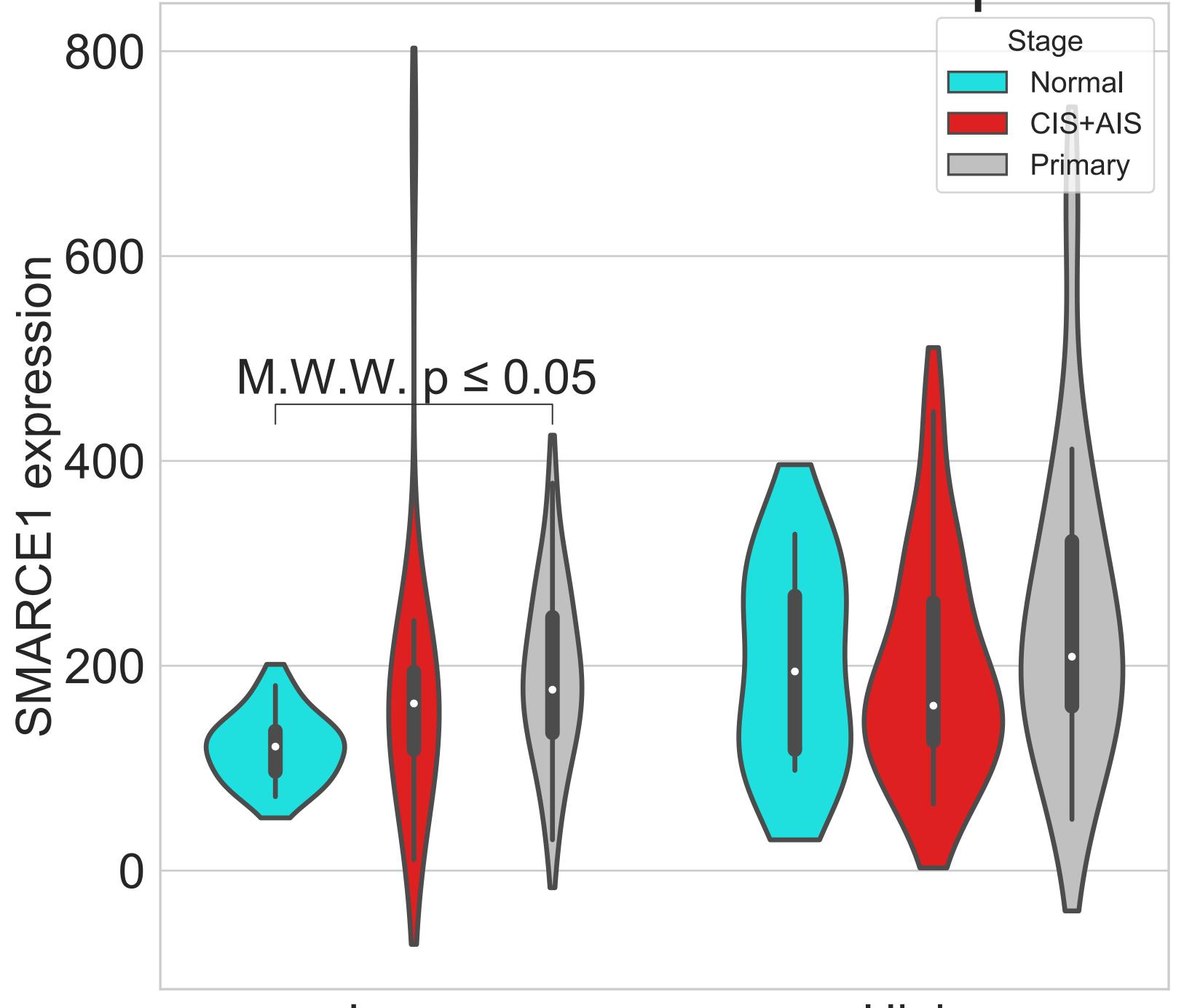
SMARCE1: Kruskal-Wallis p=0.039



Lower Higher Mutation Shared Count per TMB