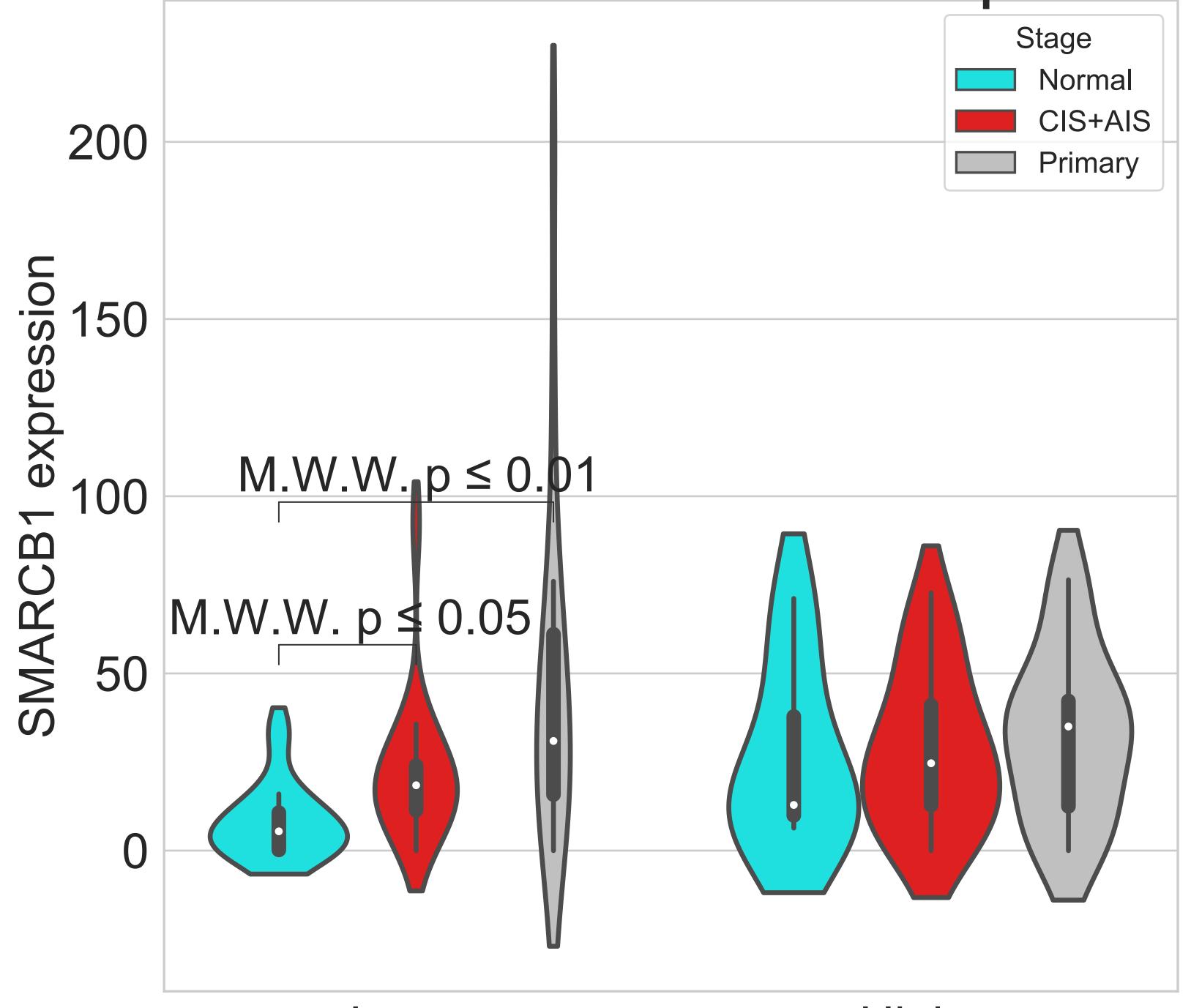
SMARCB1: Kruskal-Wallis p=0.015



Lower Higher Mutation Shared Count per TMB