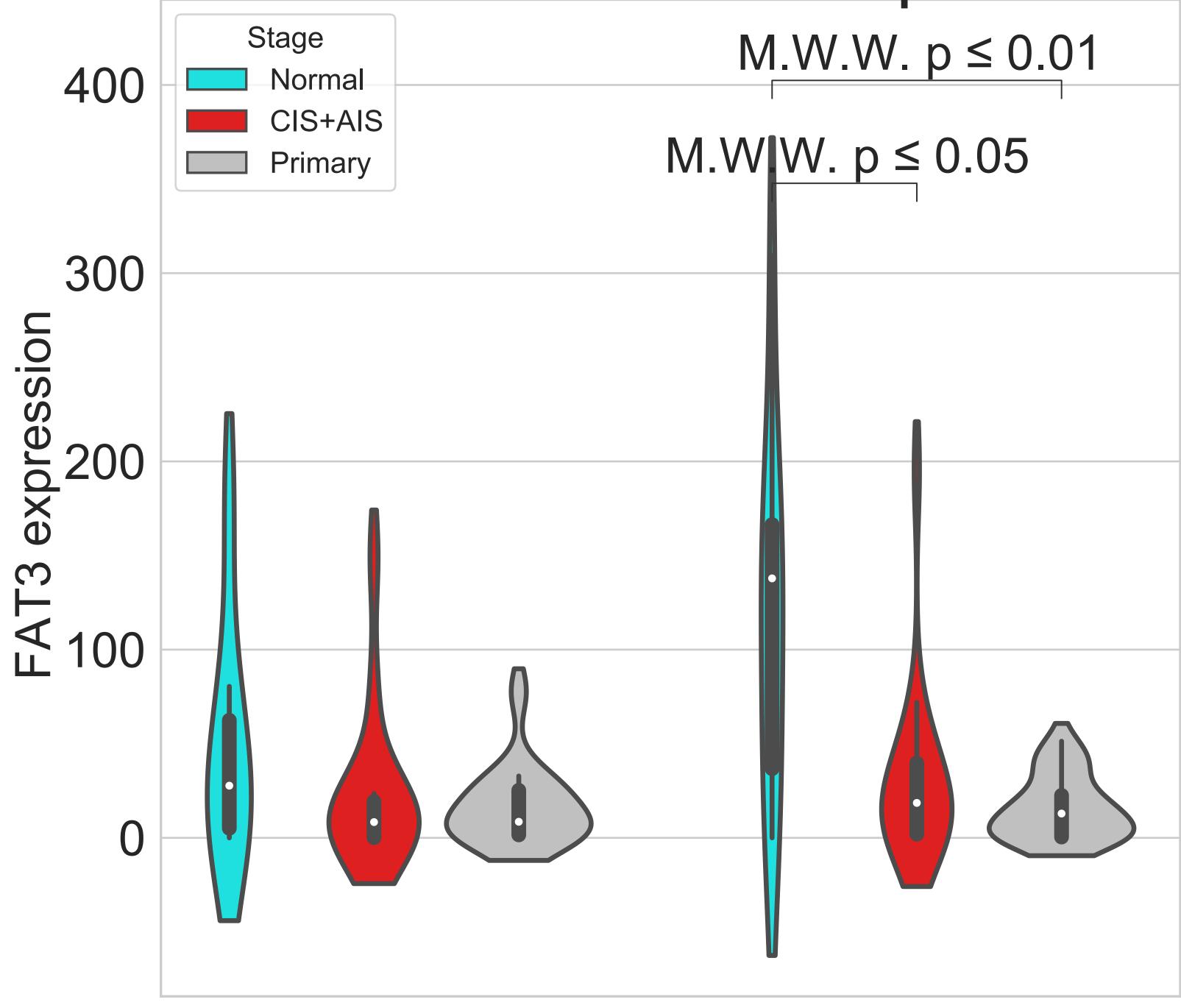
FAT3: Kruskal-Wallis p=0.049



Lower
Mutation Shared Count per TMB