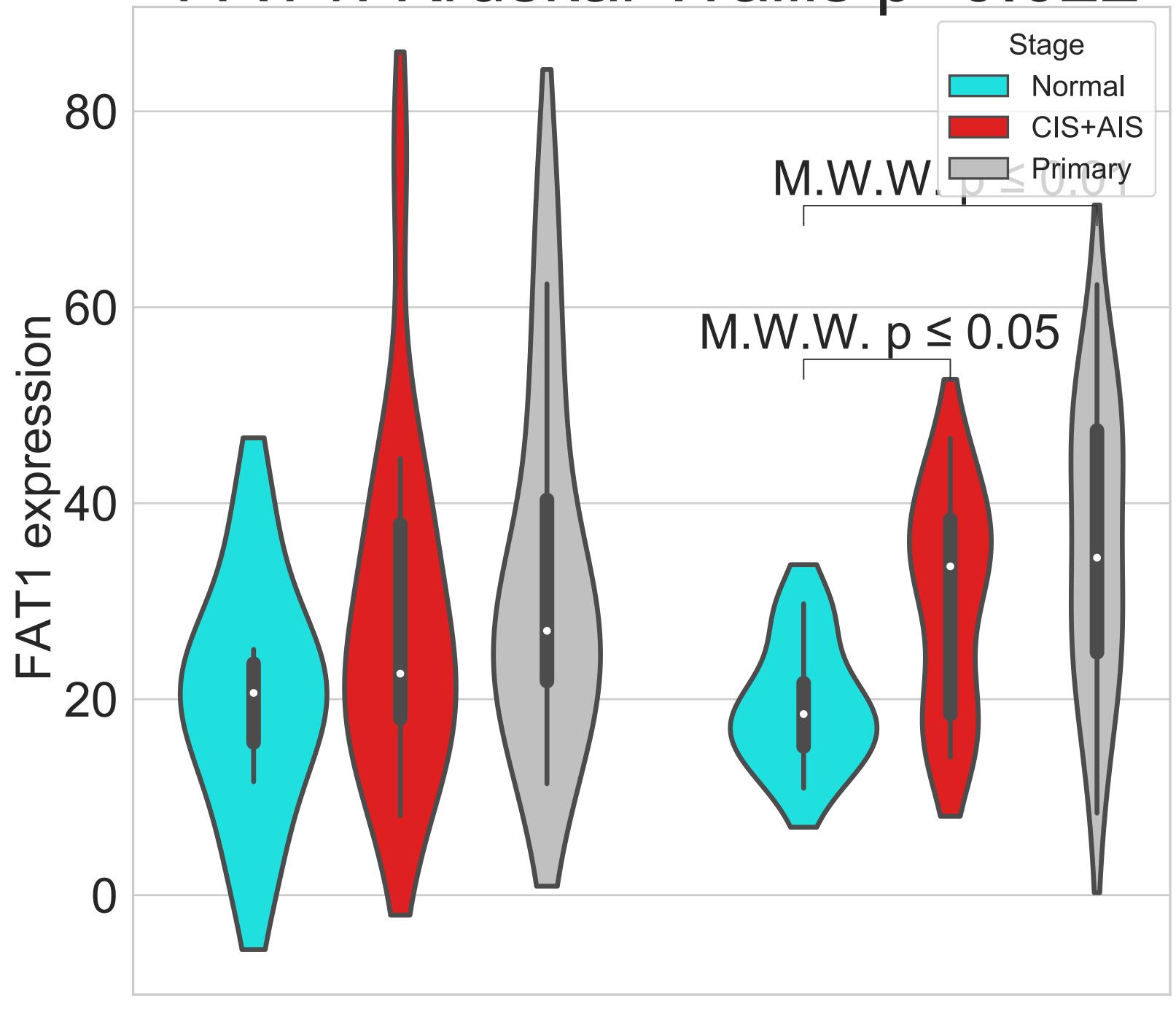
FAT1: Kruskal-Wallis p=0.022



Lower Higher Mutation Shared Proportion