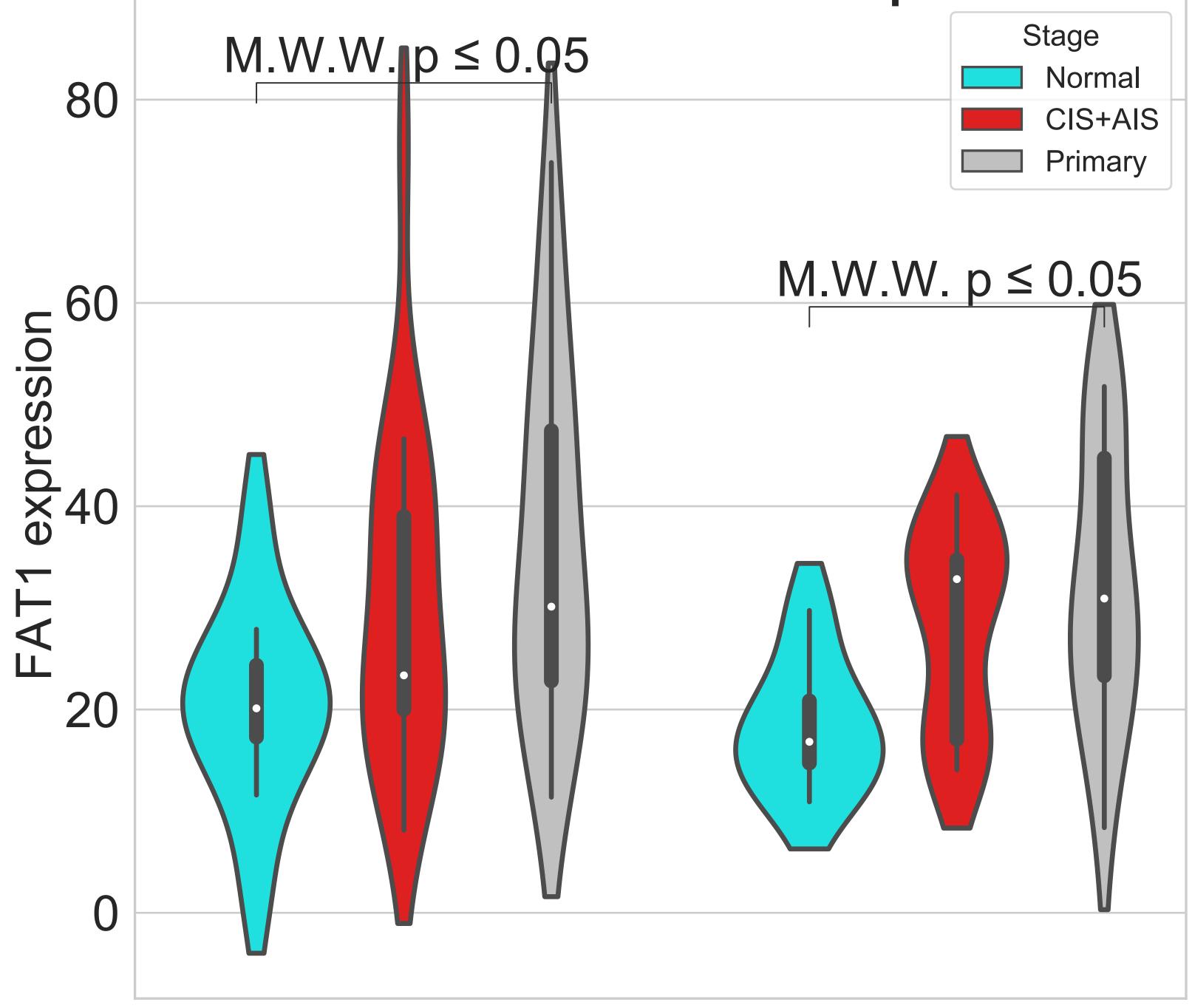
FAT1: Kruskal-Wallis p=0.033



Lower Higher Mutation Shared Proportion