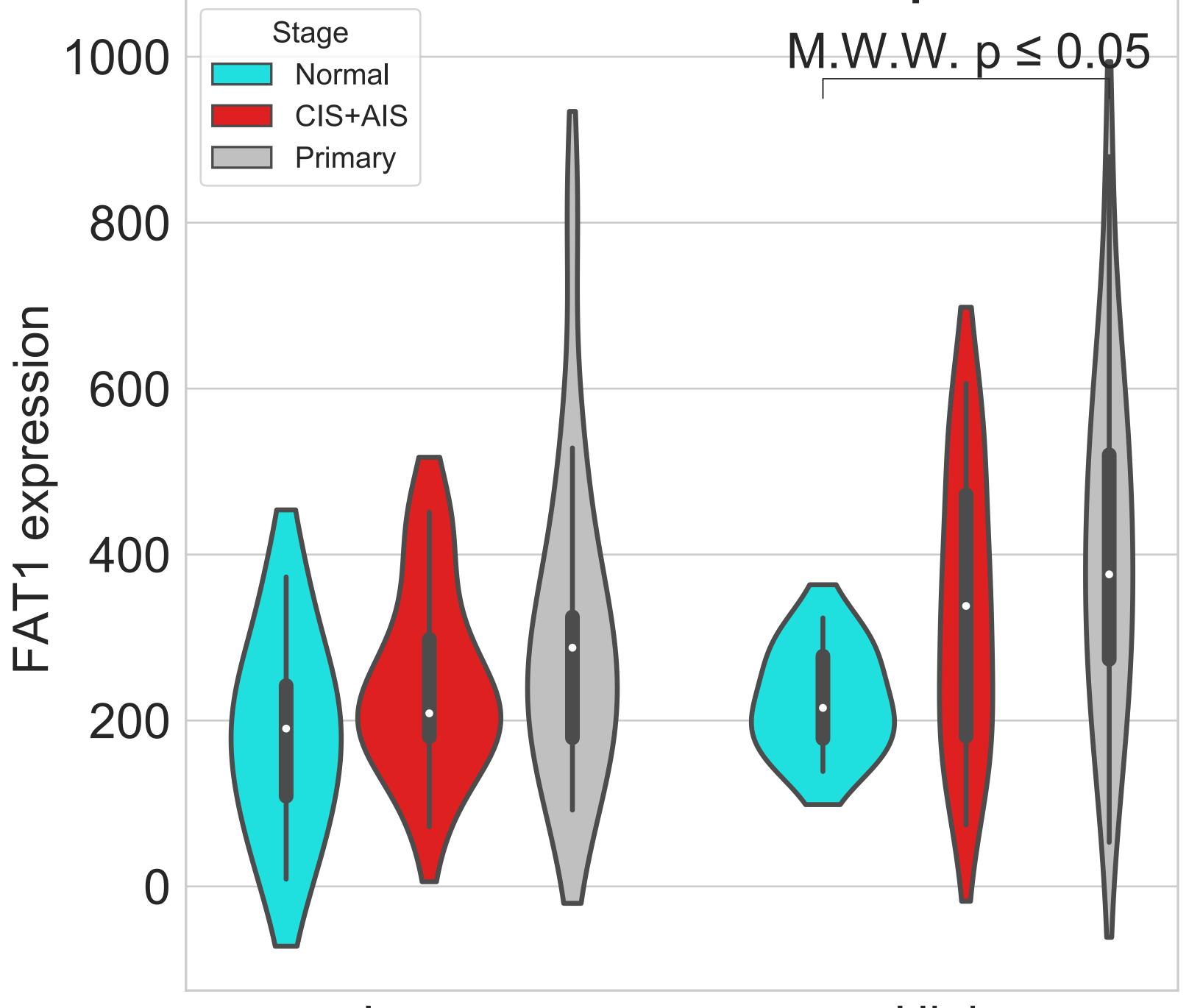
FAT1: Kruskal-Wallis p=0.025



Lower Higher Mutation Shared Proportion (SYN)