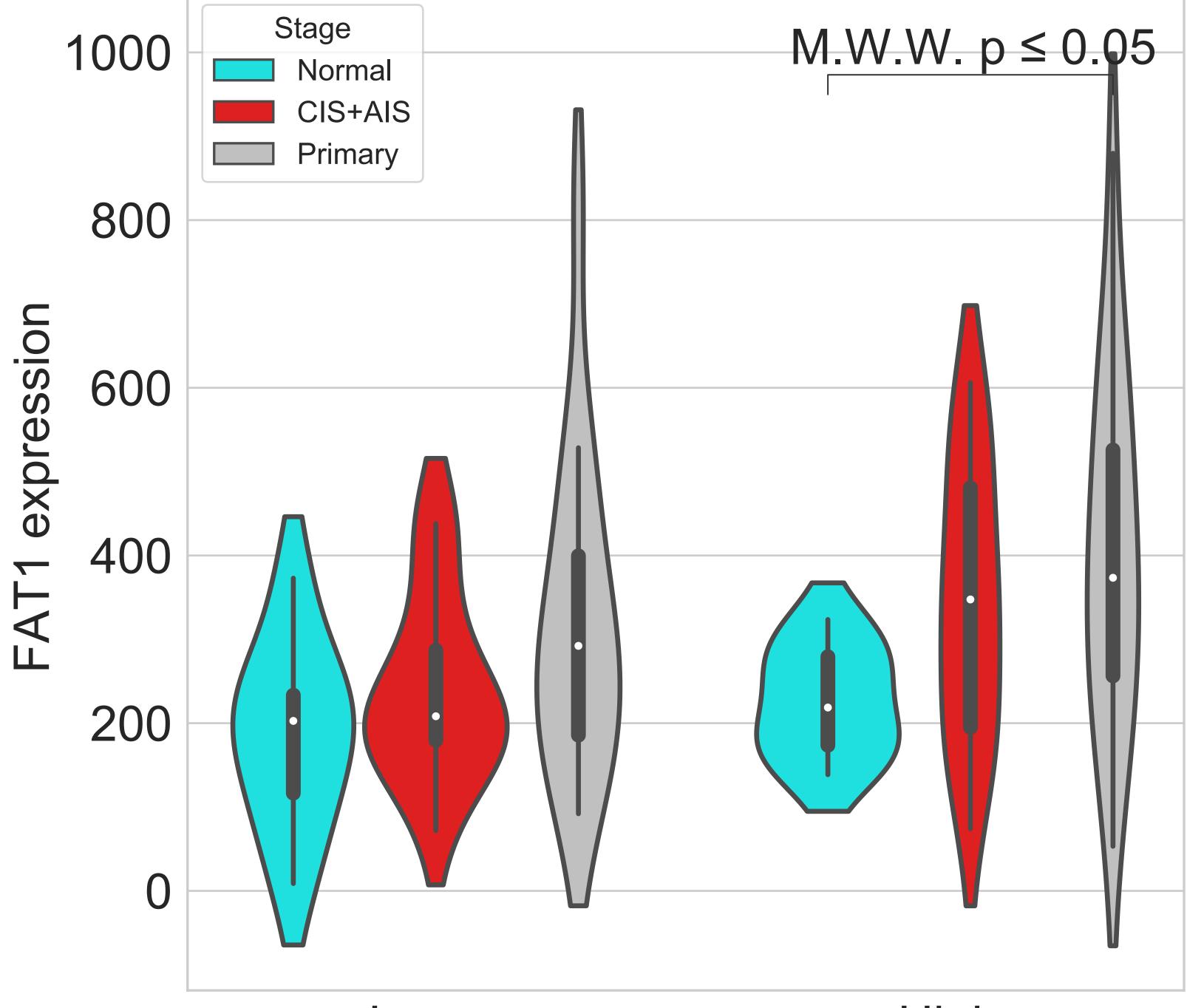
FAT1: Kruskal-Wallis p=0.021



Lower Higher Mutation Shared Proportion (Union & SYN)