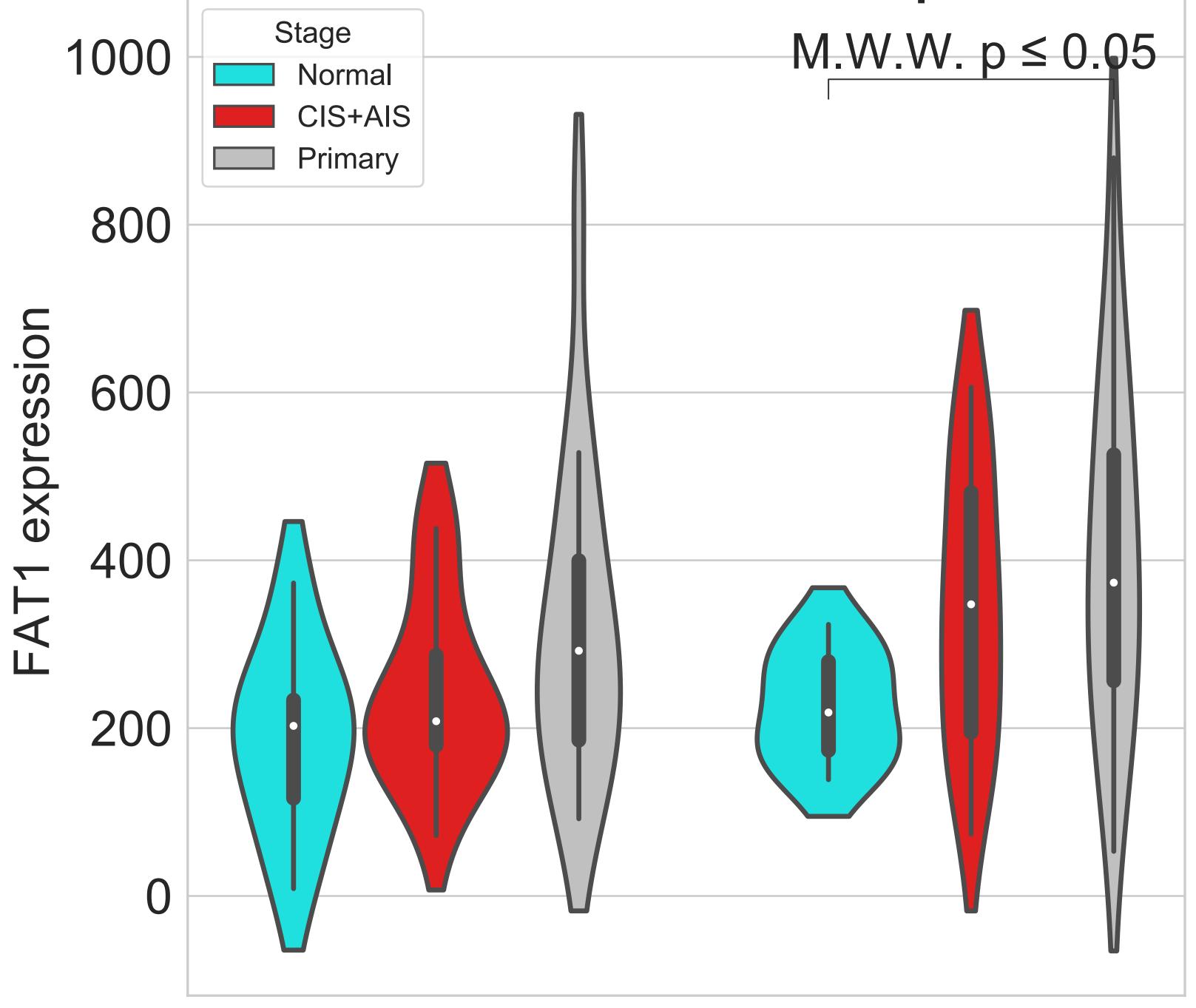
FAT1: Kruskal-Wallis p=0.021



Lower
Mutation Shared Proportion (Union)