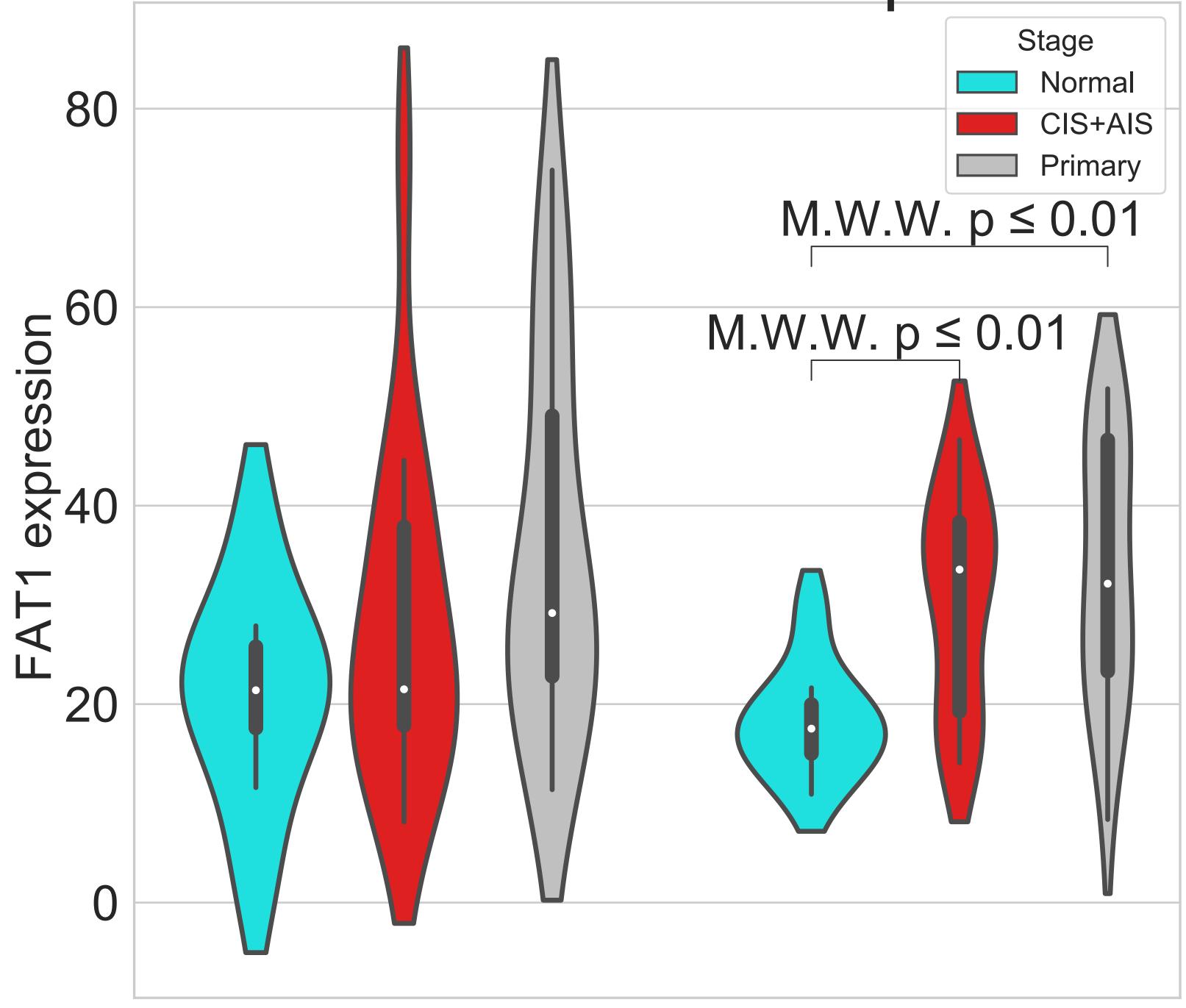
FAT1: Kruskal-Wallis p=0.022



Lower
Mutation Shared Proportion (Union)