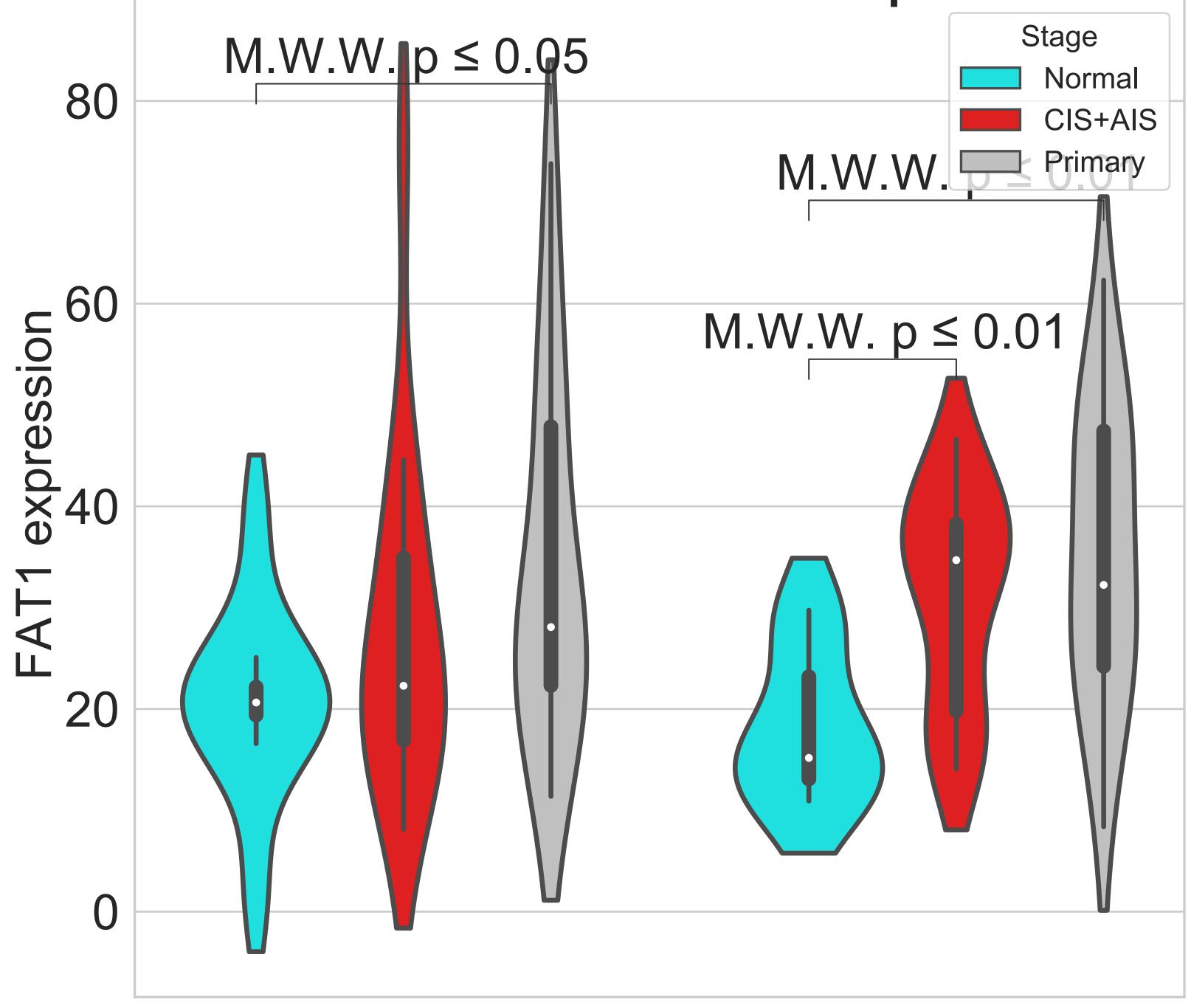
FAT1: Kruskal-Wallis p=0.016



Lower
Mutation Shared Count (SYN)