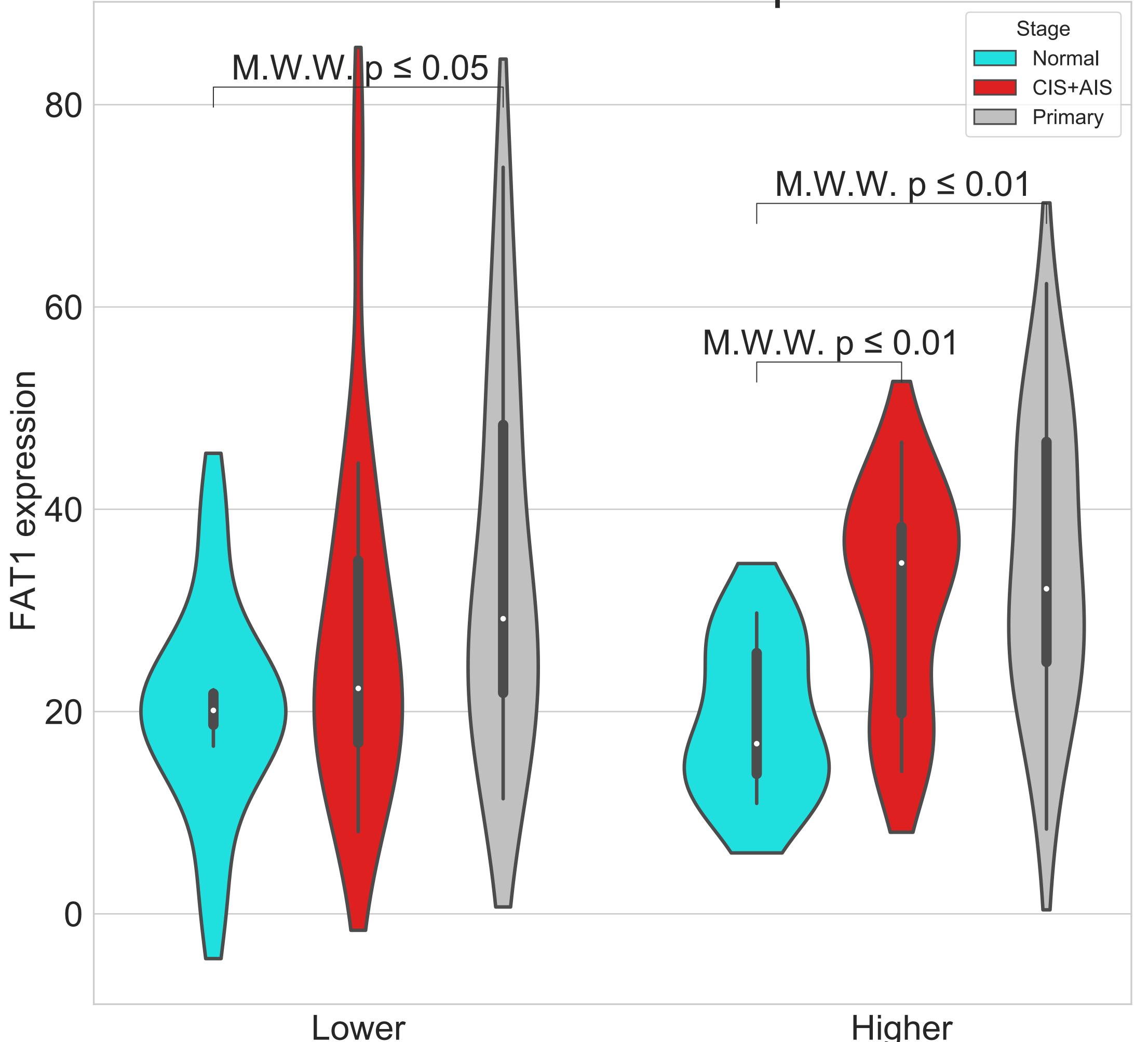
FAT1: Kruskal-Wallis p=0.019



wer
Mutation Shared Count (SYN)