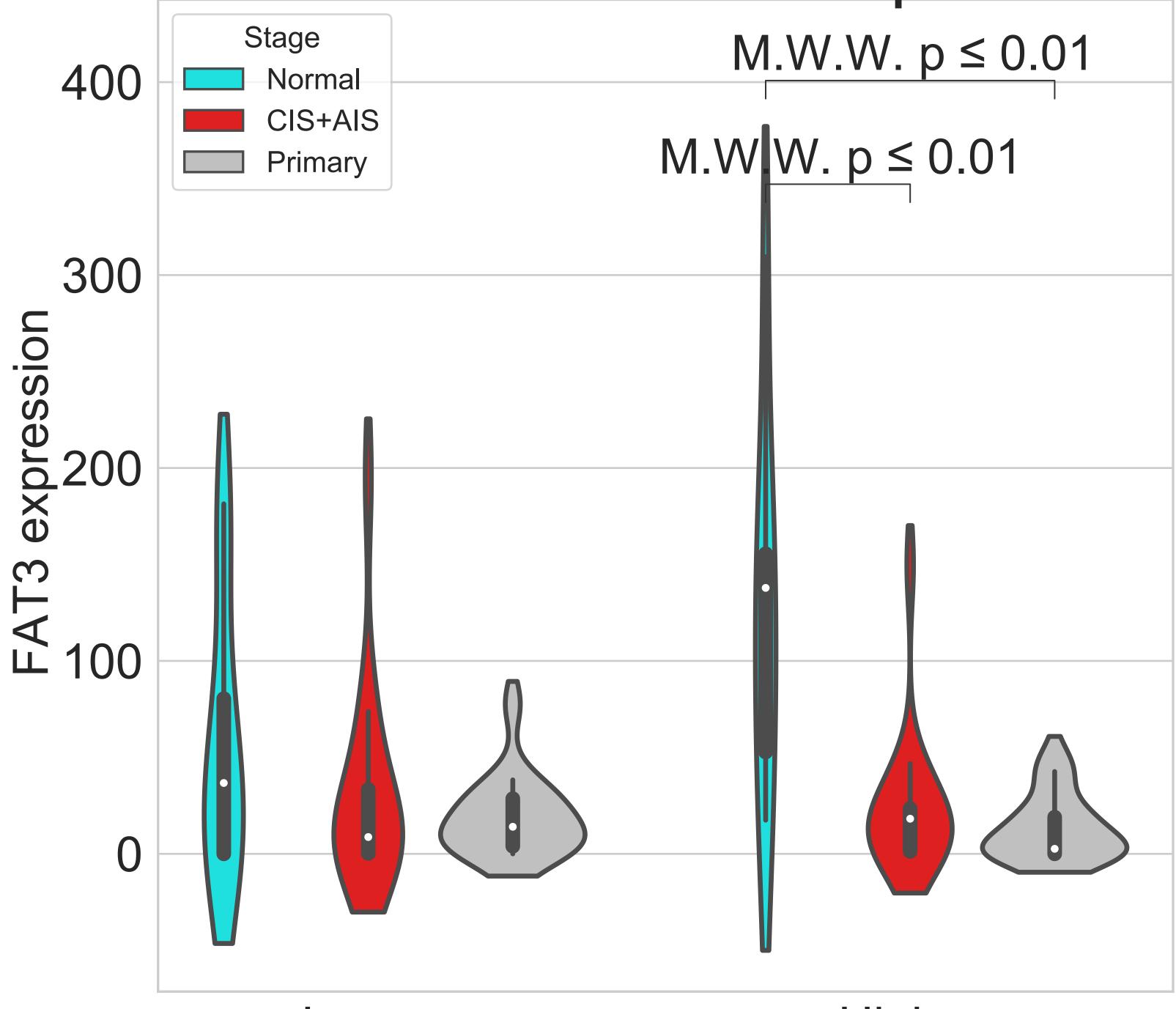
FAT3: Kruskal-Wallis p=0.027



Lower
Mutation Shared Count (SYN)