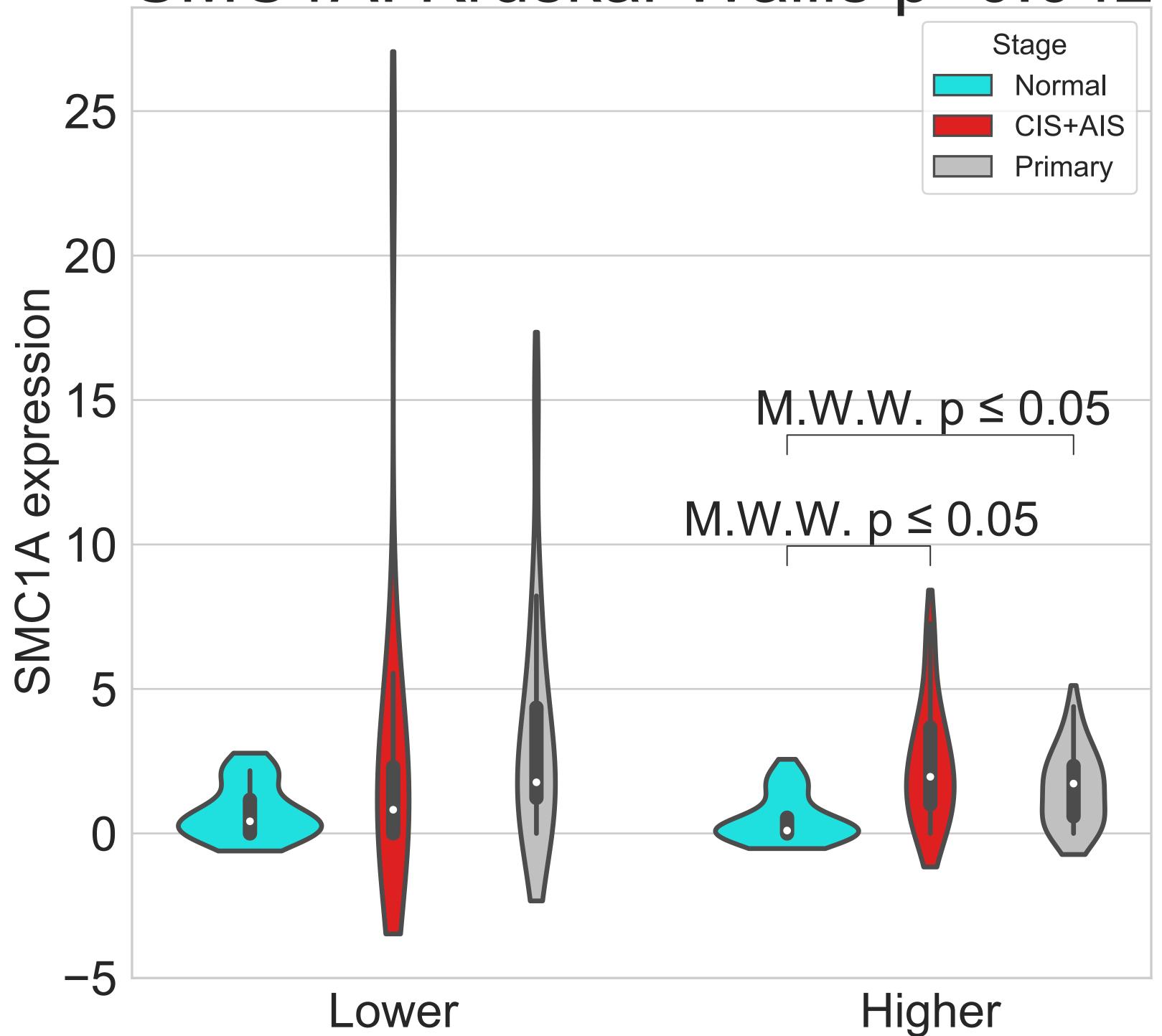
SMC1A: Kruskal-Wallis p=0.042



Mutation Shared Count (SYN)