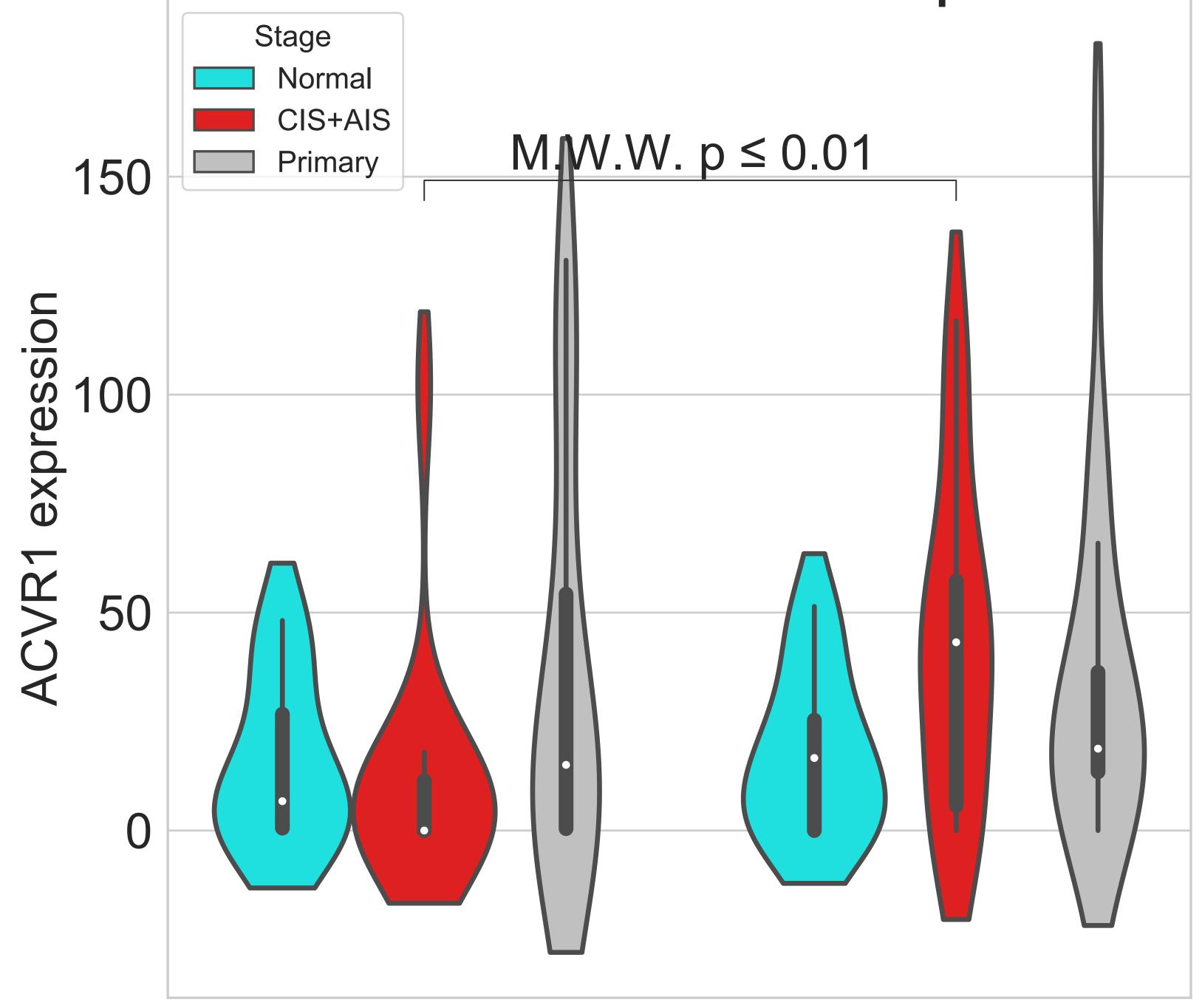
ACVR1: Kruskal-Wallis p=0.042



Lower Higher Mutation Shared Proportion (SYN)