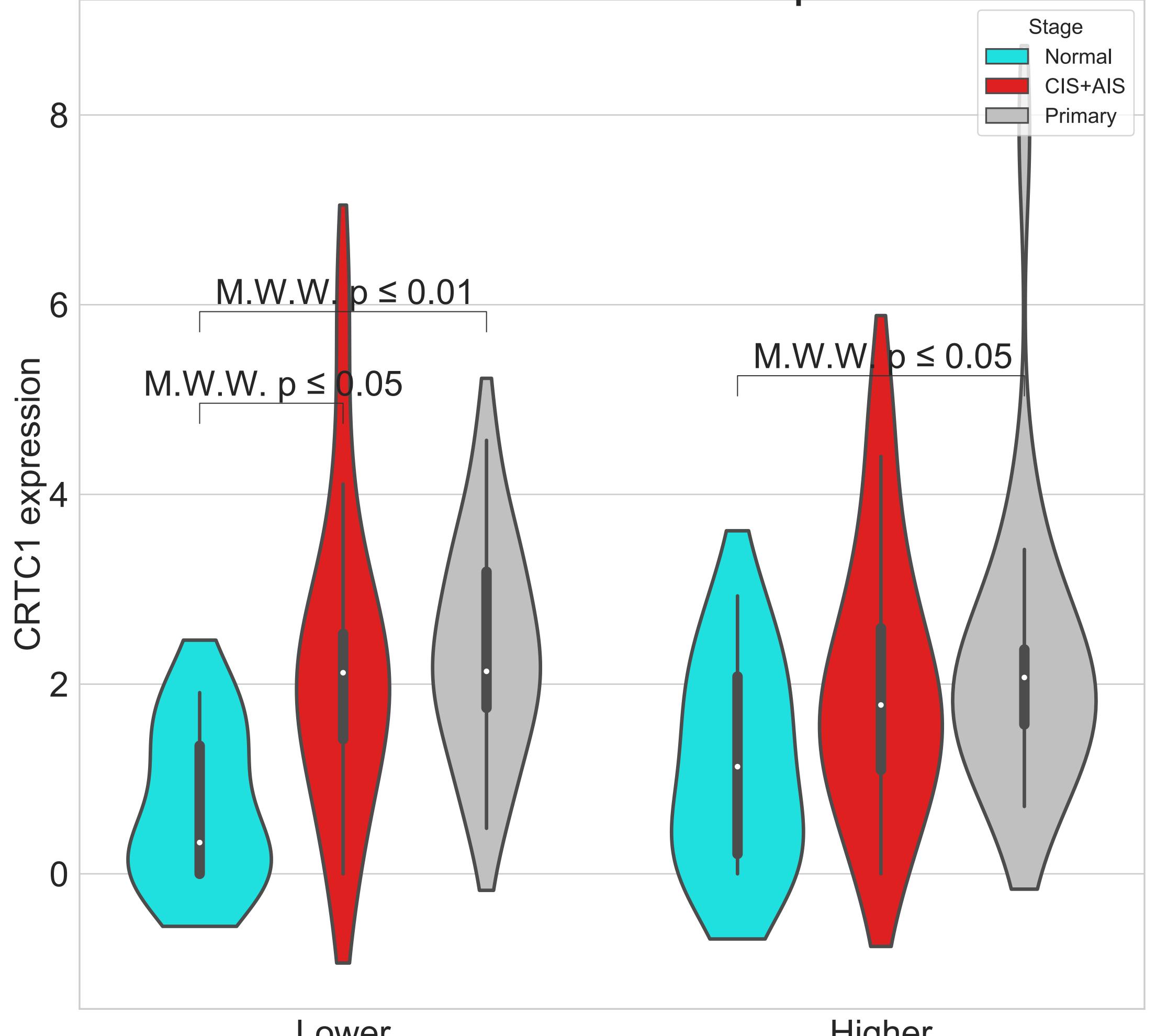
CRTC1: Kruskal-Wallis p=0.017



Higher Lower Mutation Shared Proportion (SYN)