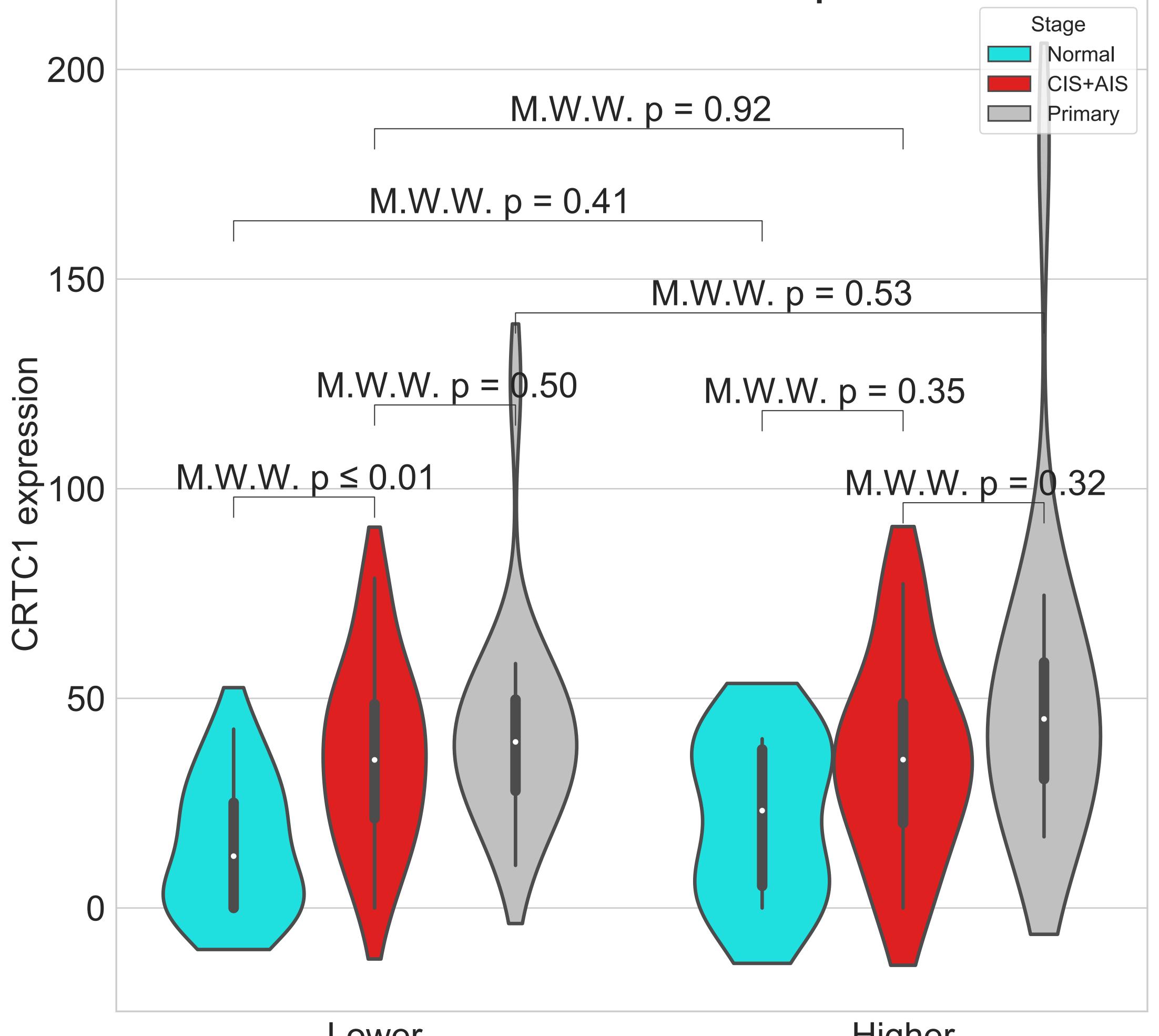
CRTC1: Kruskal-Wallis p=0.008



Higher Lower Mutation Shared Proportion (SYN)