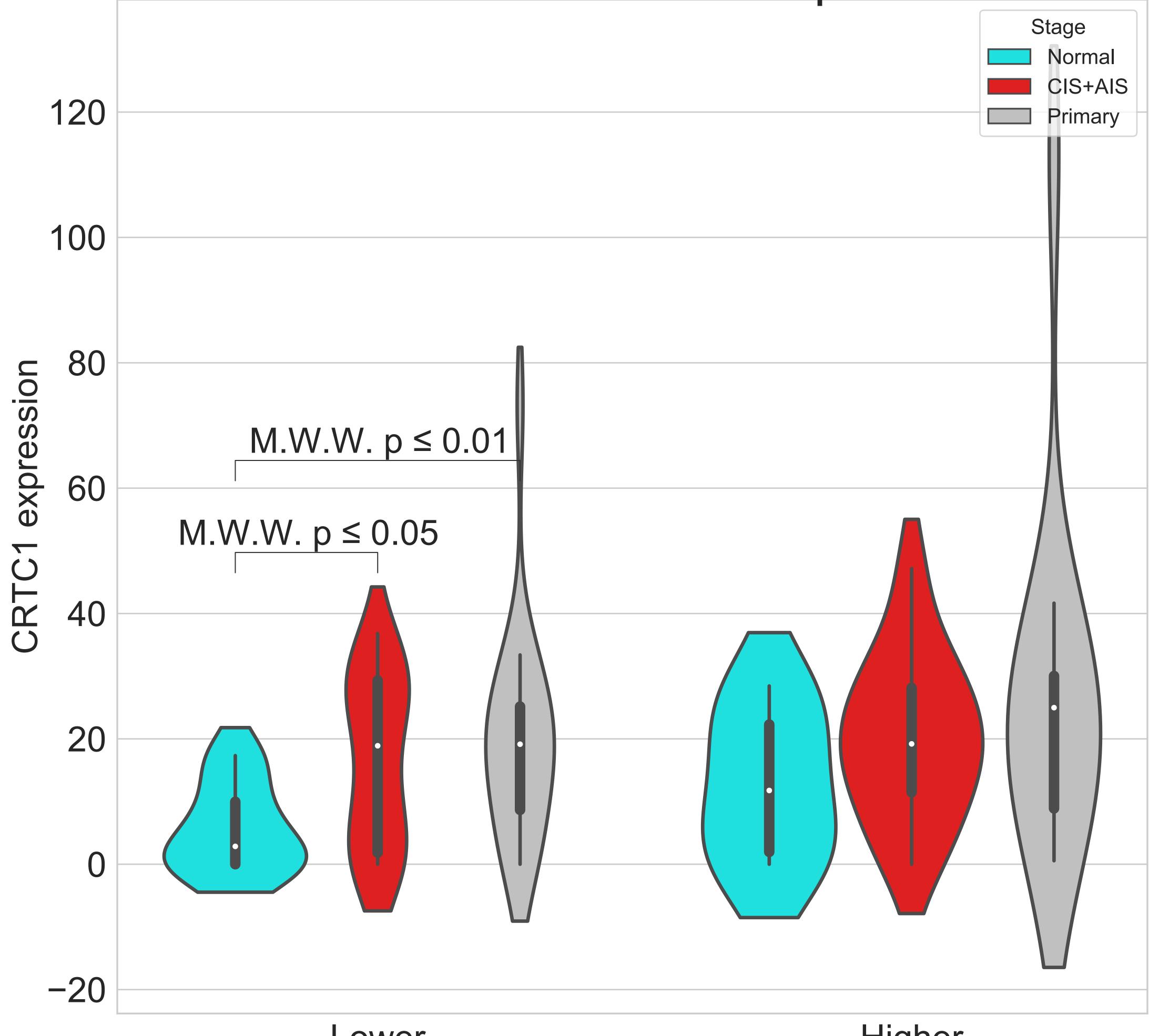
CRTC1: Kruskal-Wallis p=0.032



Higher Lower Mutation Shared Proportion (Union)