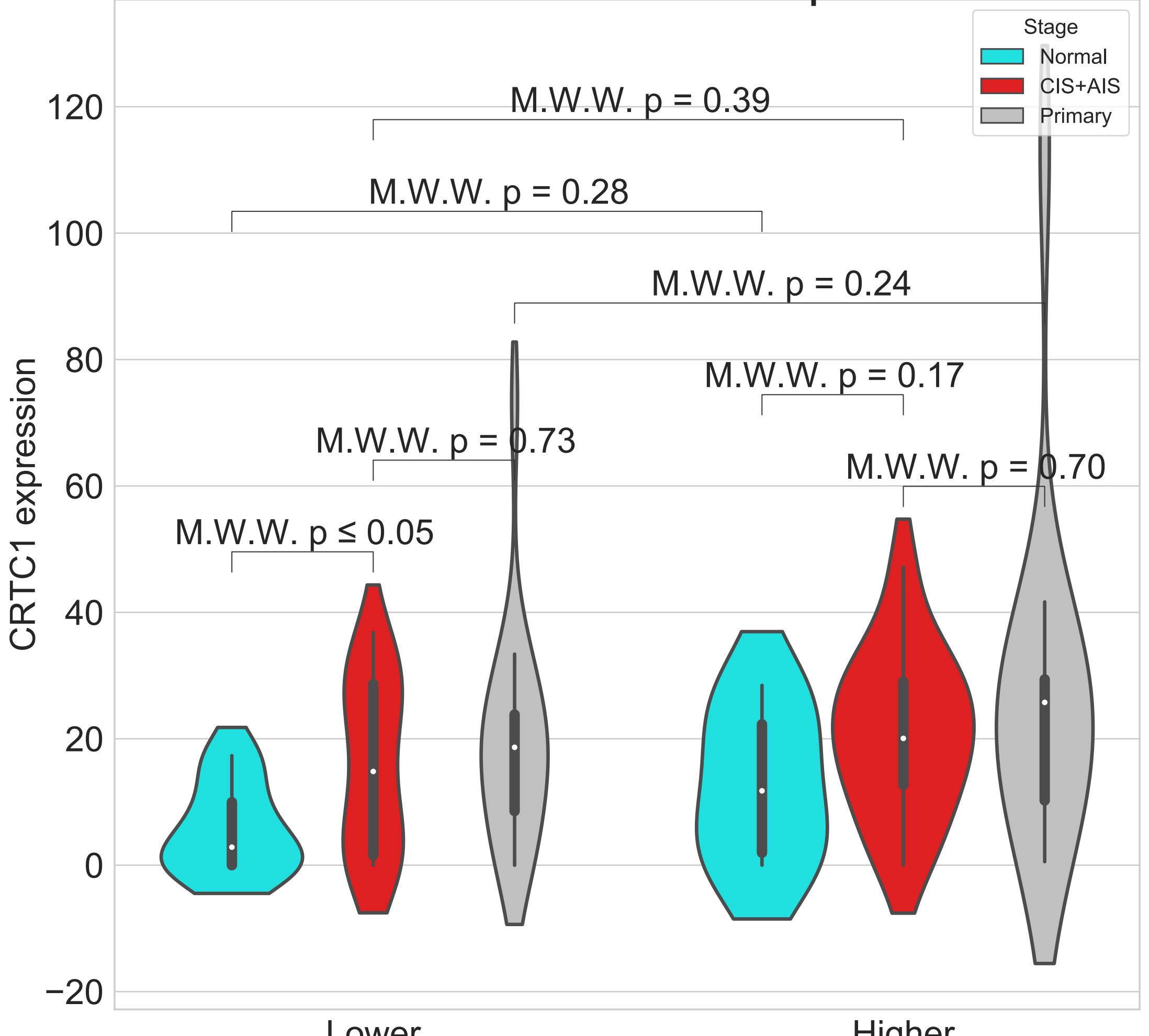
CRTC1: Kruskal-Wallis p=0.022



Higher Lower Mutation Shared Count per TMB