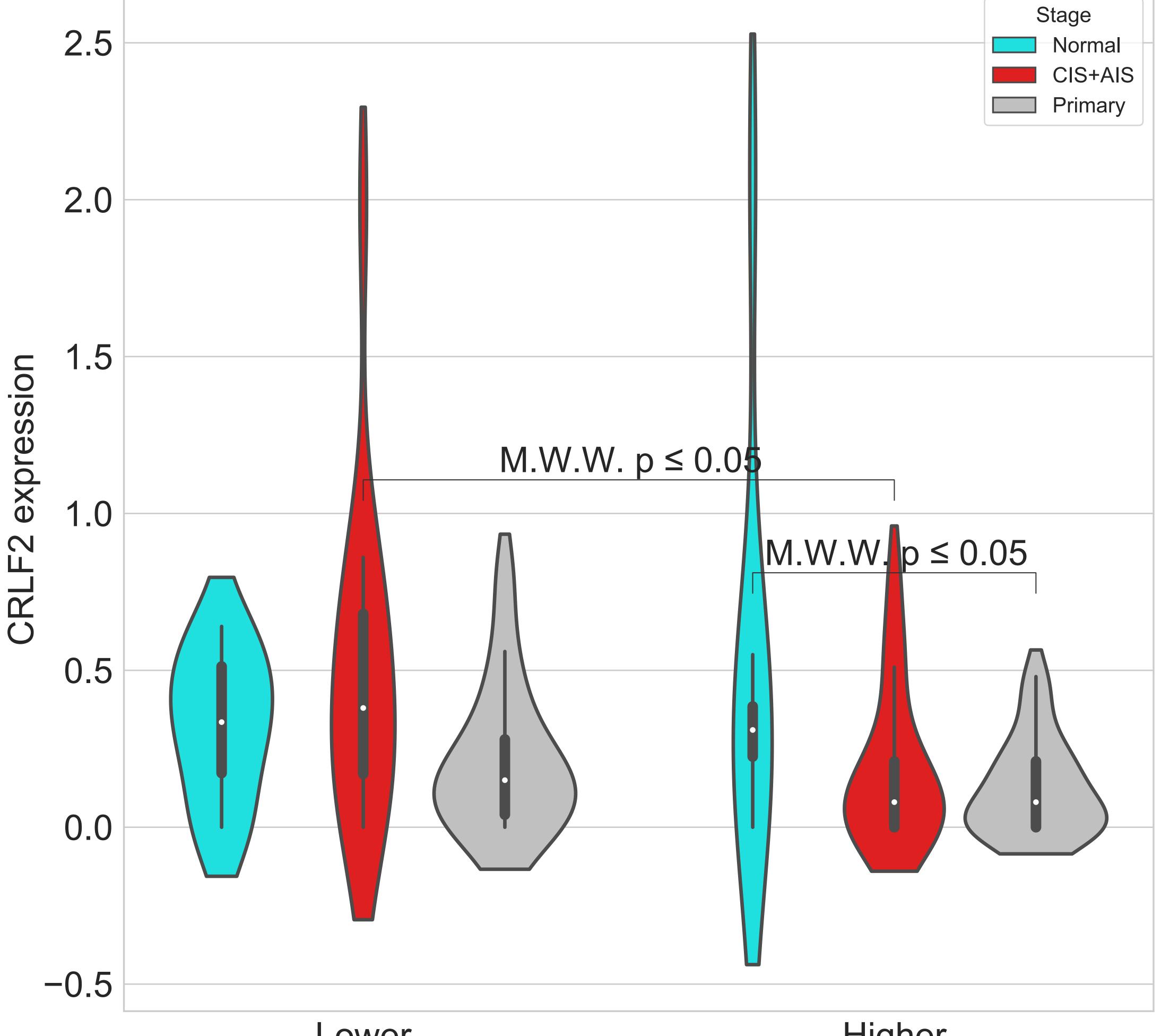
CRLF2: Kruskal-Wallis p=0.022



Higher Lower Mutation Shared Count (SYN)