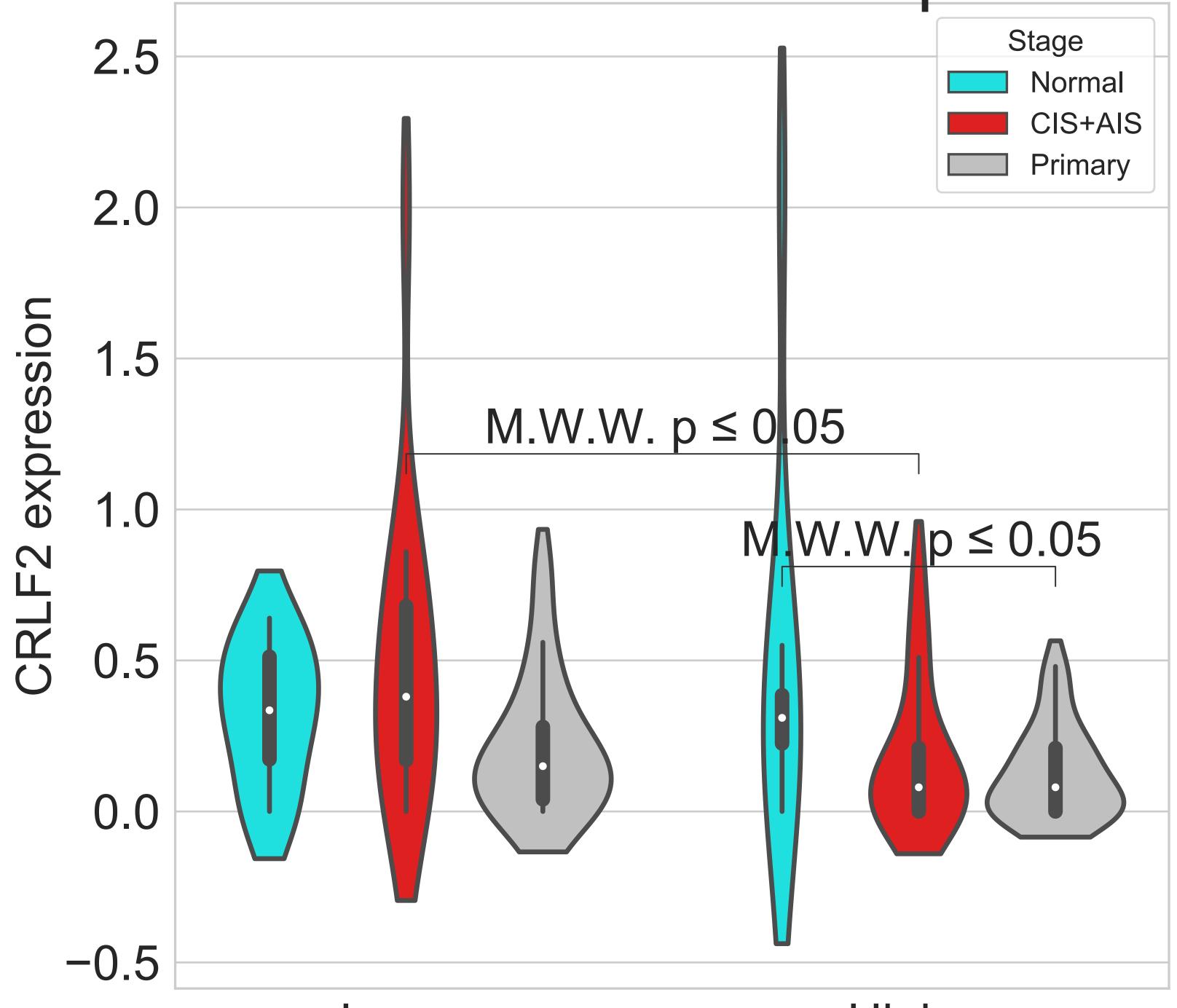
CRLF2: Kruskal-Wallis p=0.022



Lower Higher Mutation Shared Count (SYN)