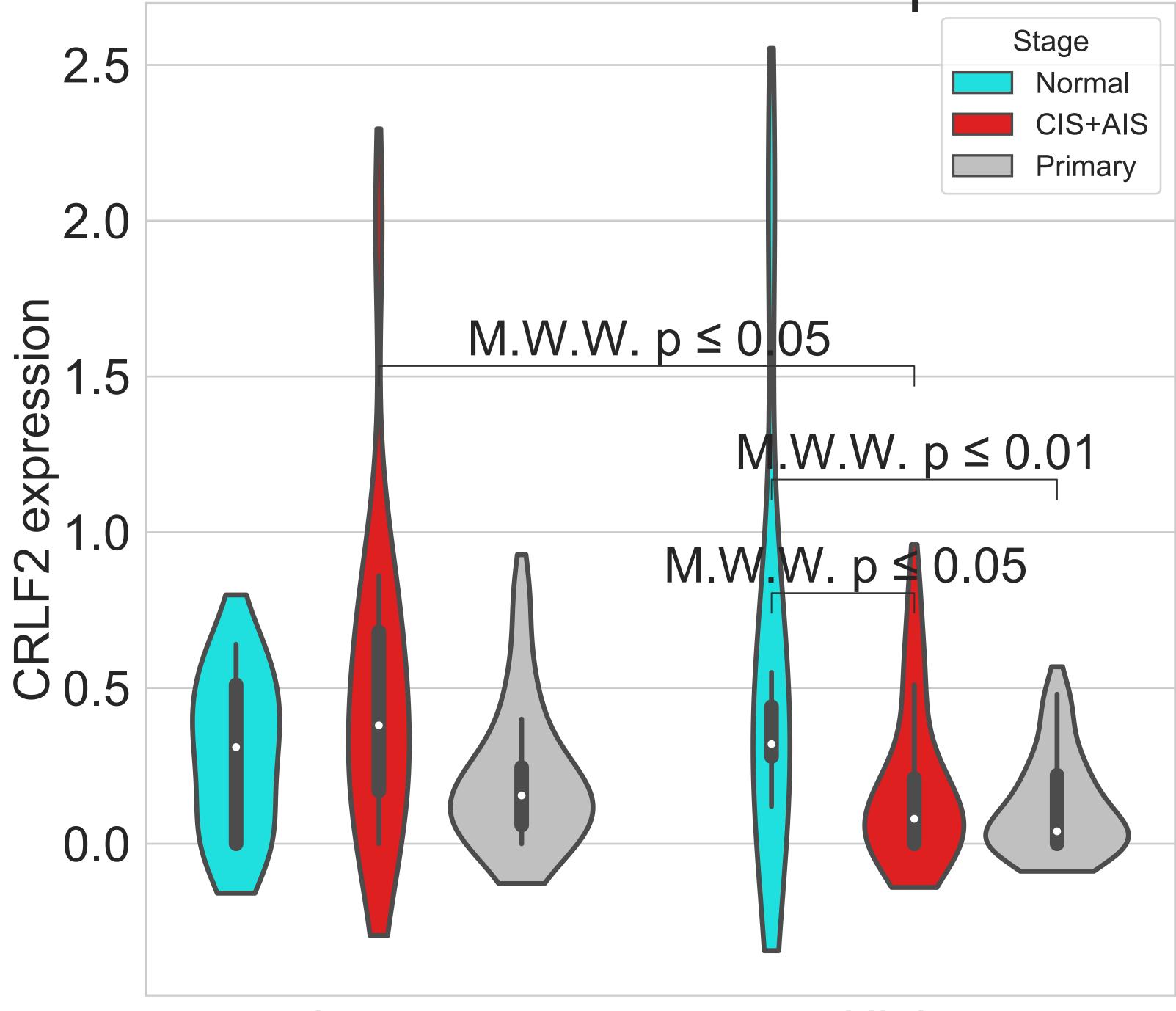
CRLF2: Kruskal-Wallis p=0.013



Lower
Mutation Shared Count (SYN)