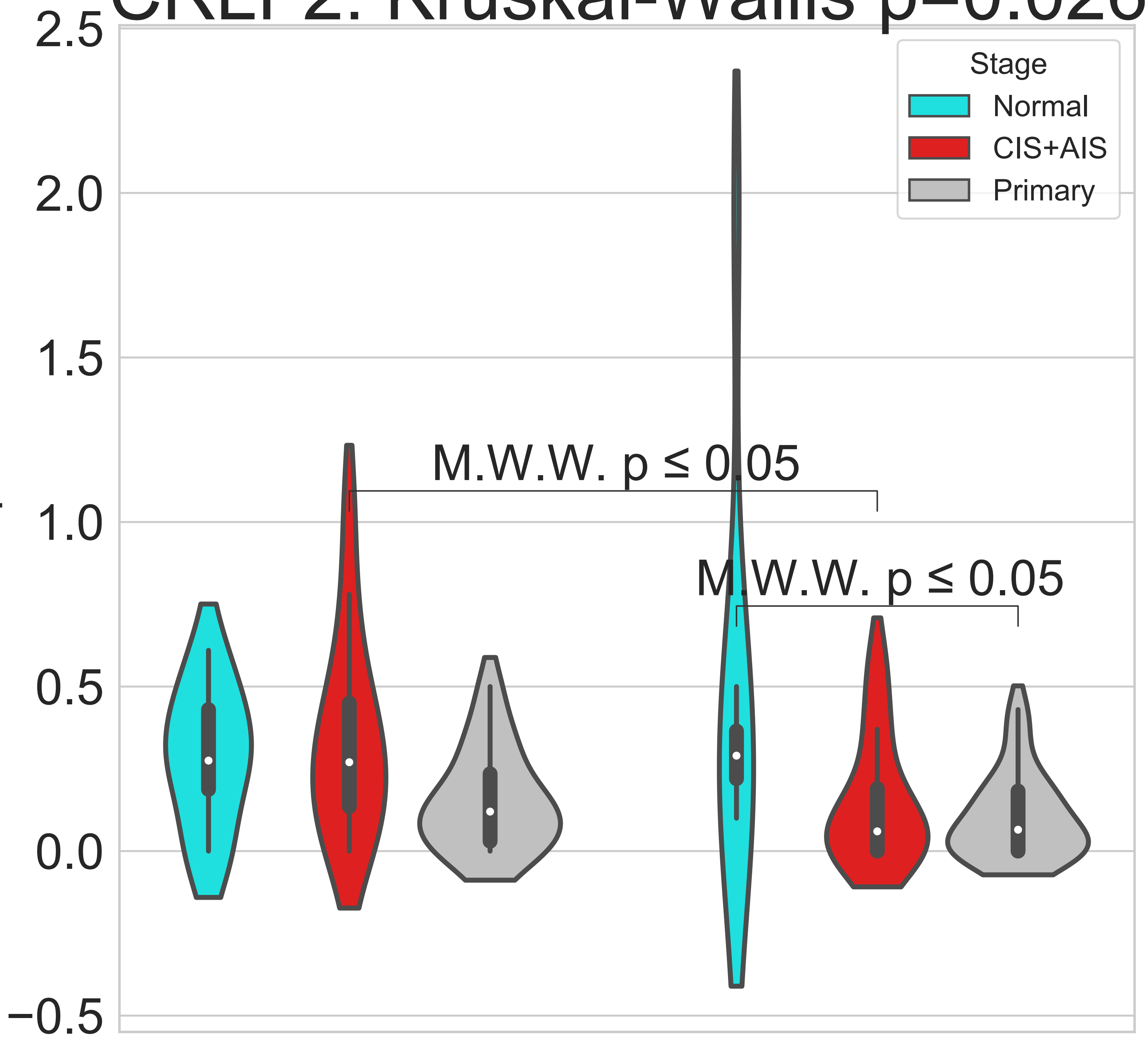


CRLF2: Kruskal-Wallis p=0.026

CRLF2 expression



Lower Higher
Mutation Shared Count (SYN)