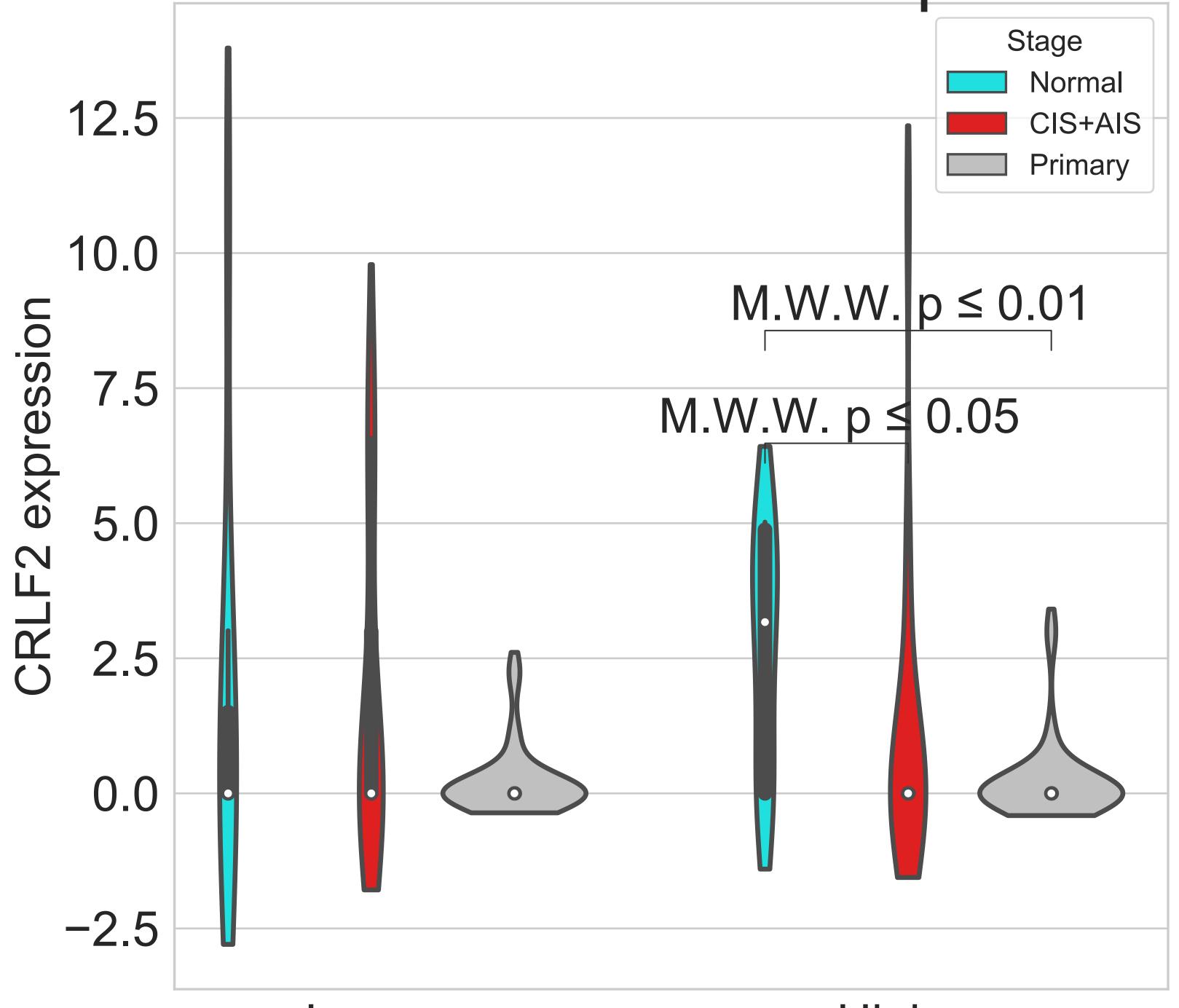
CRLF2: Kruskal-Wallis p=0.023



Lower Higher Mutation Shared Proportion