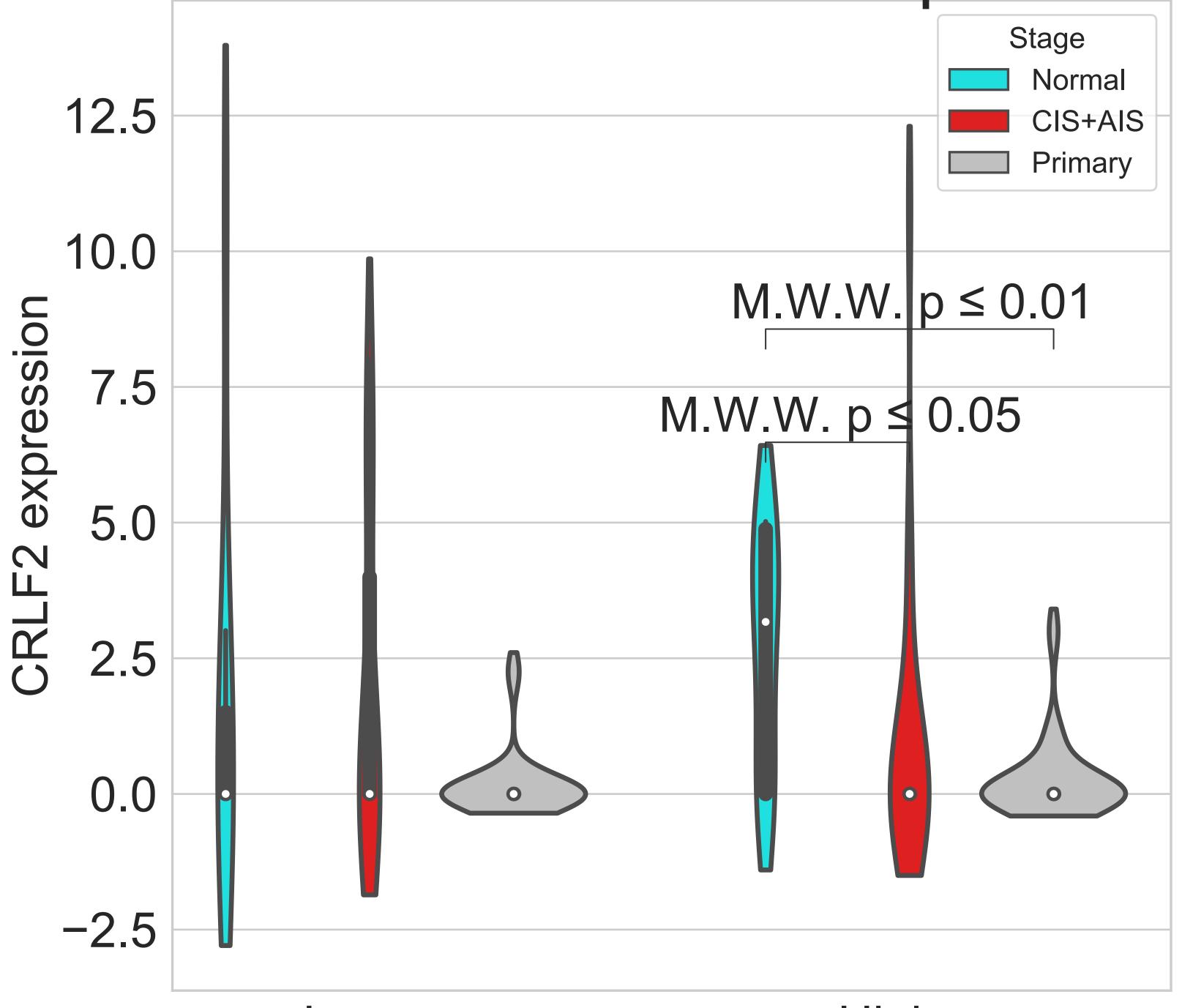
CRLF2: Kruskal-Wallis p=0.018



Lower
Mutation Shared Proportion (SYN)