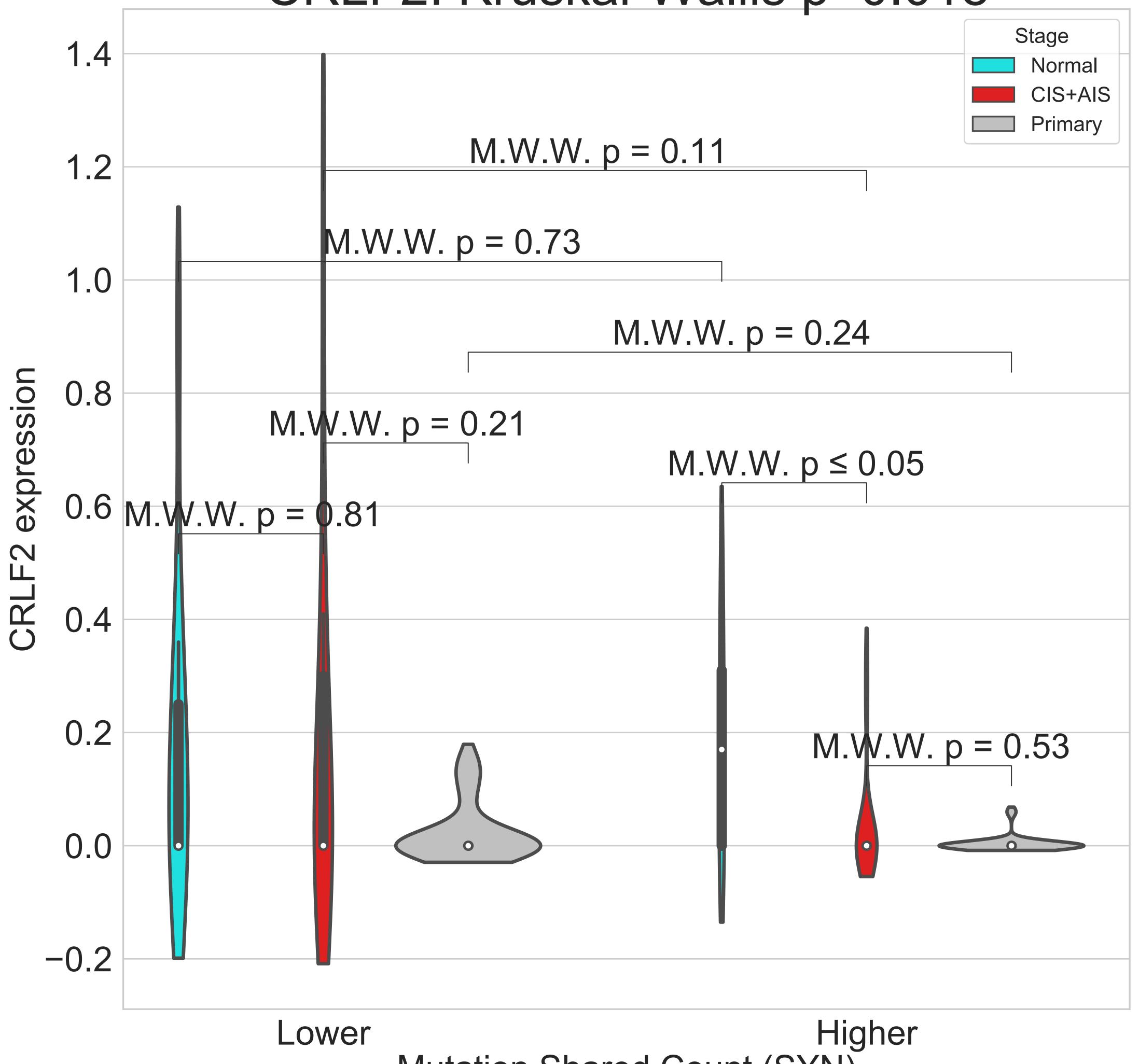
CRLF2: Kruskal-Wallis p=0.018



Mutation Shared Count (SYN)