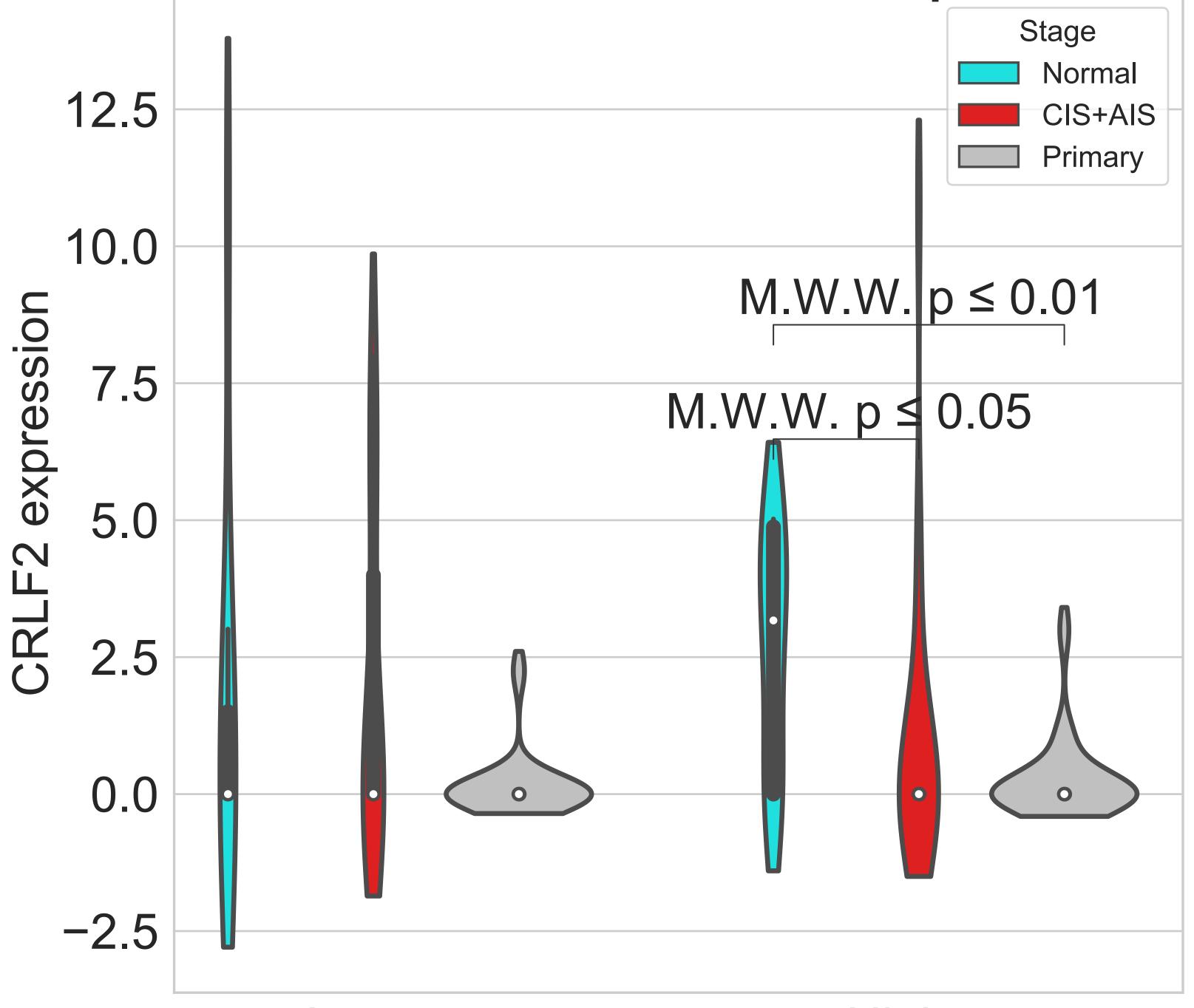
CRLF2: Kruskal-Wallis p=0.018

Stage
Normal



Lower Higher Mutation Shared Count (SYN) per TMB