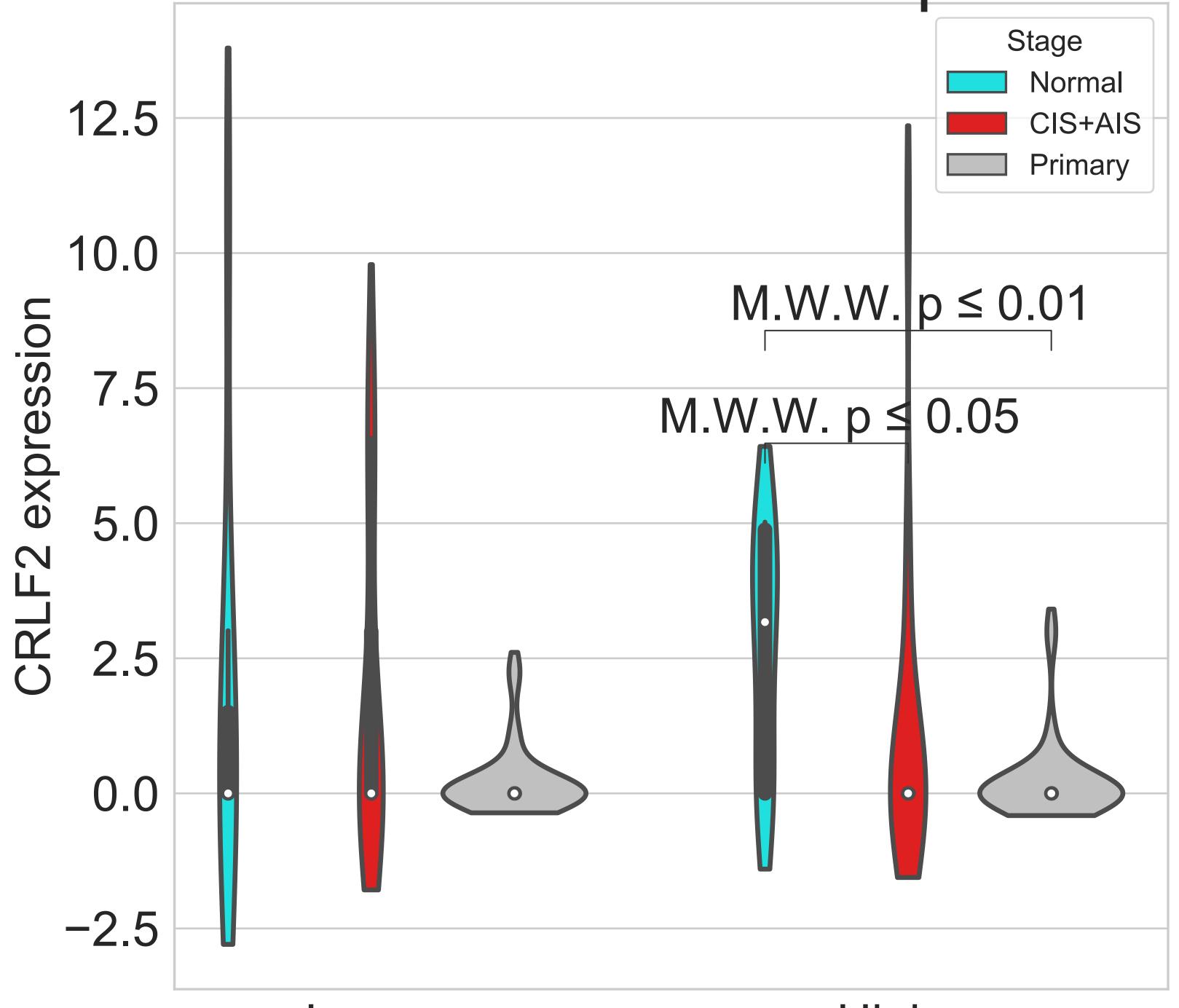
CRLF2: Kruskal-Wallis p=0.023



Lower
Mutation Shared Count per TMB