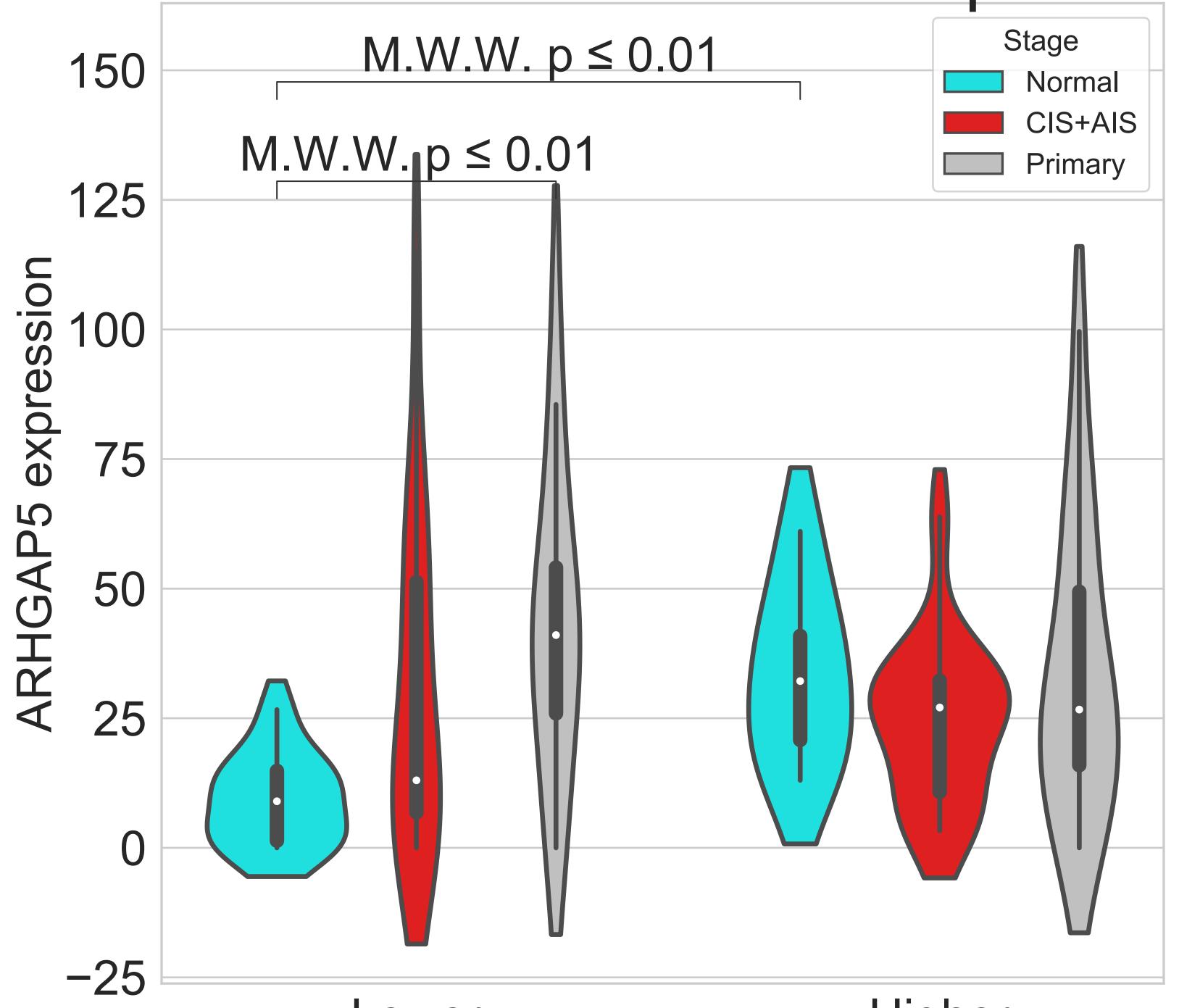
ARHGAP5: Kruskal-Wallis p=0.022



Lower
Mutation Shared Count (SYN) per TMB