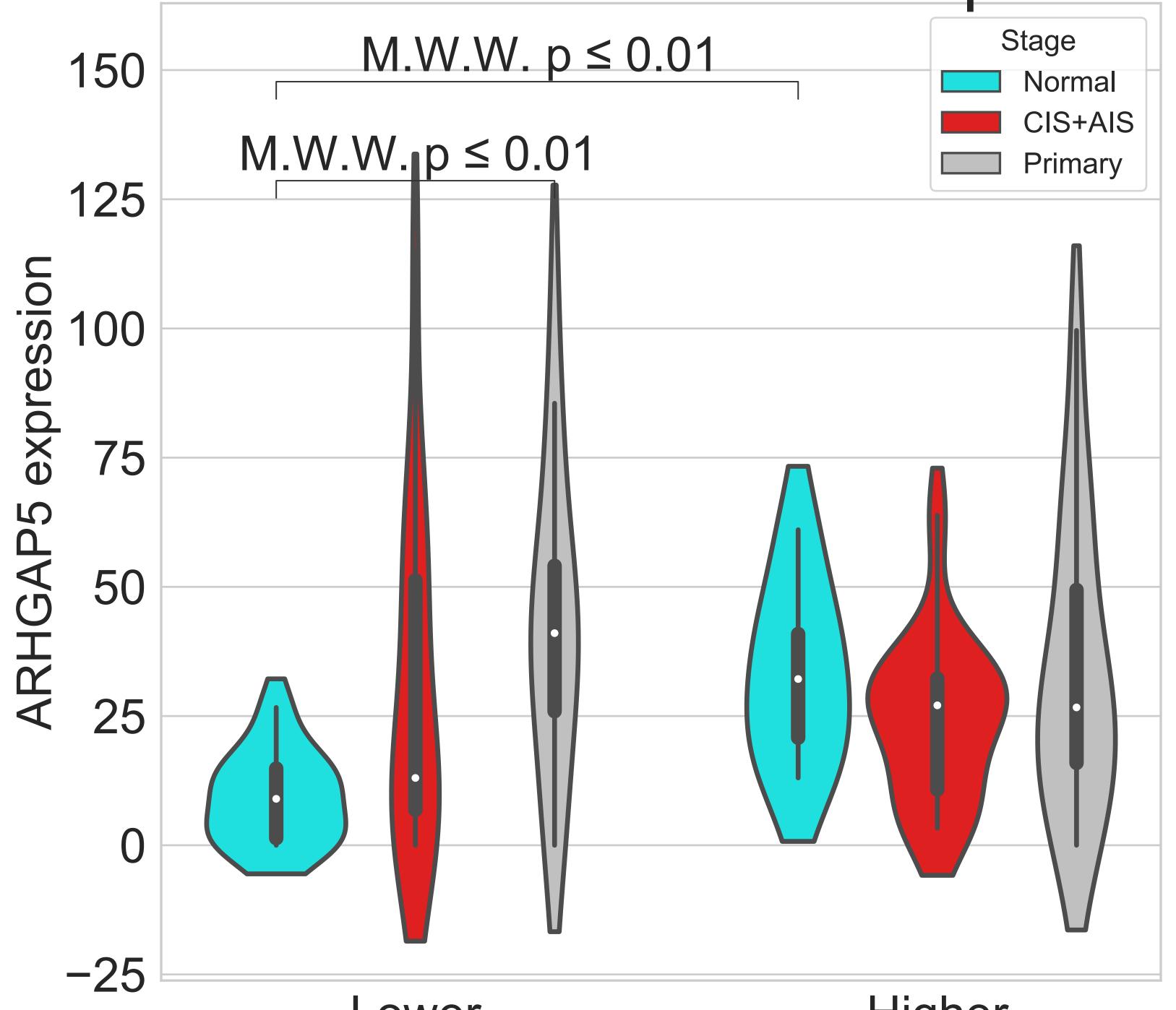
ARHGAP5: Kruskal-Wallis p=0.022



Lower
Mutation Shared Proportion (SYN)