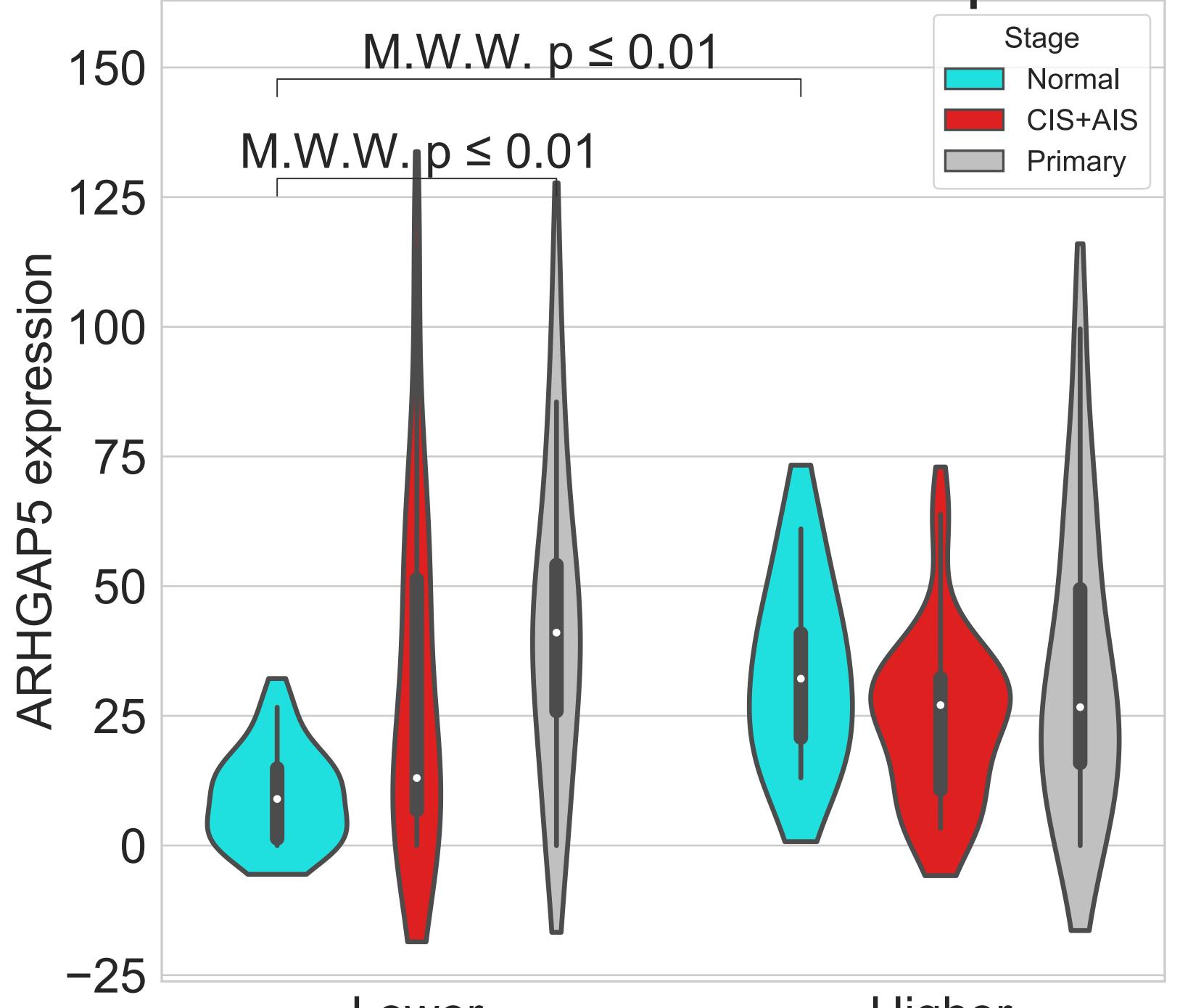
ARHGAP5: Kruskal-Wallis p=0.022



Lower Higher Mutation Shared Proportion (Union & SYN)