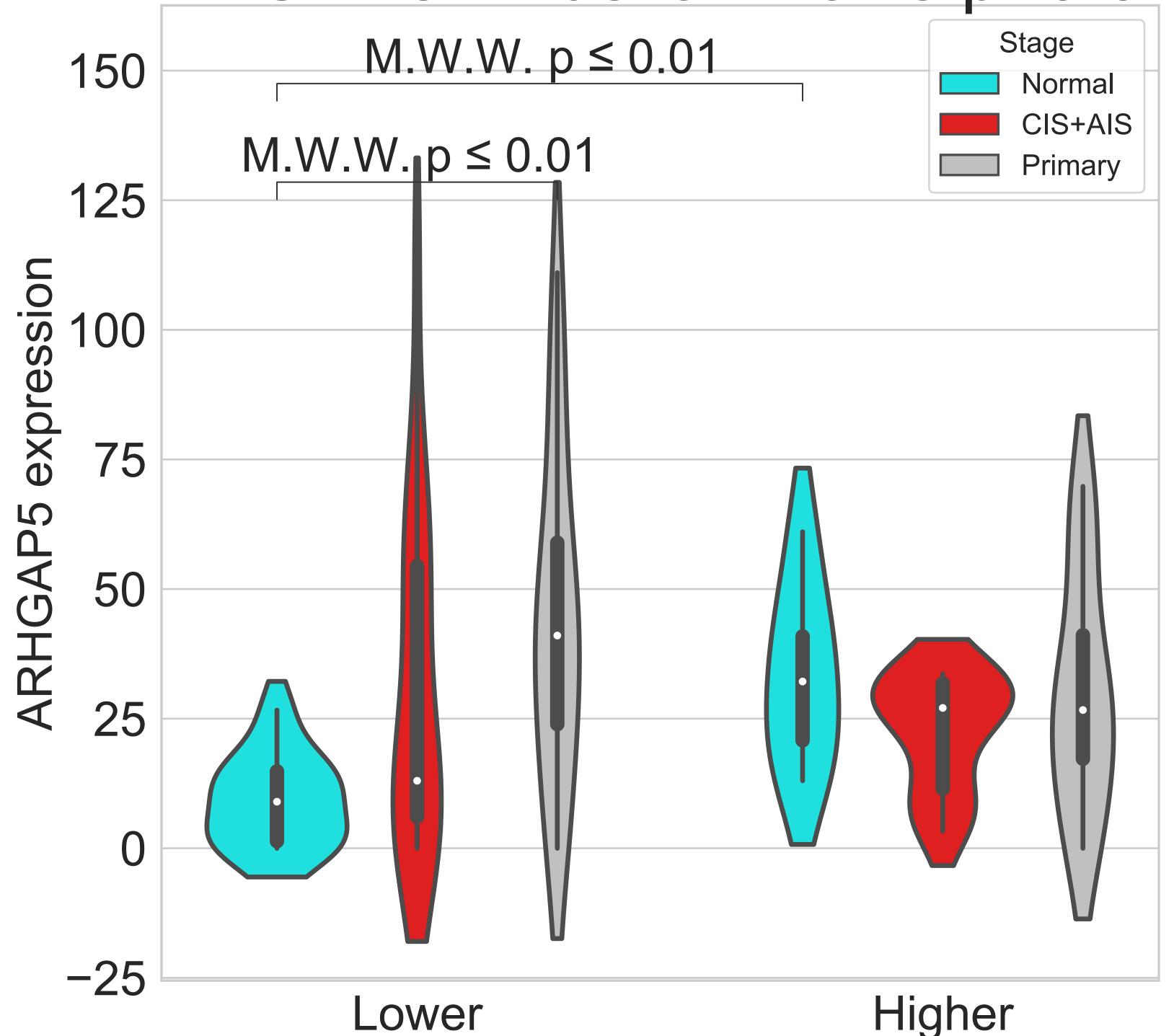
ARHGAP5: Kruskal-Wallis p=0.019



Mutation Shared Proportion (Union)