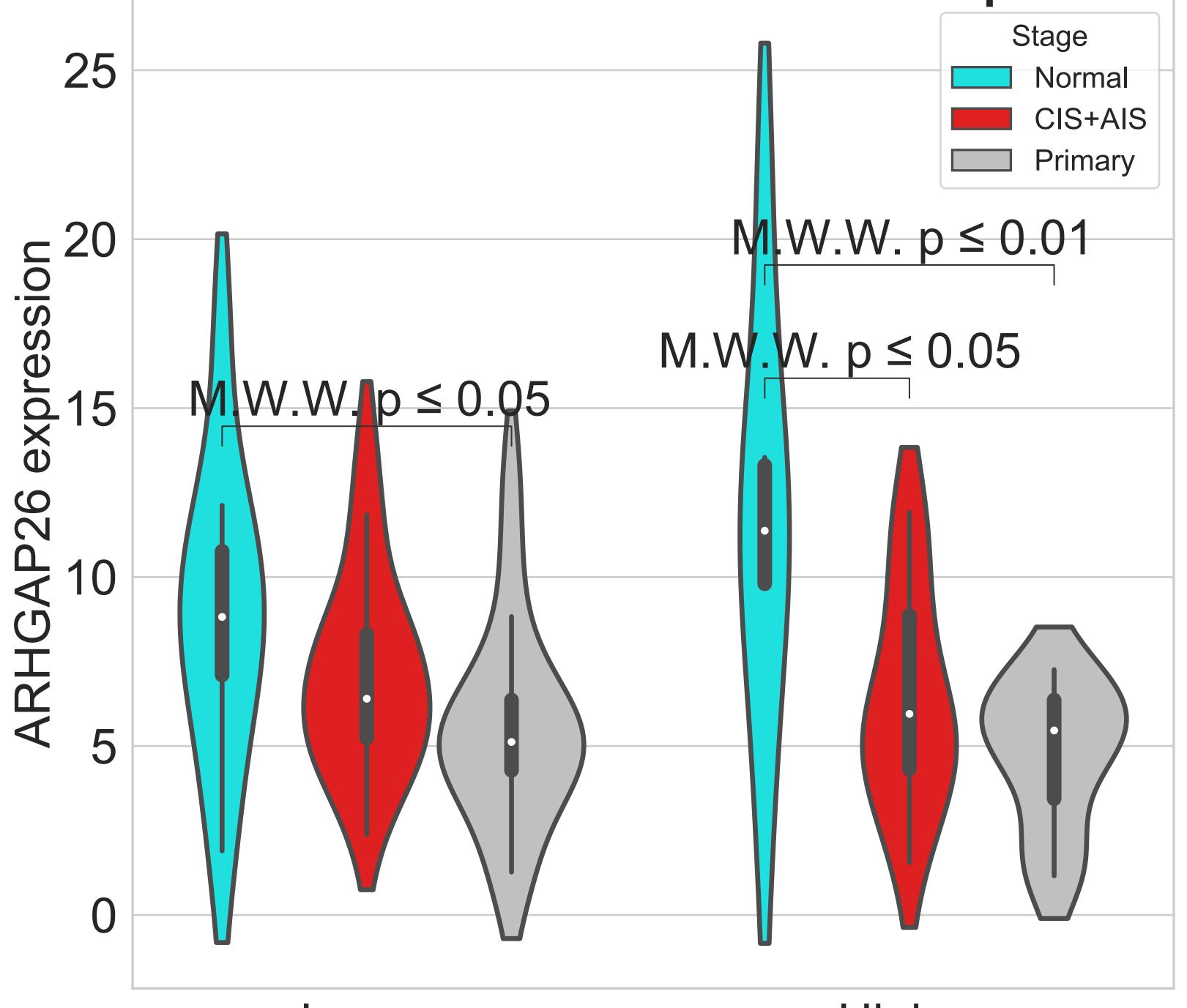
ARHGAP26: Kruskal-Wallis p=0.011



Lower Higher Mutation Shared Proportion (Union)