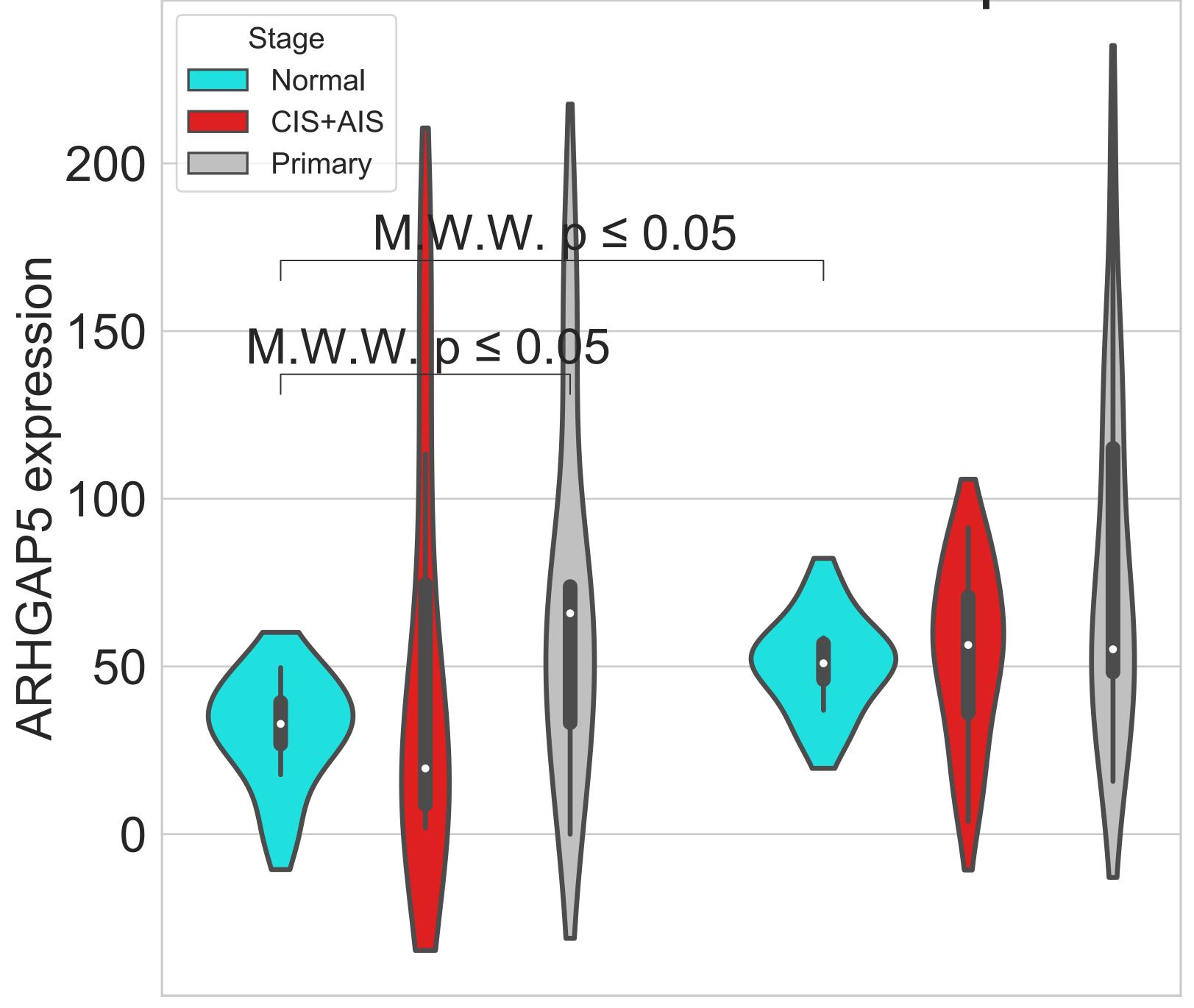
ARHGAP5: Kruskal-Wallis p=0.035



Lower
Mutation Shared Proportion (Union & SYN)