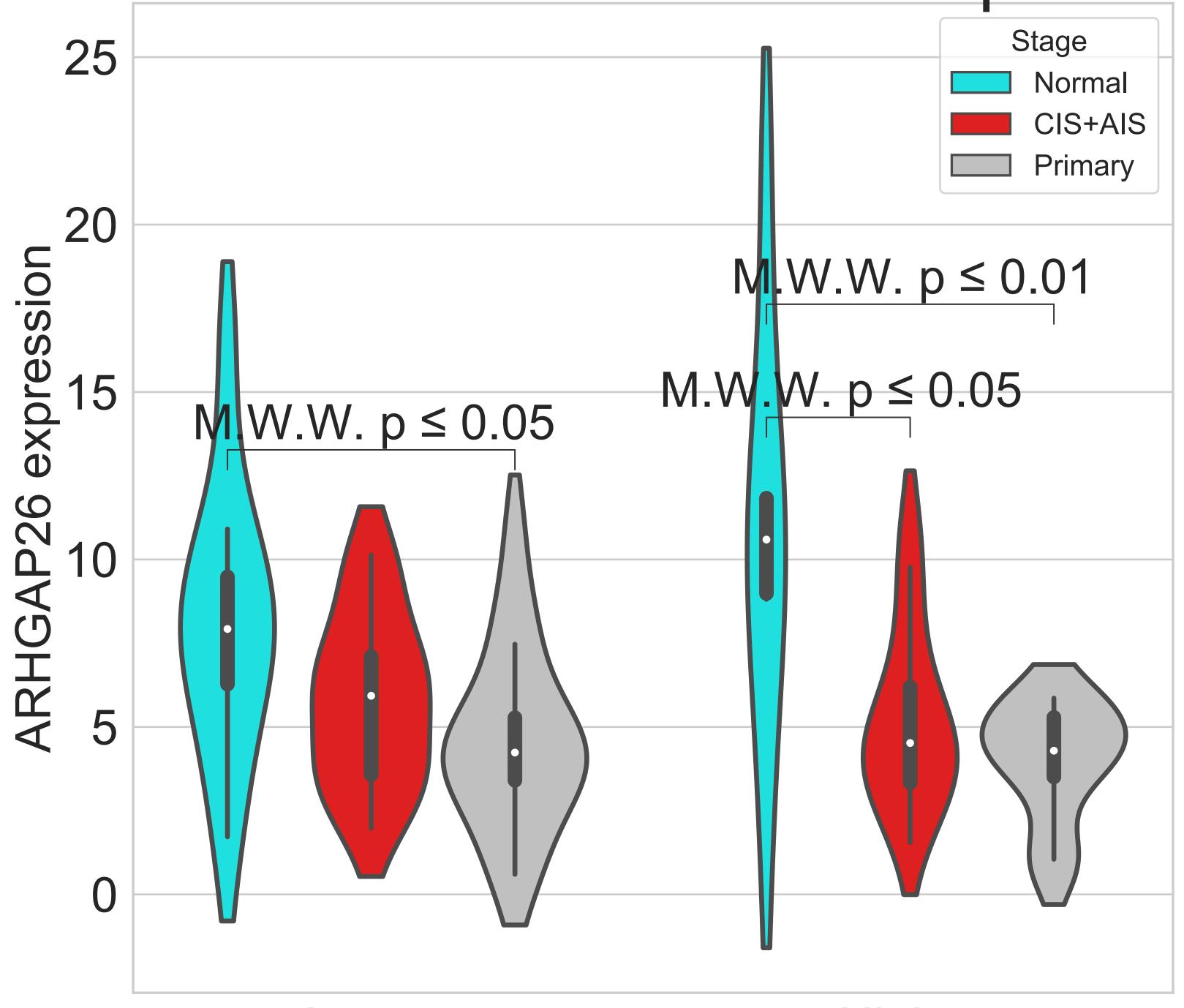
ARHGAP26: Kruskal-Wallis p=0.008



Lower

Mutation Shared Proportion (Union & SYN)