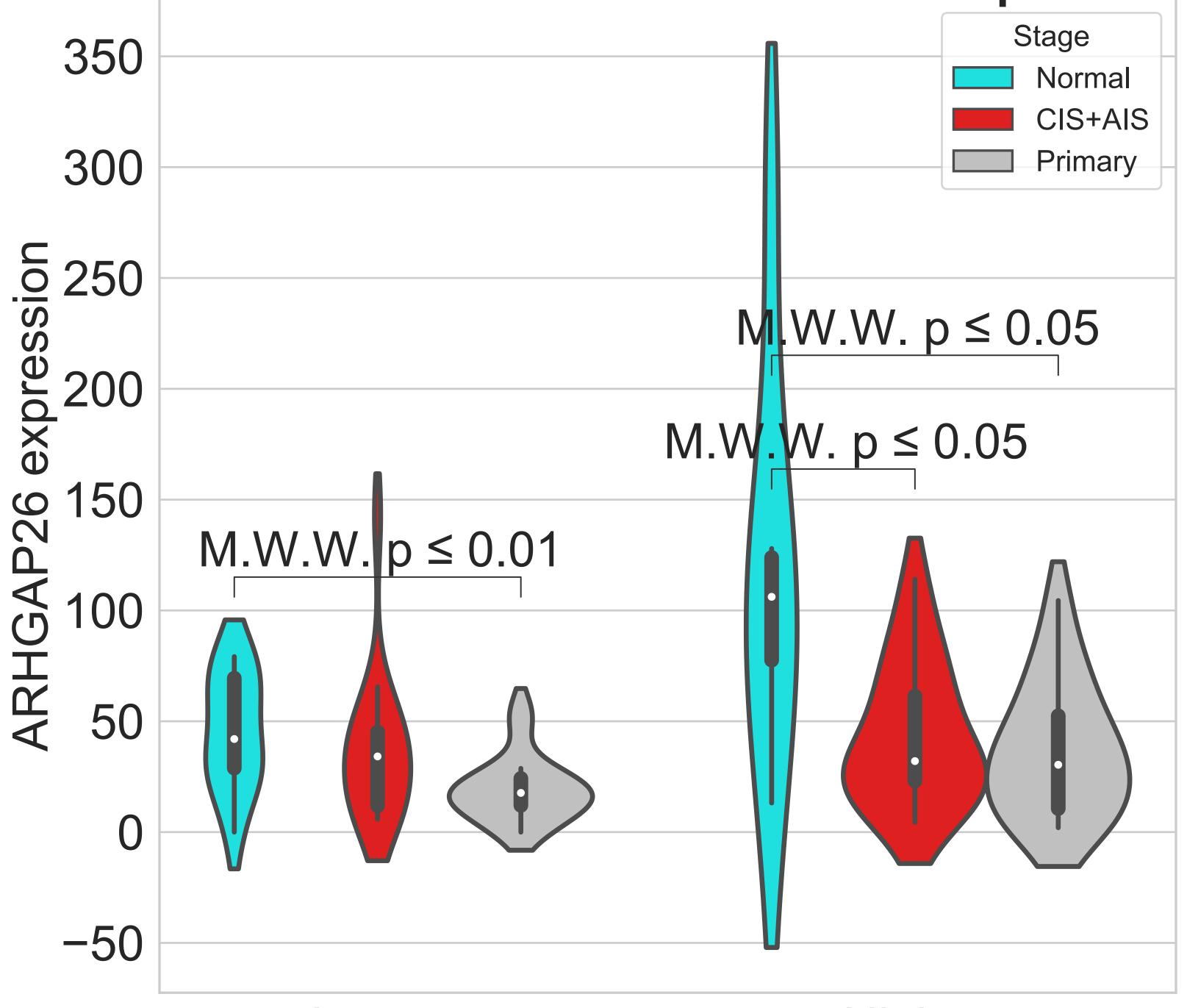
ARHGAP26: Kruskal-Wallis p=0.00



Lower Higher Mutation Shared Count (SYN) per TMB