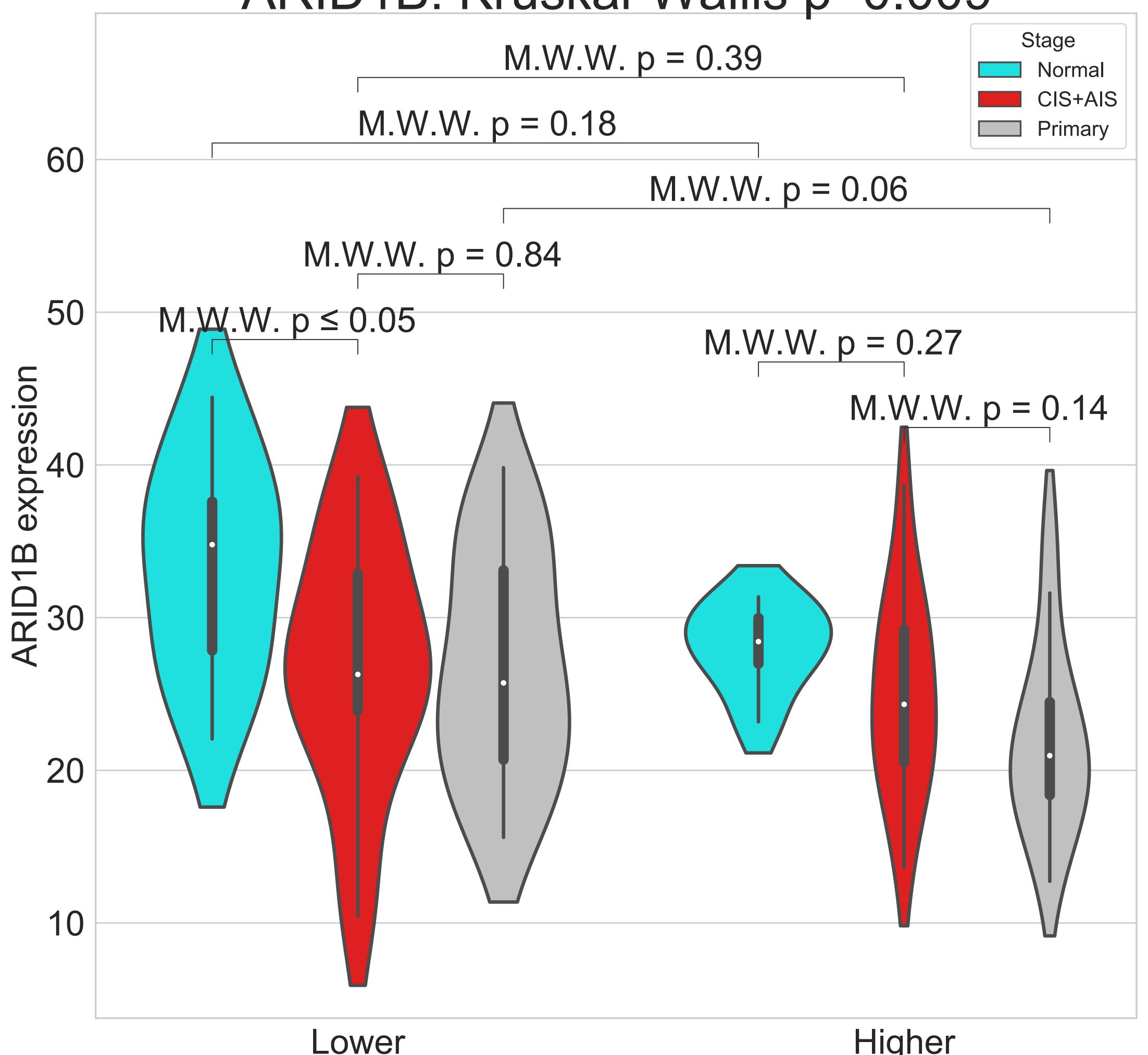
ARID1B: Kruskal-Wallis p=0.009



Higher Mutation Shared Proportion (SYN)