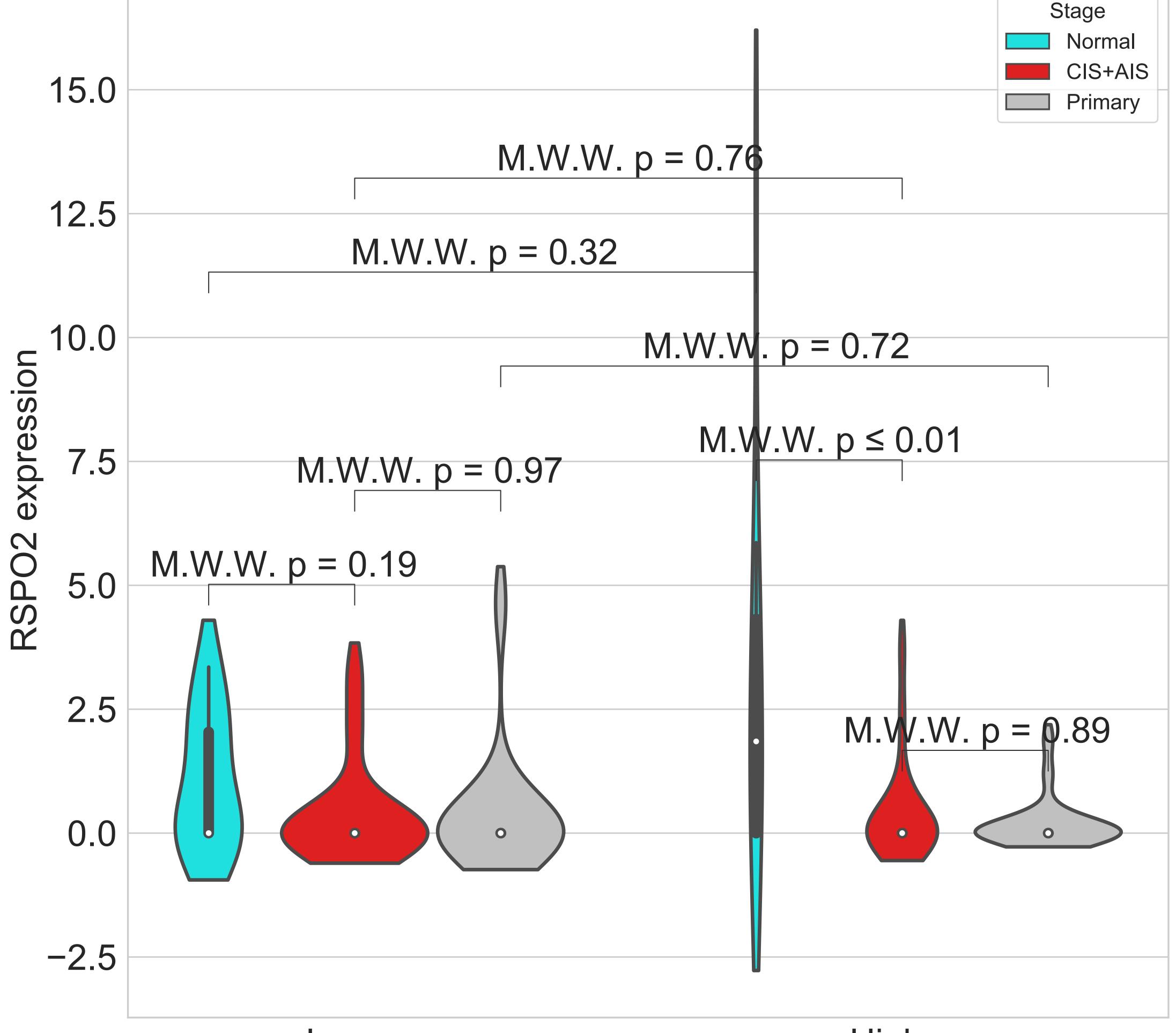
RSP02: Kruskal-Wallis p=0.019



Higher Lower Mutation Shared Proportion (SYN)