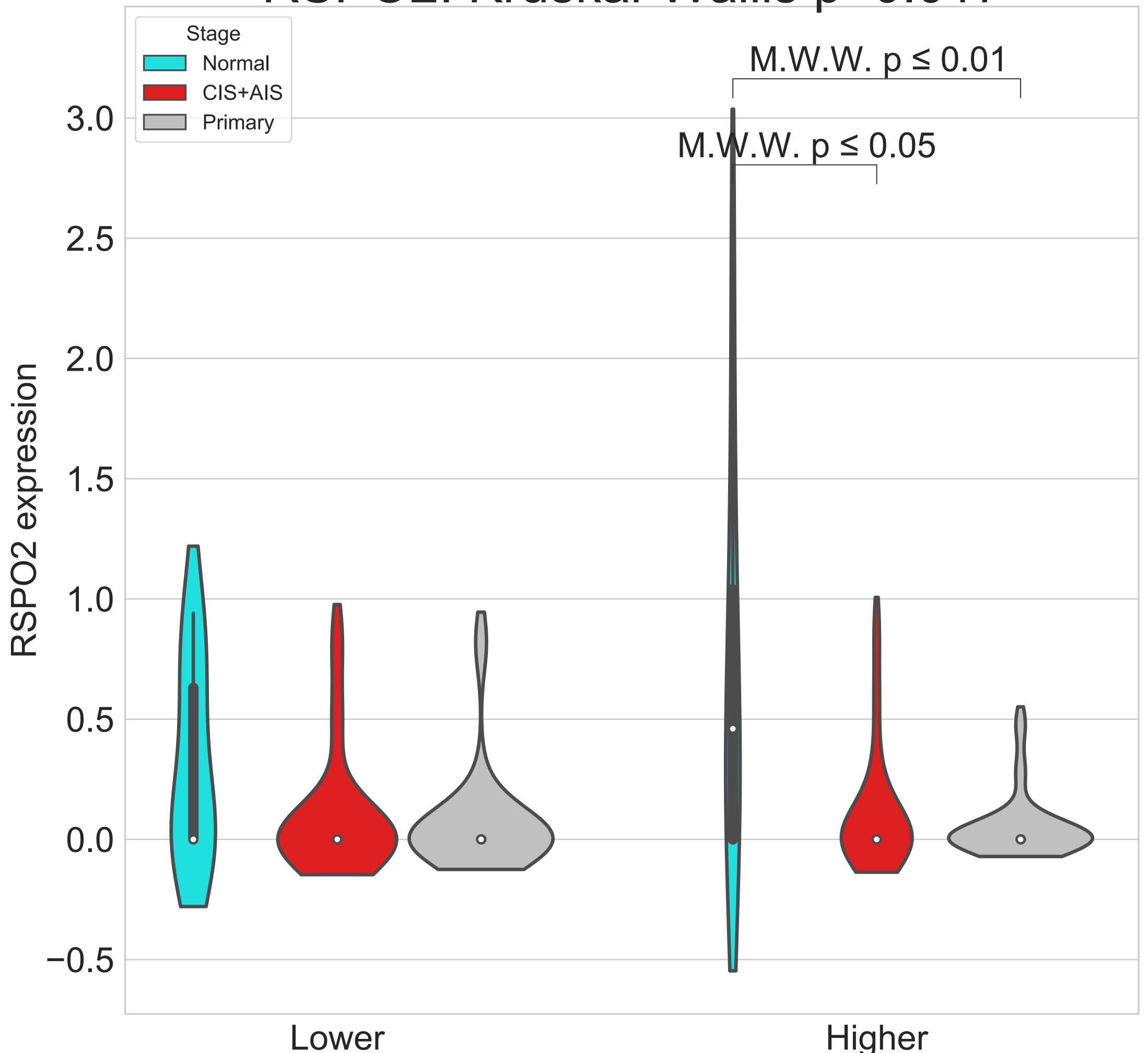
RSP02: Kruskal-Wallis p=0.017



Higher Mutation Shared Proportion