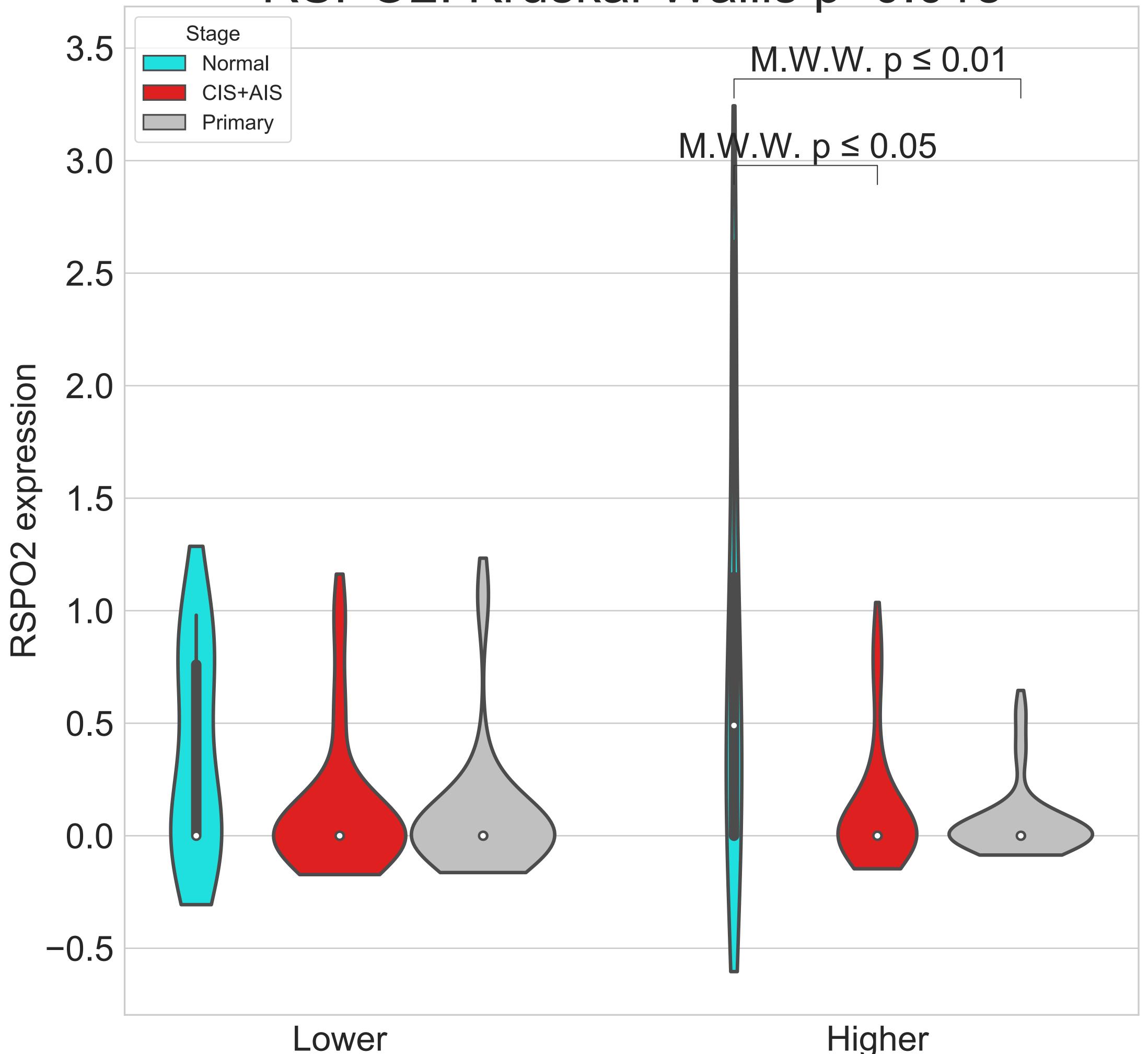
RSP02: Kruskal-Wallis p=0.018



Higher Mutation Shared Proportion