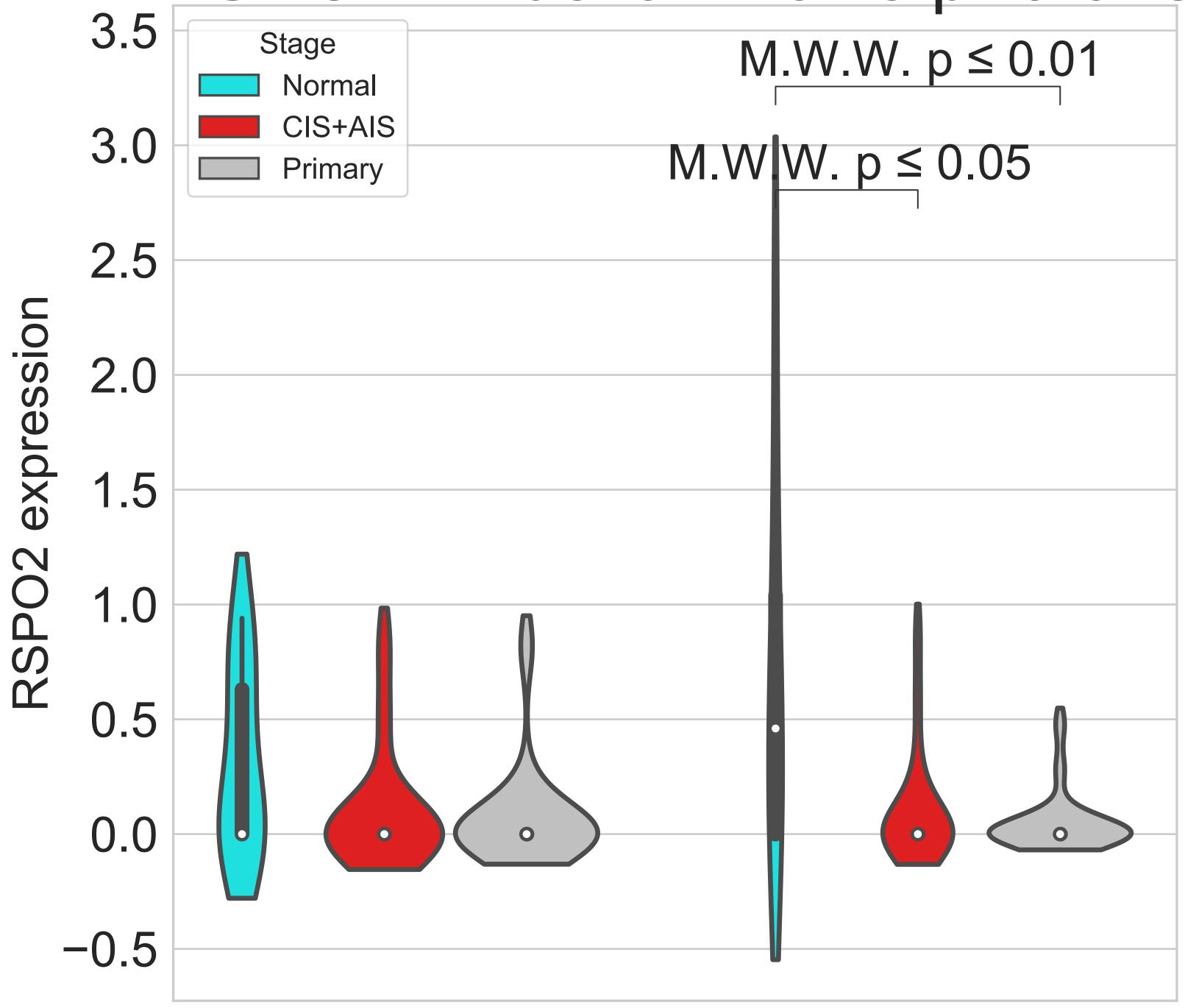
RSP02: Kruskal-Wallis p=0.018



Lower Higher Mutation Shared Proportion (SYN)