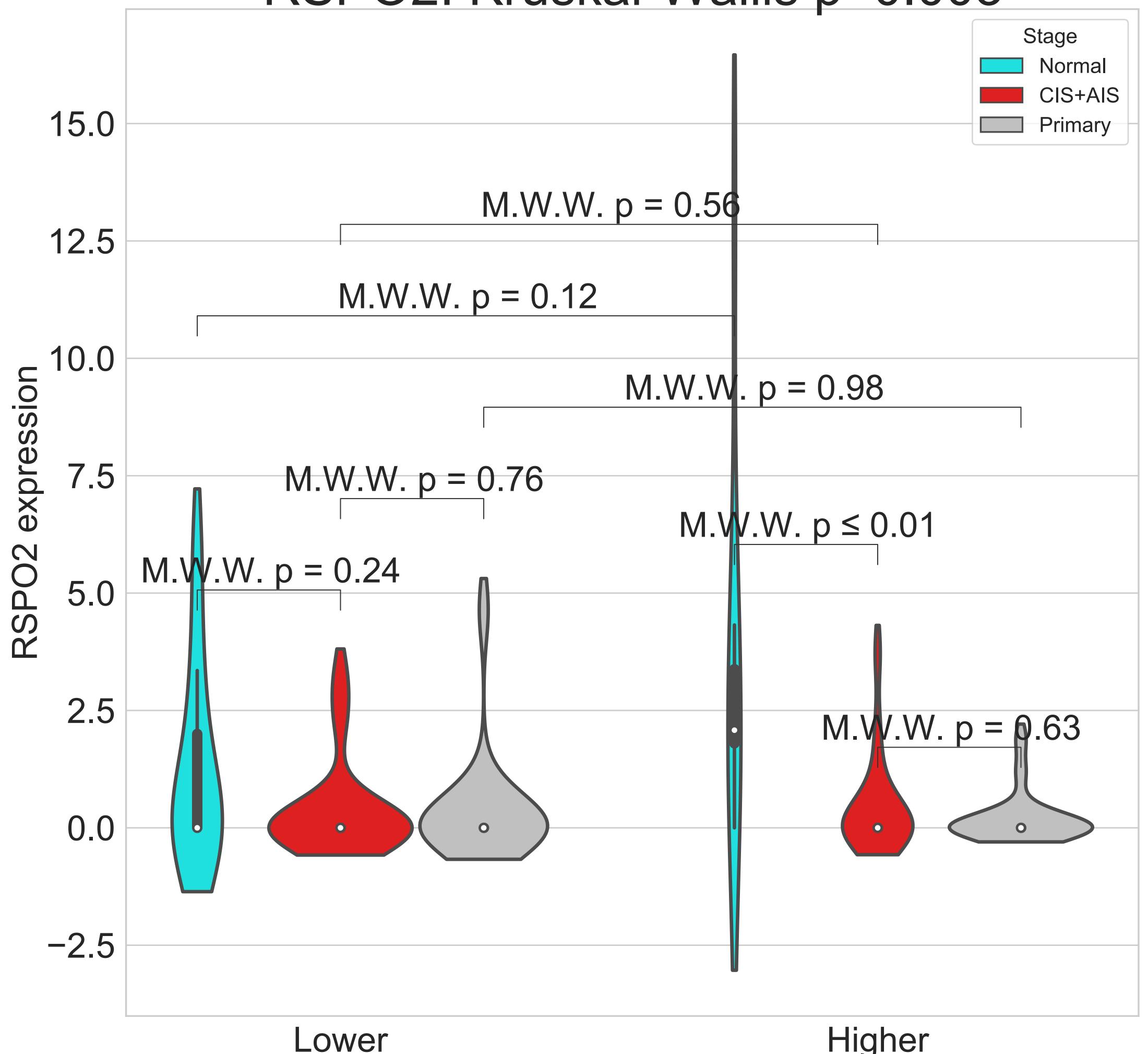
RSP02: Kruskal-Wallis p=0.003



wer
Mutation Shared Count (SYN)