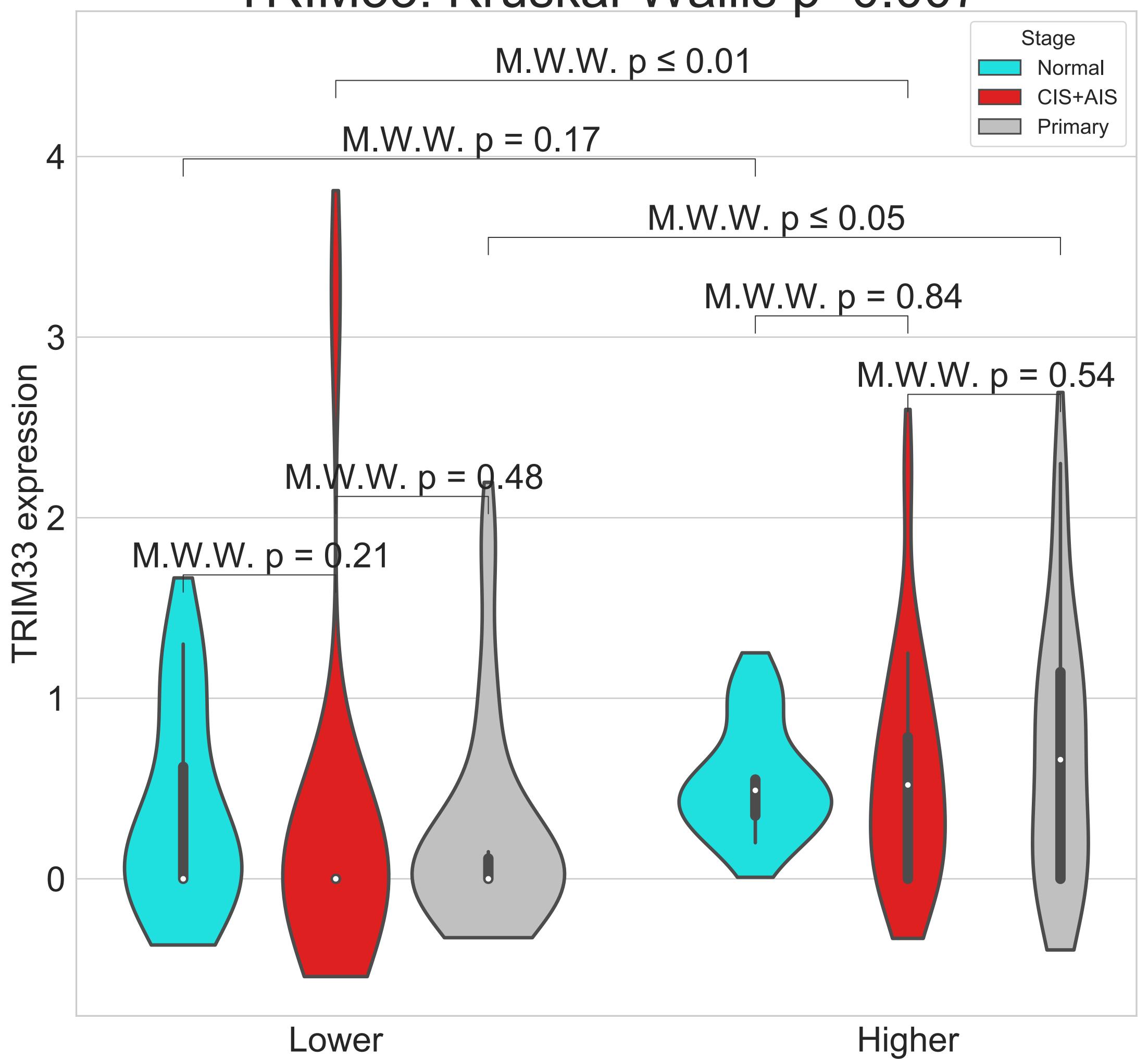
TRIM33: Kruskal-Wallis p=0.007



wer
Mutation Shared Proportion (SYN)