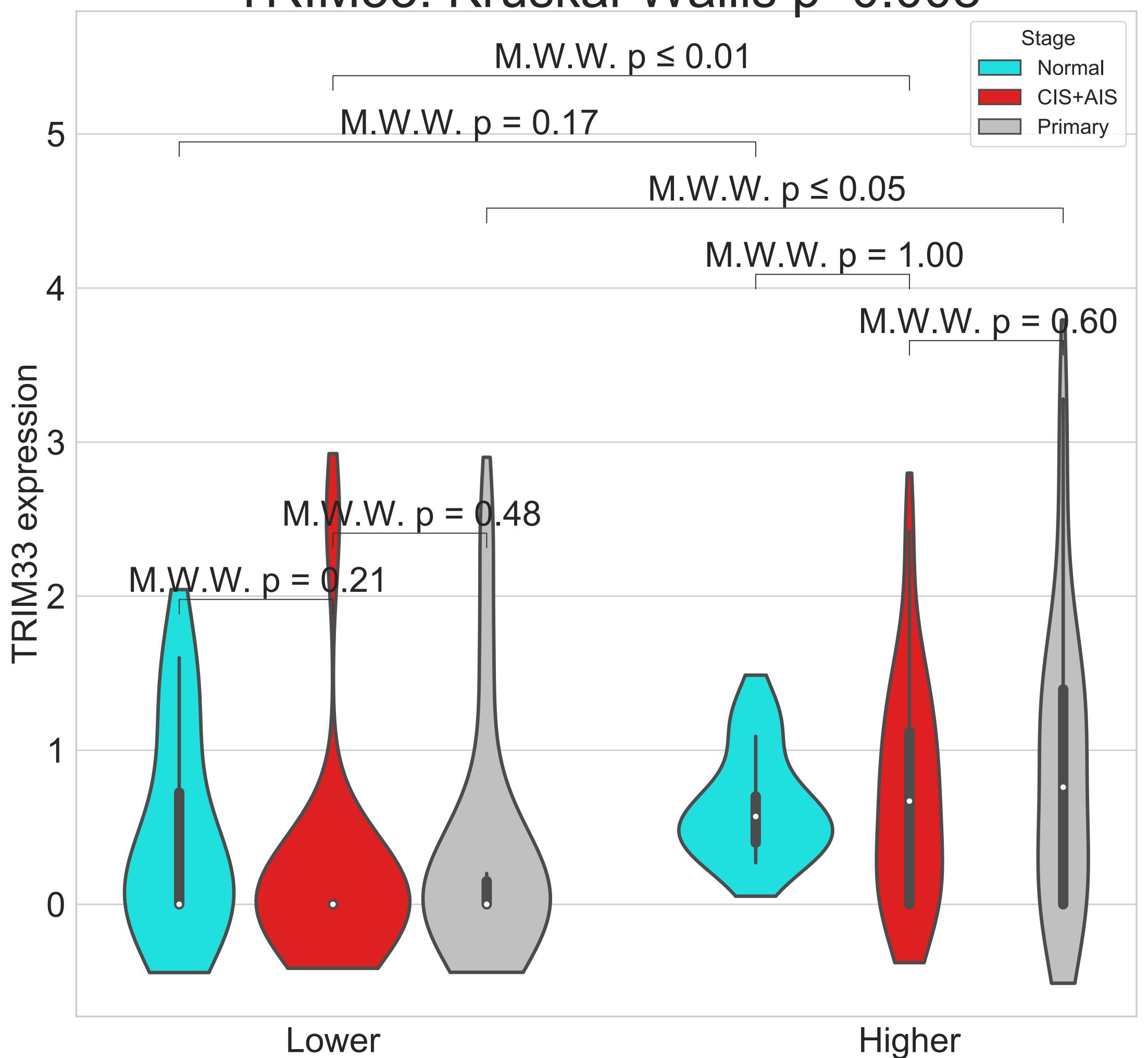
TRIM33: Kruskal-Wallis p=0.008



Mutation Shared Proportion (SYN)