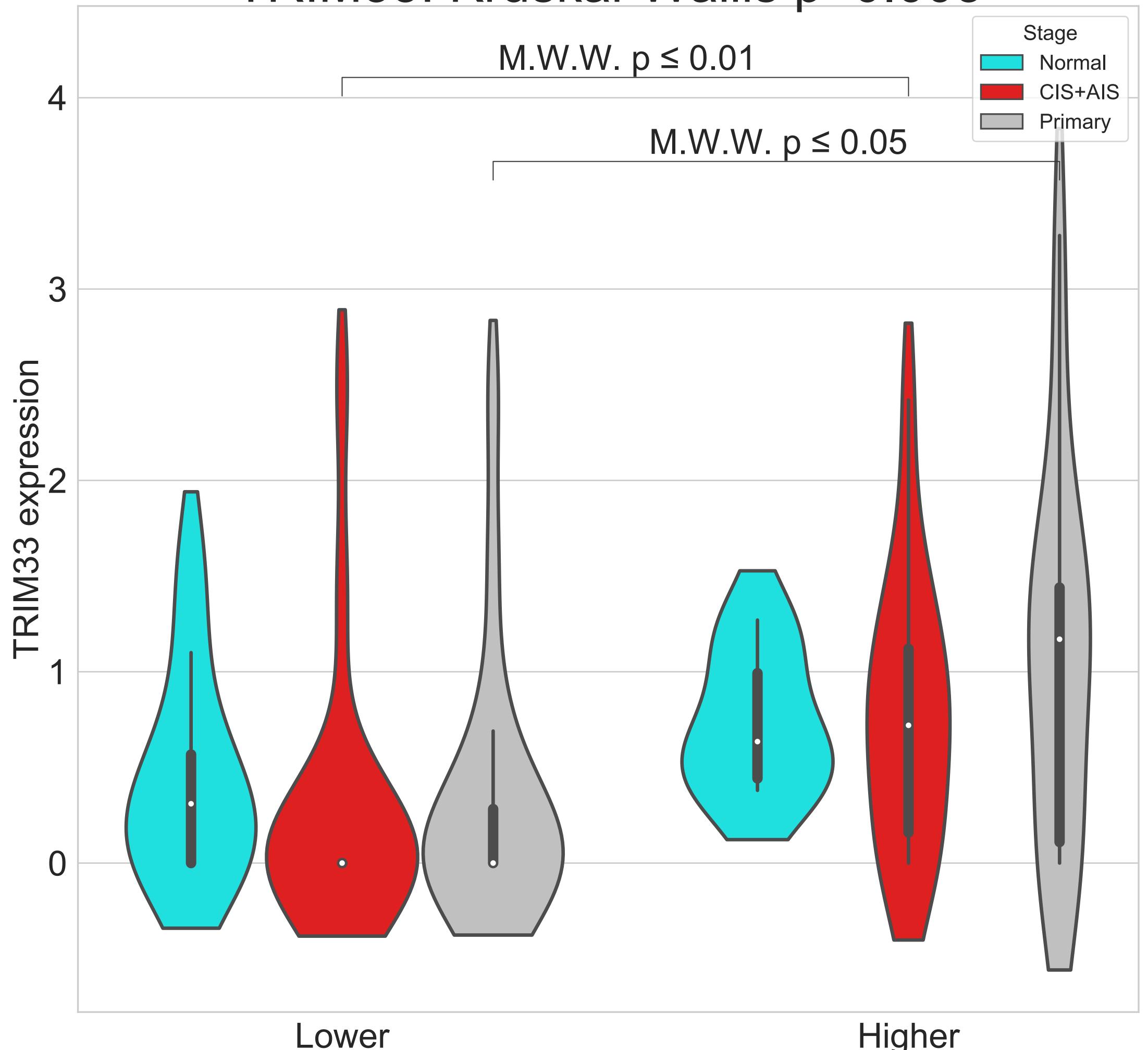
TRIM33: Kruskal-Wallis p=0.003



Wer Higher Mutation Shared Proportion (SYN)