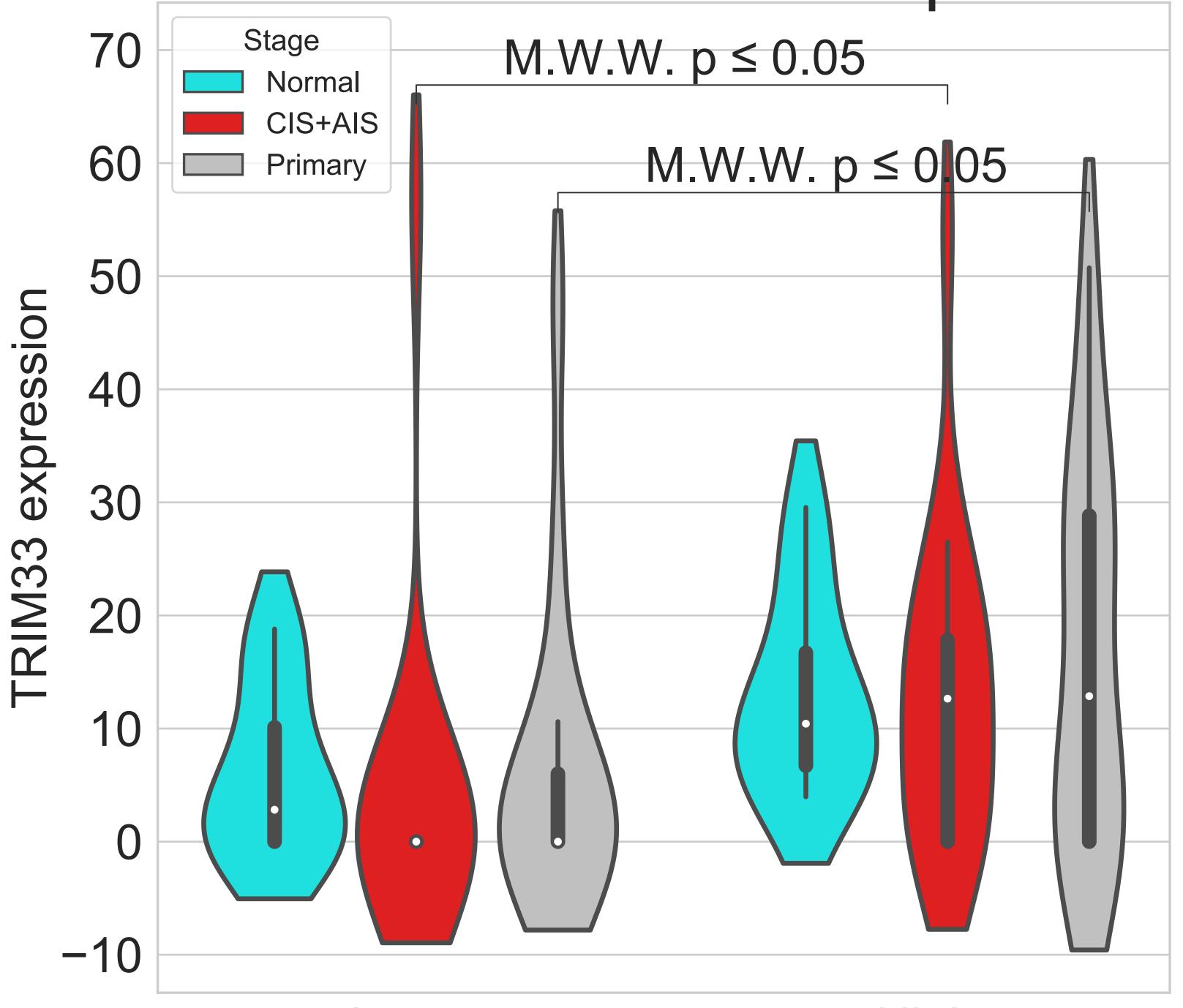
TRIM33: Kruskal-Wallis p=0.008



Lower Higher Mutation Shared Proportion (Union & SYN)