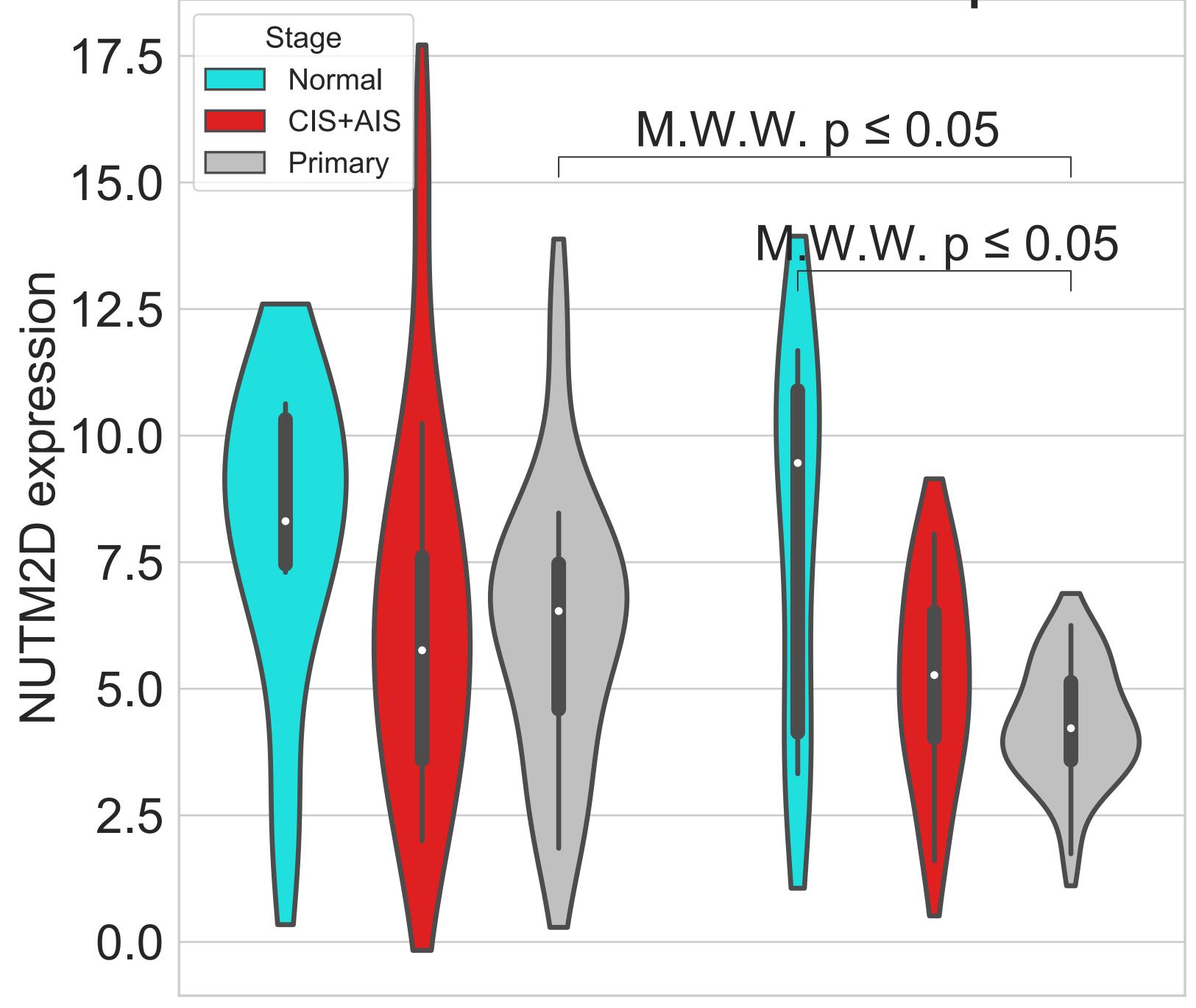
NUTM2D: Kruskal-Wallis p=0.007



Lower Higher Mutation Shared Proportion (SYN)