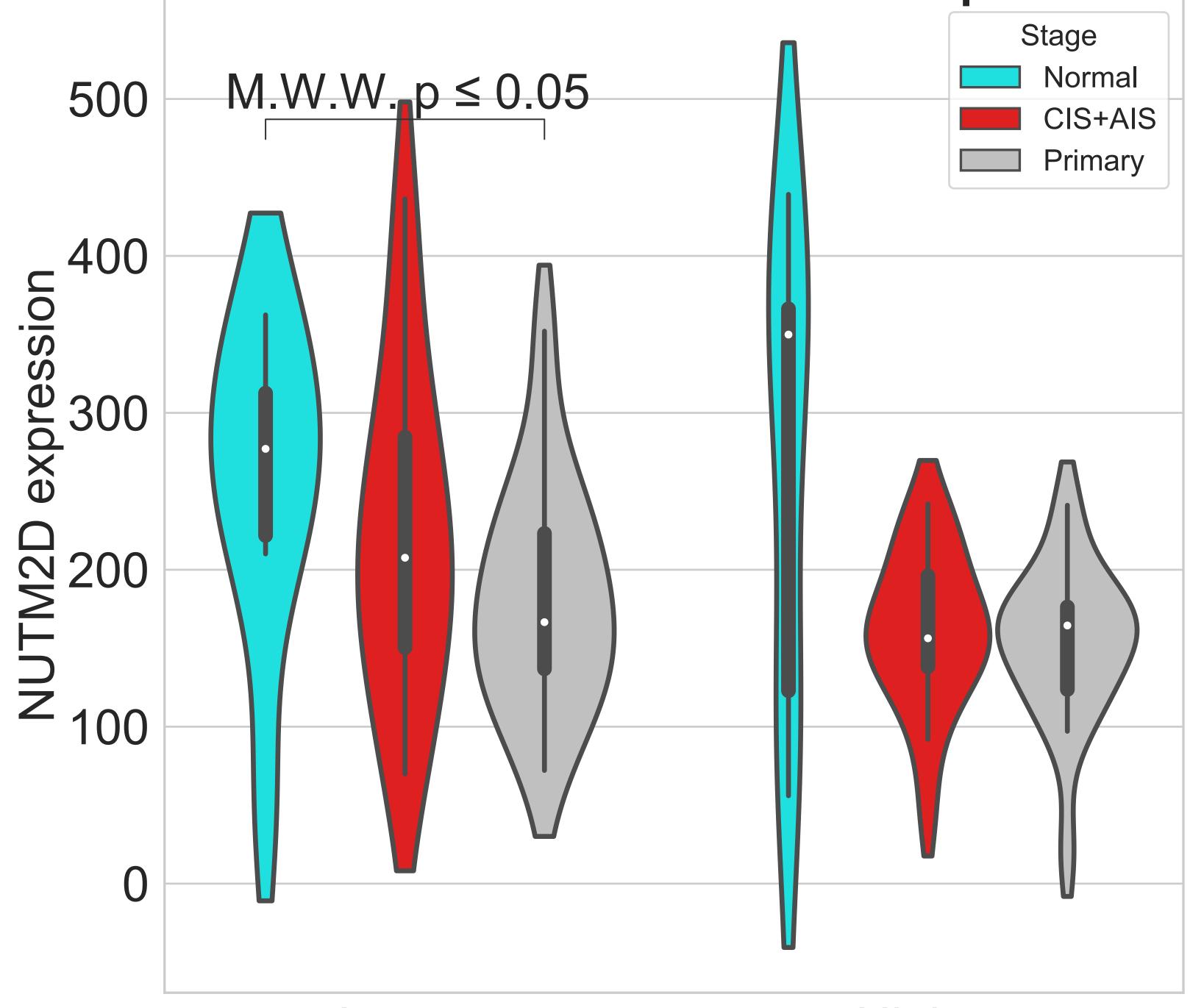
NUTM2D: Kruskal-Wallis p=0.034



Lower Higher Mutation Shared Proportion (Union & SYN)