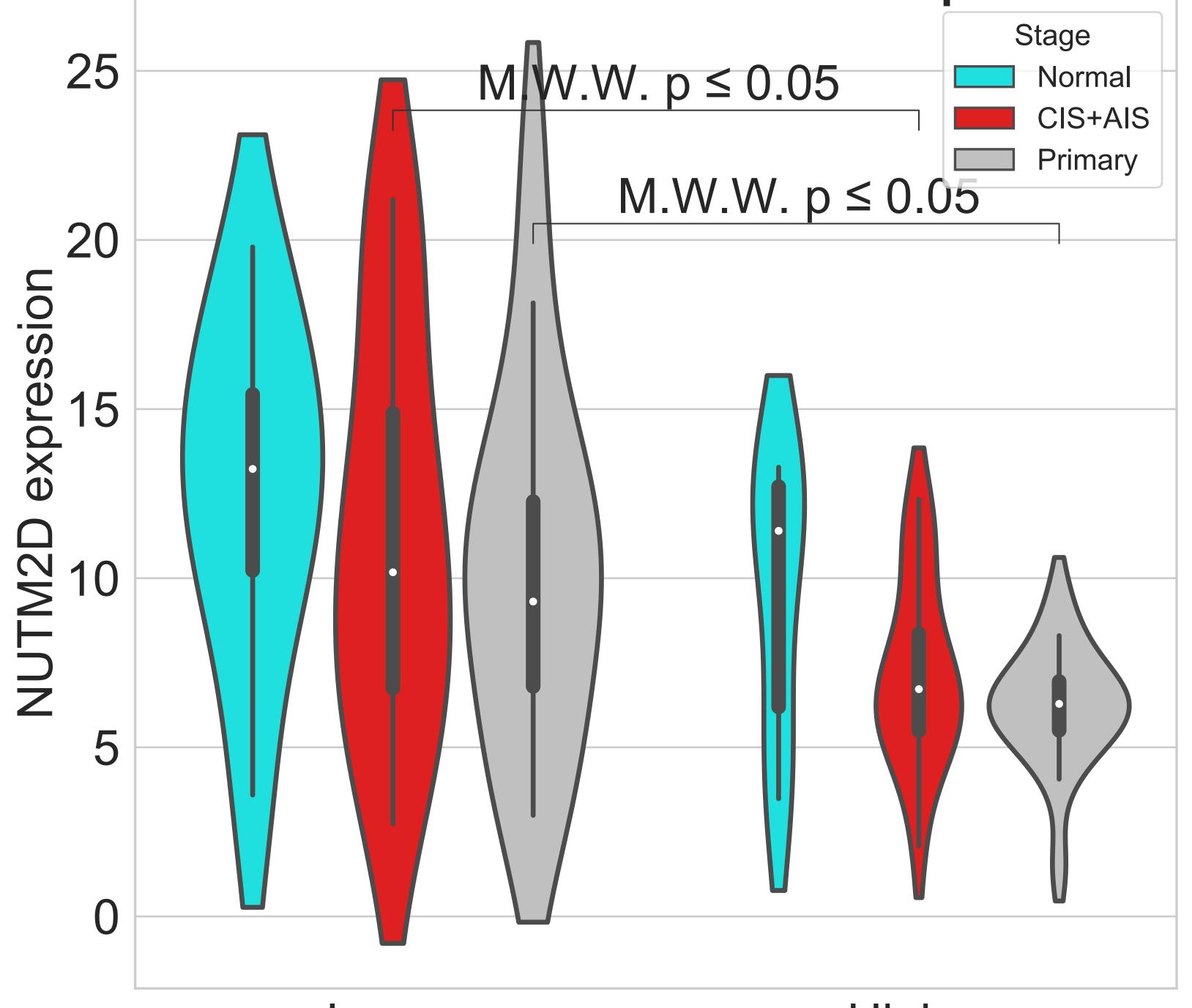
NUTM2D: Kruskal-Wallis p=0.006



Lower Higher Mutation Shared Proportion (Union & SYN)