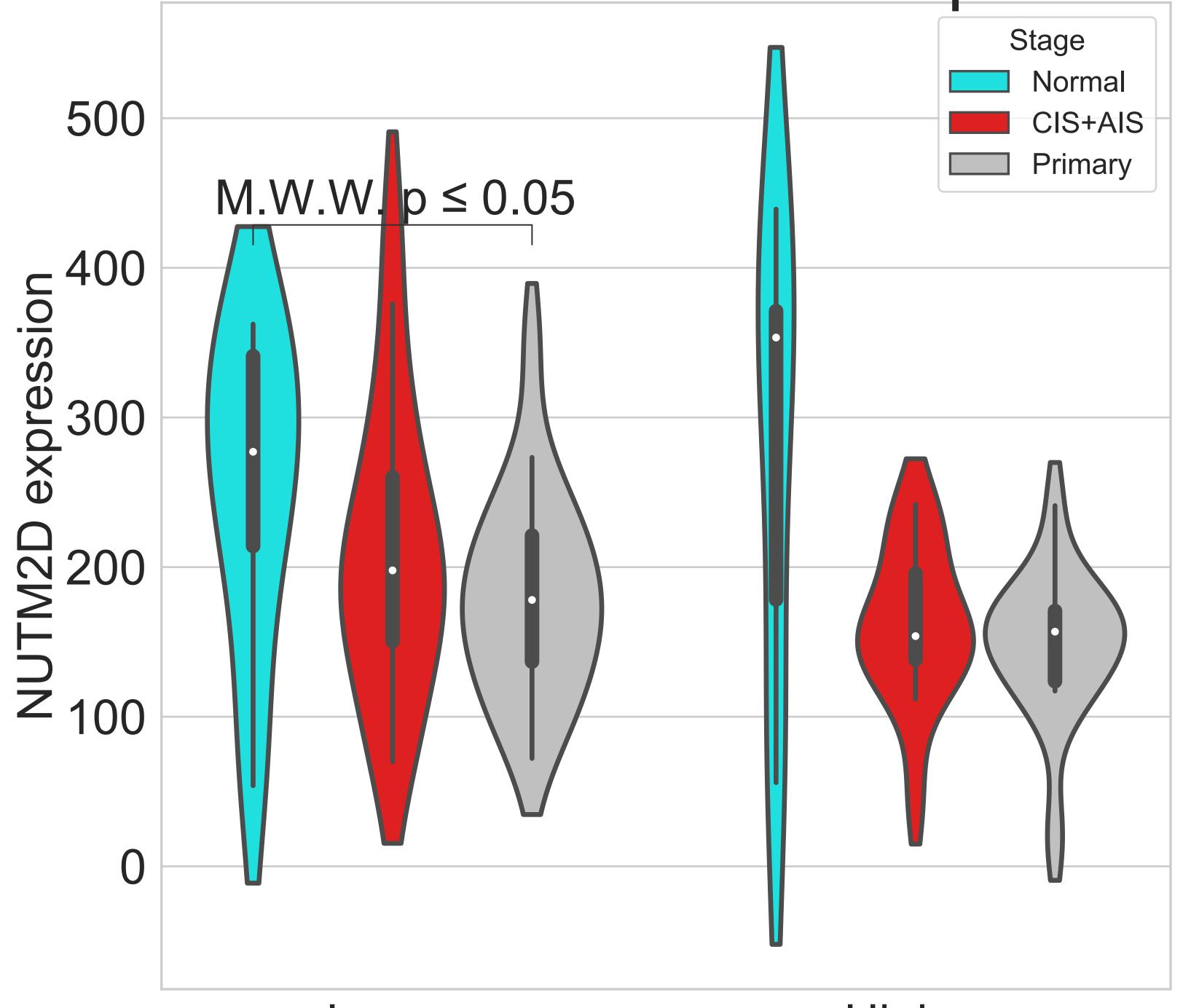
NUTM2D: Kruskal-Wallis p=0.041



Lower Higher Mutation Shared Proportion