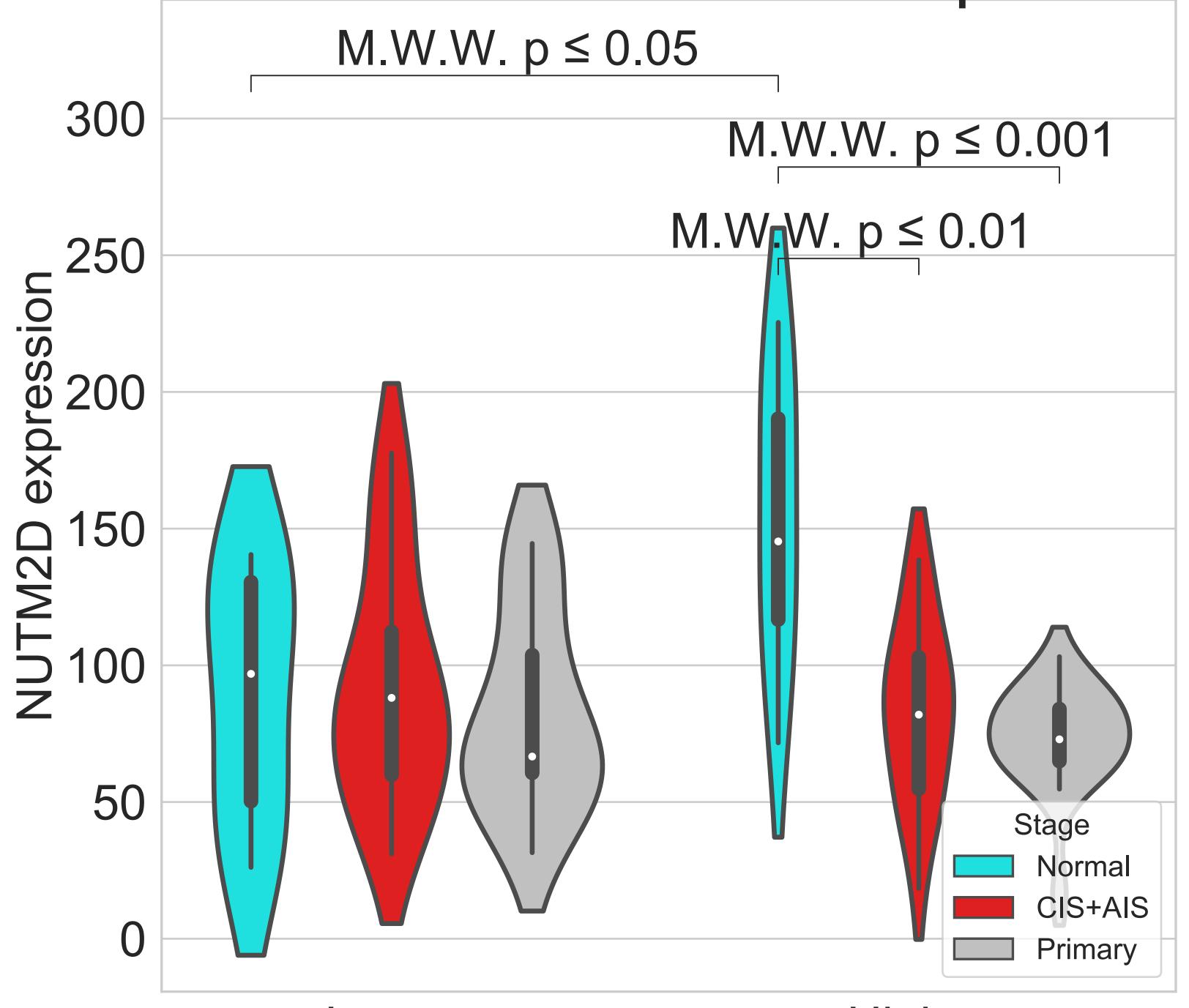
NUTM2D: Kruskal-Wallis p=0.011



Lower Higher Mutation Shared Count