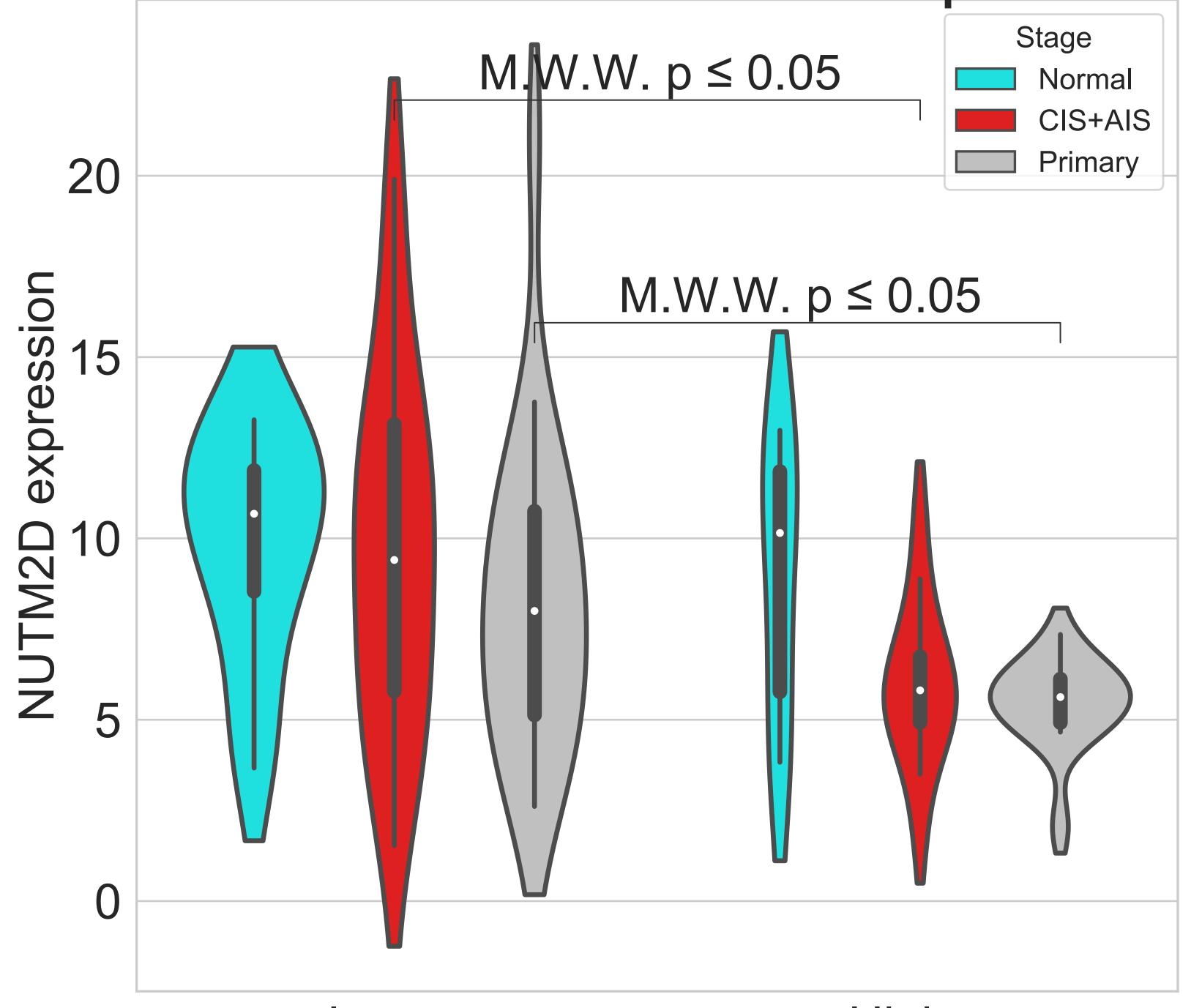
NUTM2D: Kruskal-Wallis p=0.014



Lower Higher Mutation Shared Proportion