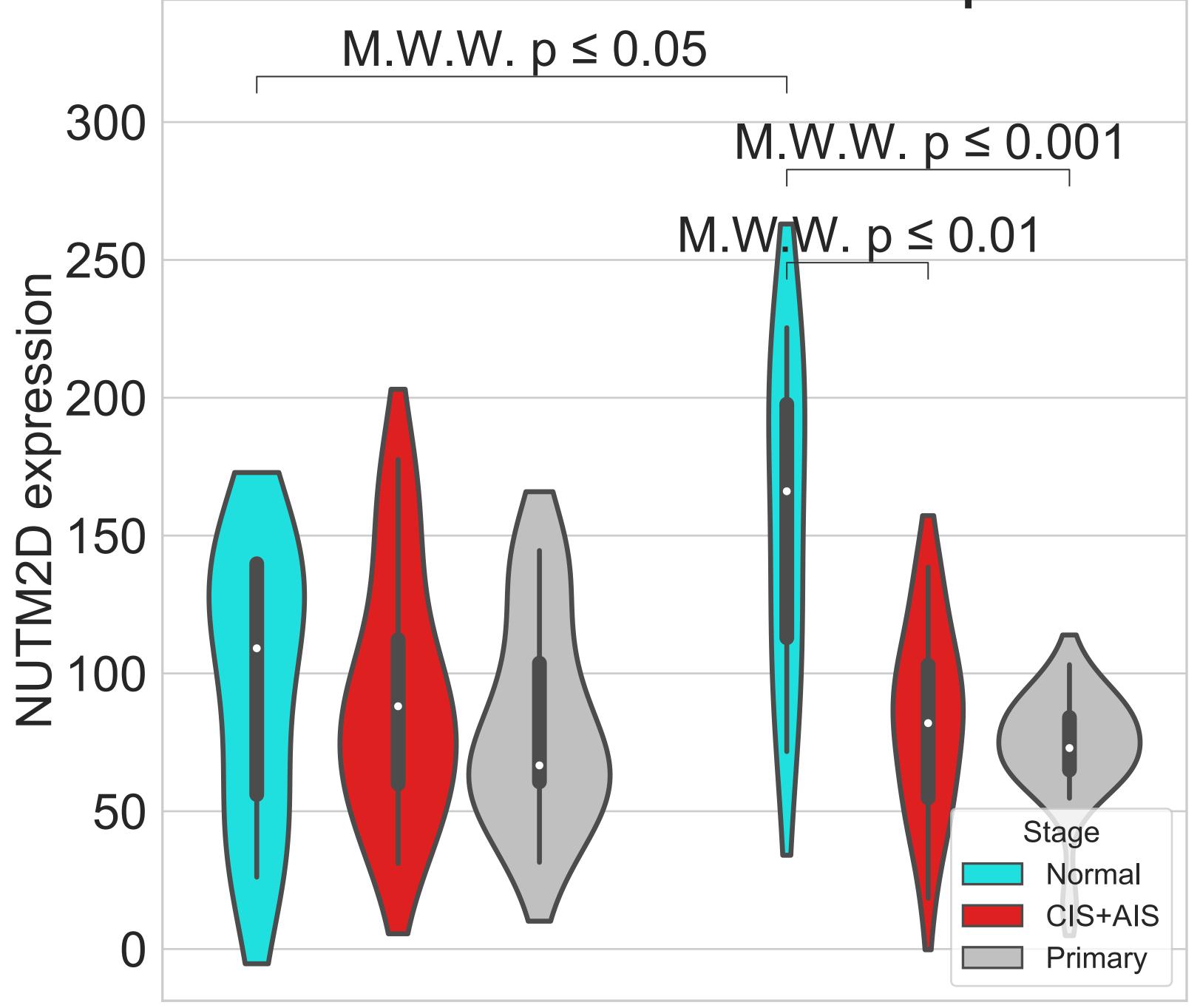
NUTM2D: Kruskal-Wallis p=0.019



Lower Higher Mutation Shared Count (SYN)