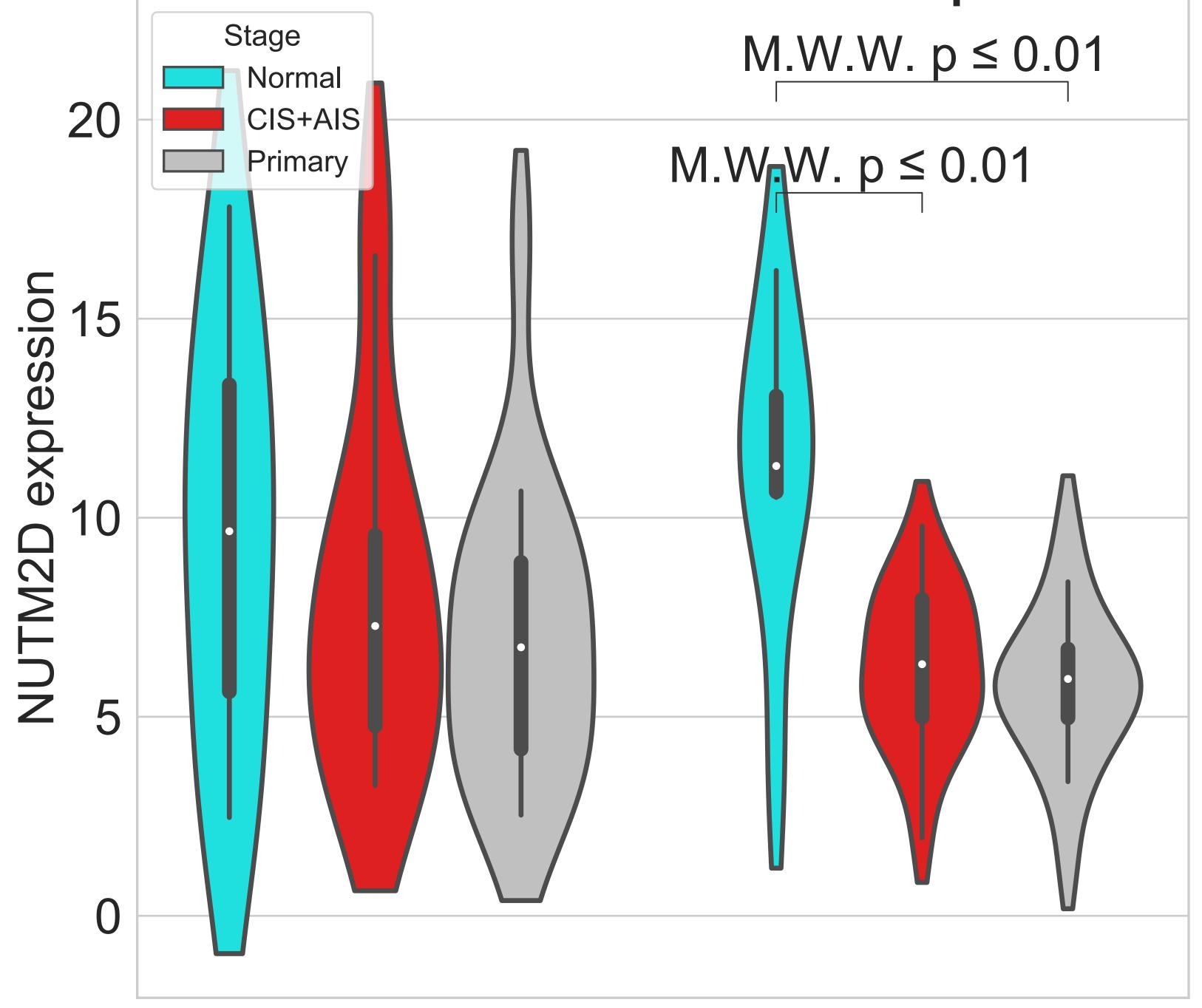
NUTM2D: Kruskal-Wallis p=0.038



Lower
Mutation Shared Count (SYN)