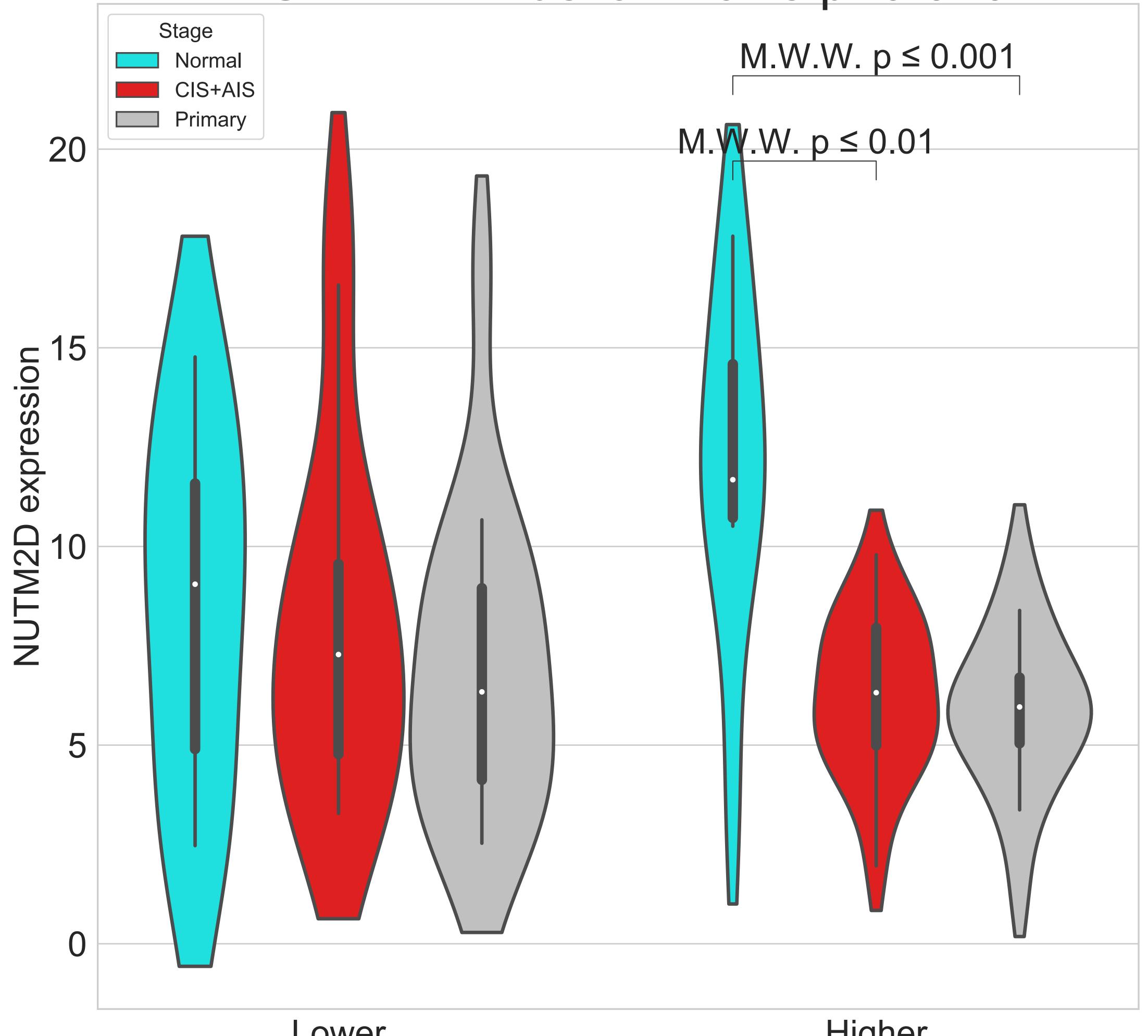
NUTM2D: Kruskal-Wallis p=0.026



Lower
Mutation Shared Count (SYN)