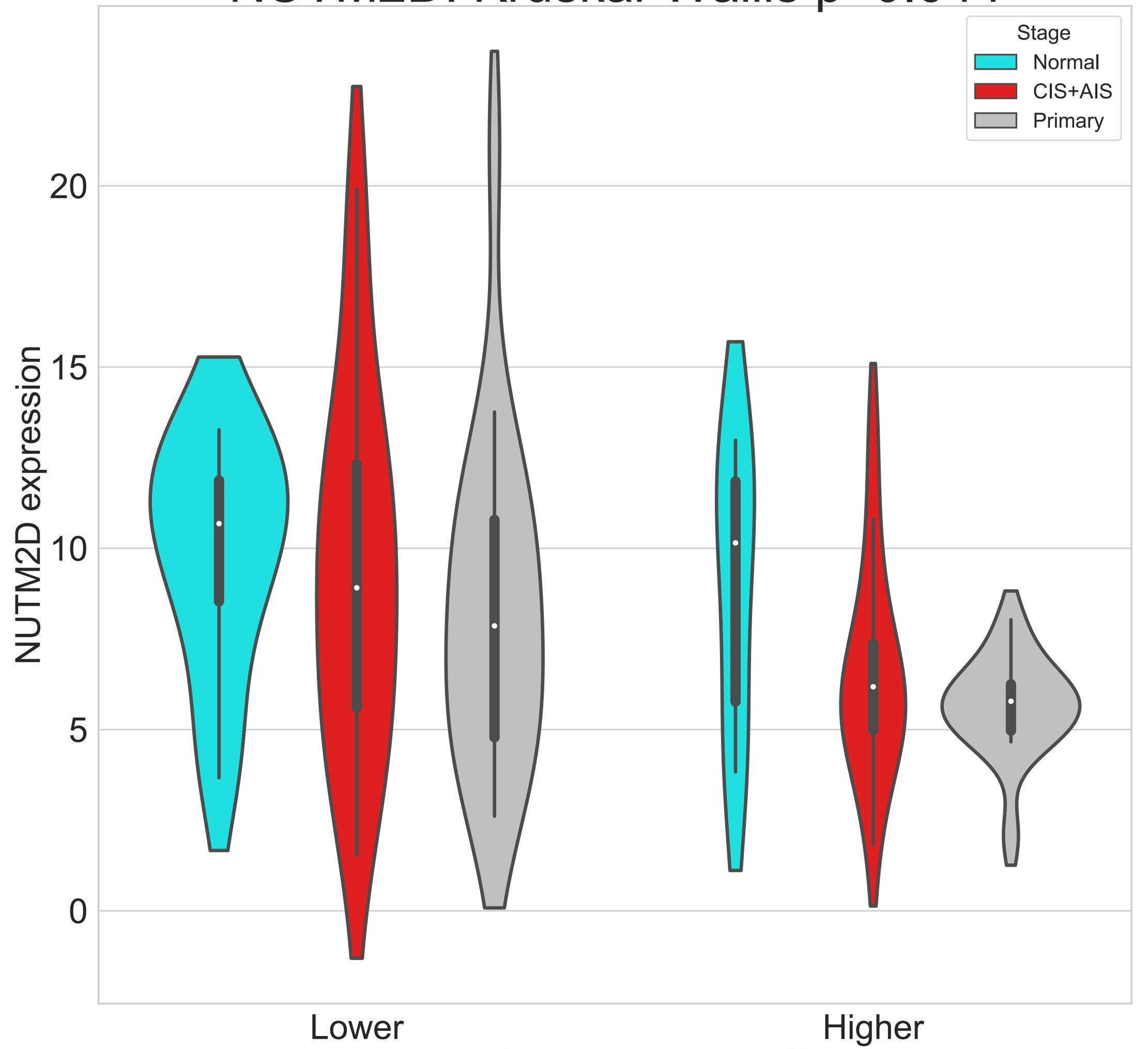
NUTM2D: Kruskal-Wallis p=0.044



Mutation Shared Proportion (SYN)