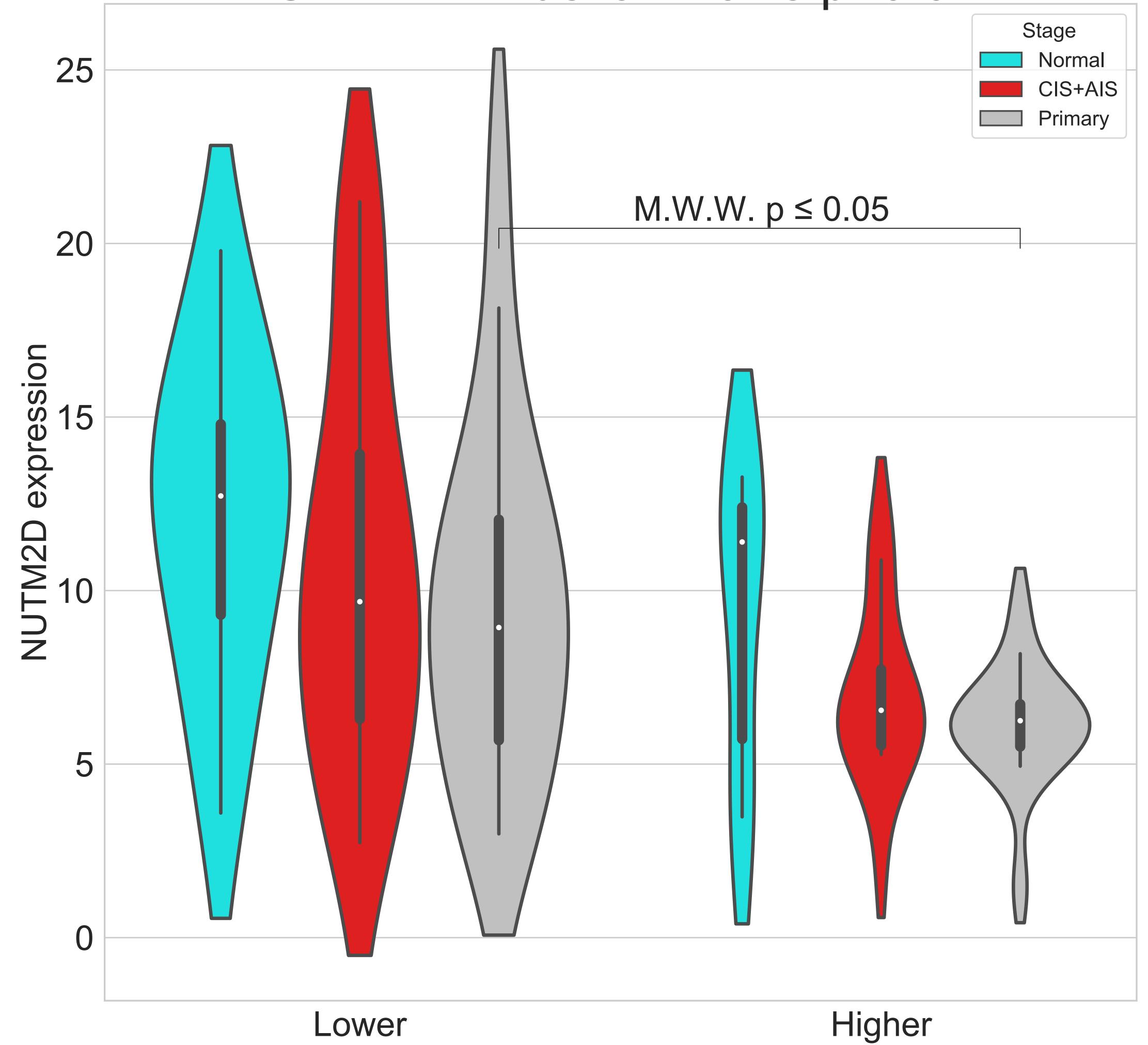
NUTM2D: Kruskal-Wallis p=0.012



Mutation Shared Proportion (SYN)