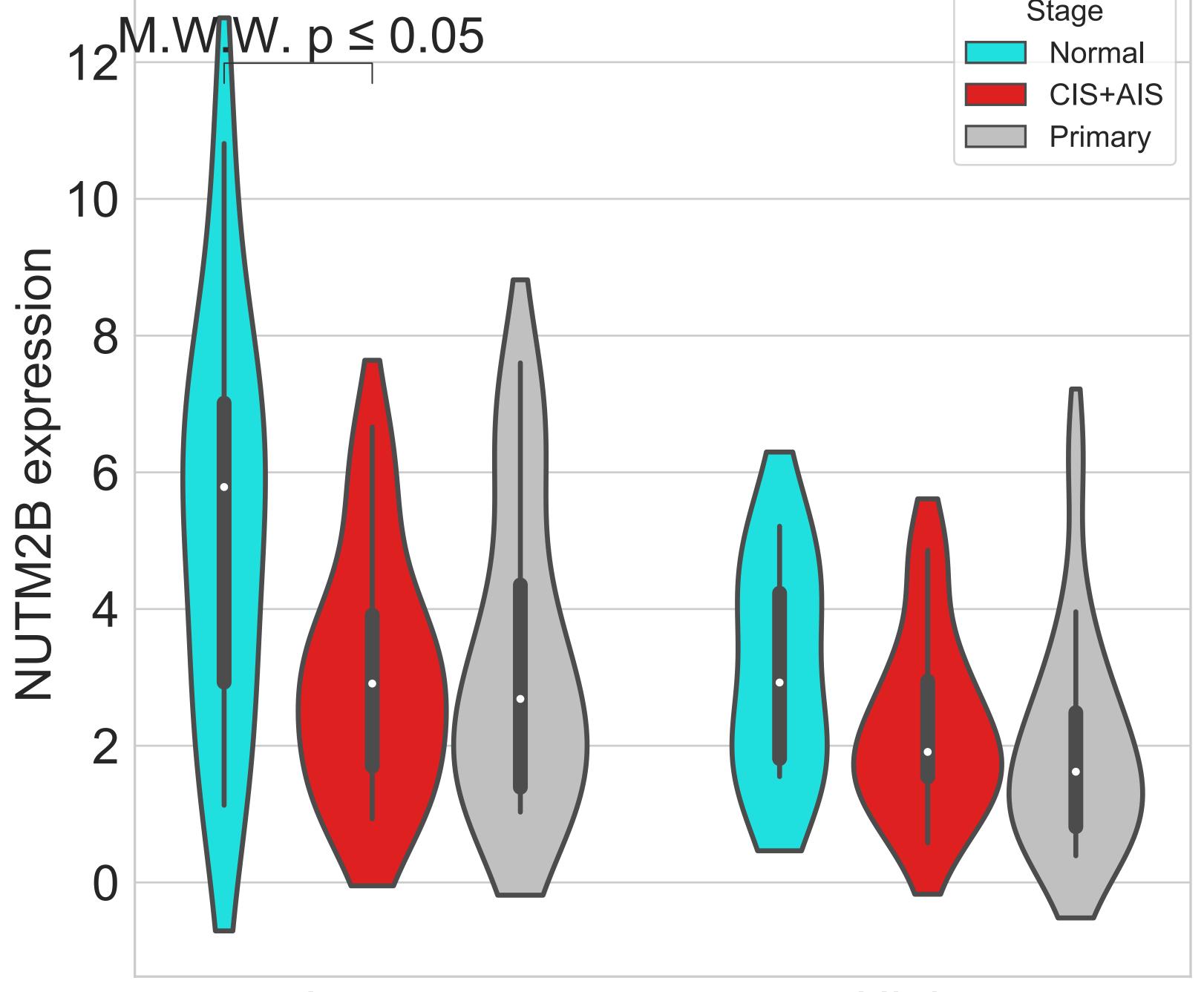
NUTM2B: Kruskal-Wallis p=0.015 Normal Normal



Lower Higher Mutation Shared Proportion (SYN)