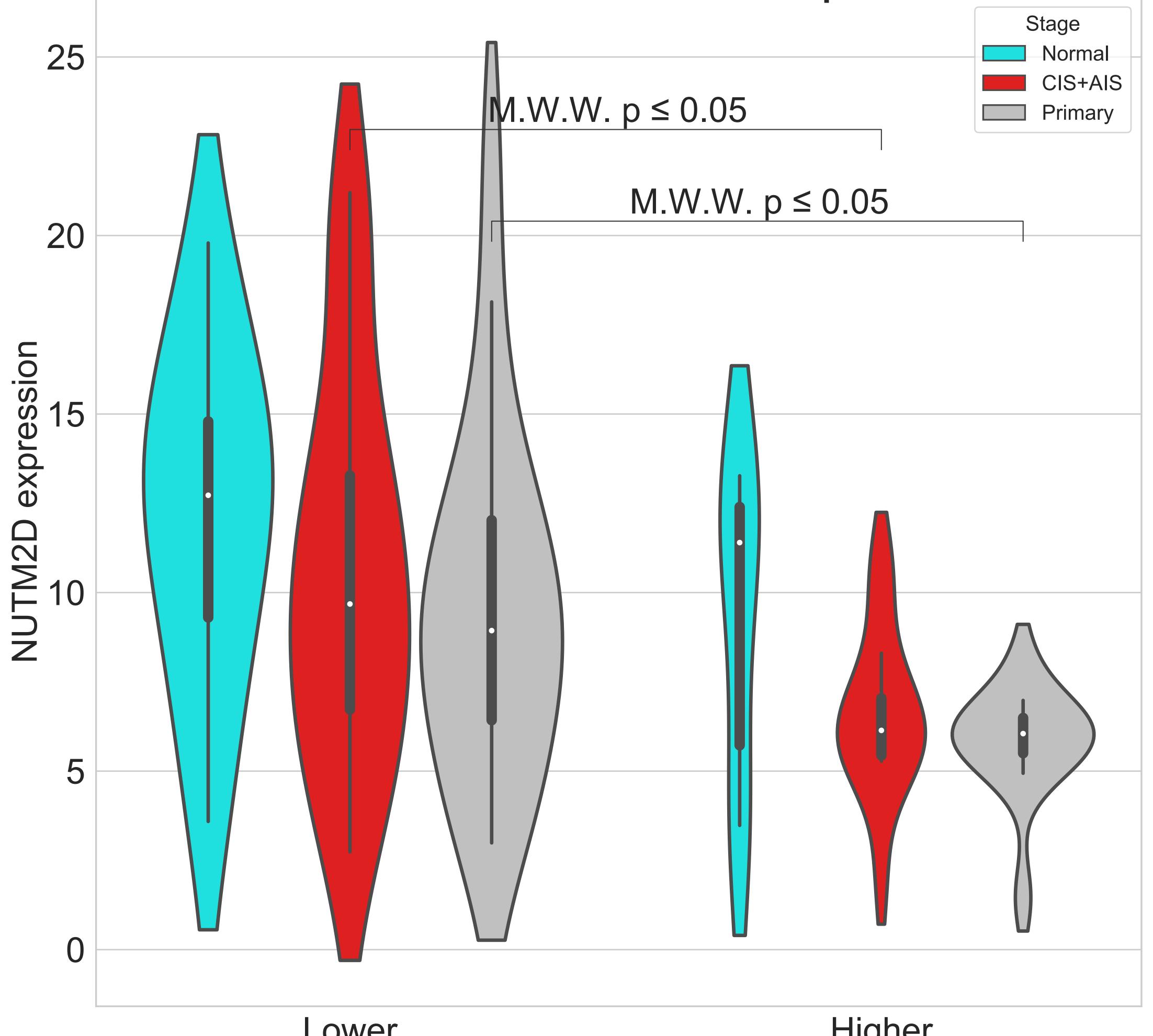
NUTM2D: Kruskal-Wallis p=0.004



Higher Lower Mutation Shared Proportion (Union)