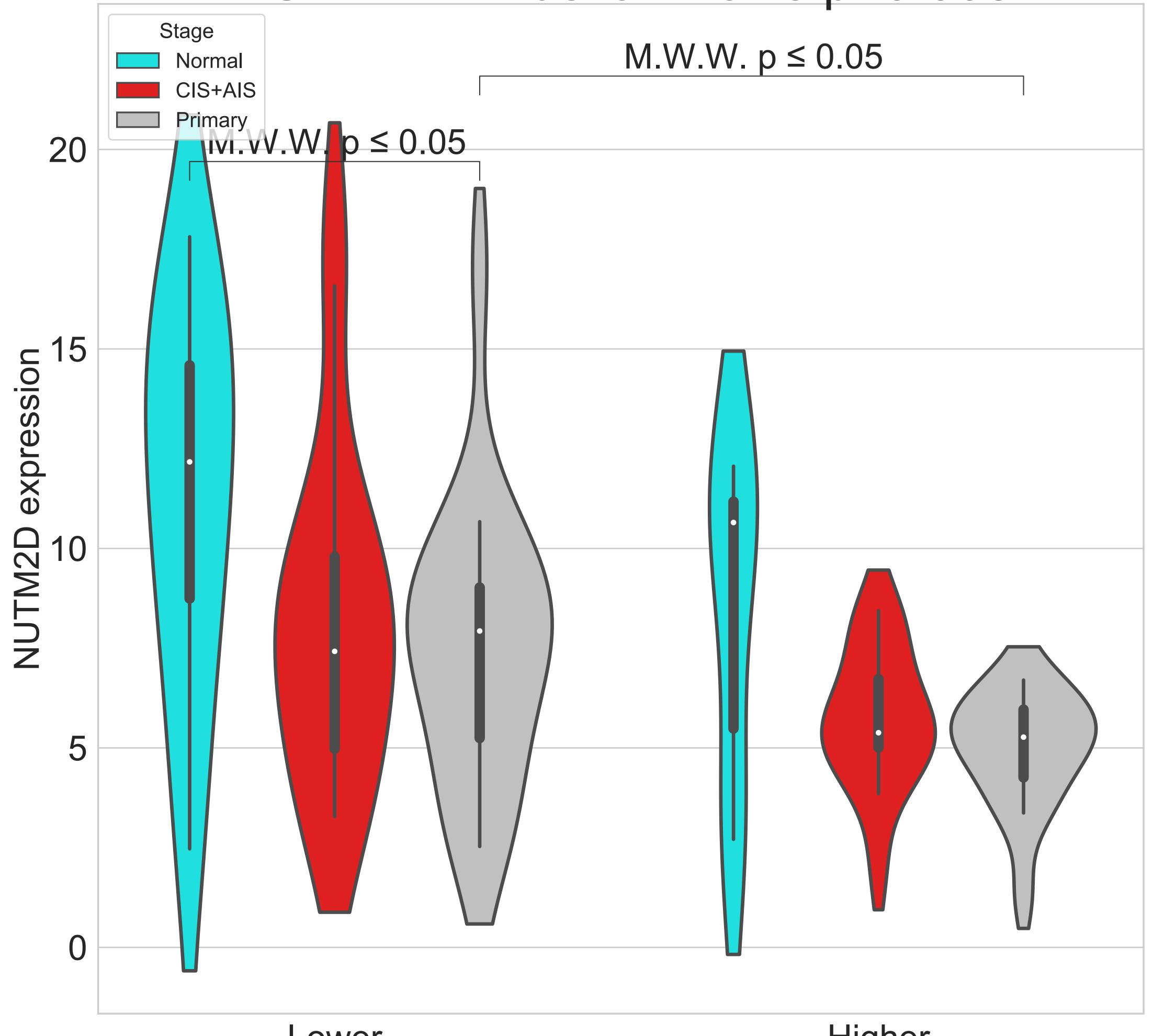
NUTM2D: Kruskal-Wallis p=0.003



Lower
Mutation Shared Proportion (Union & SYN)