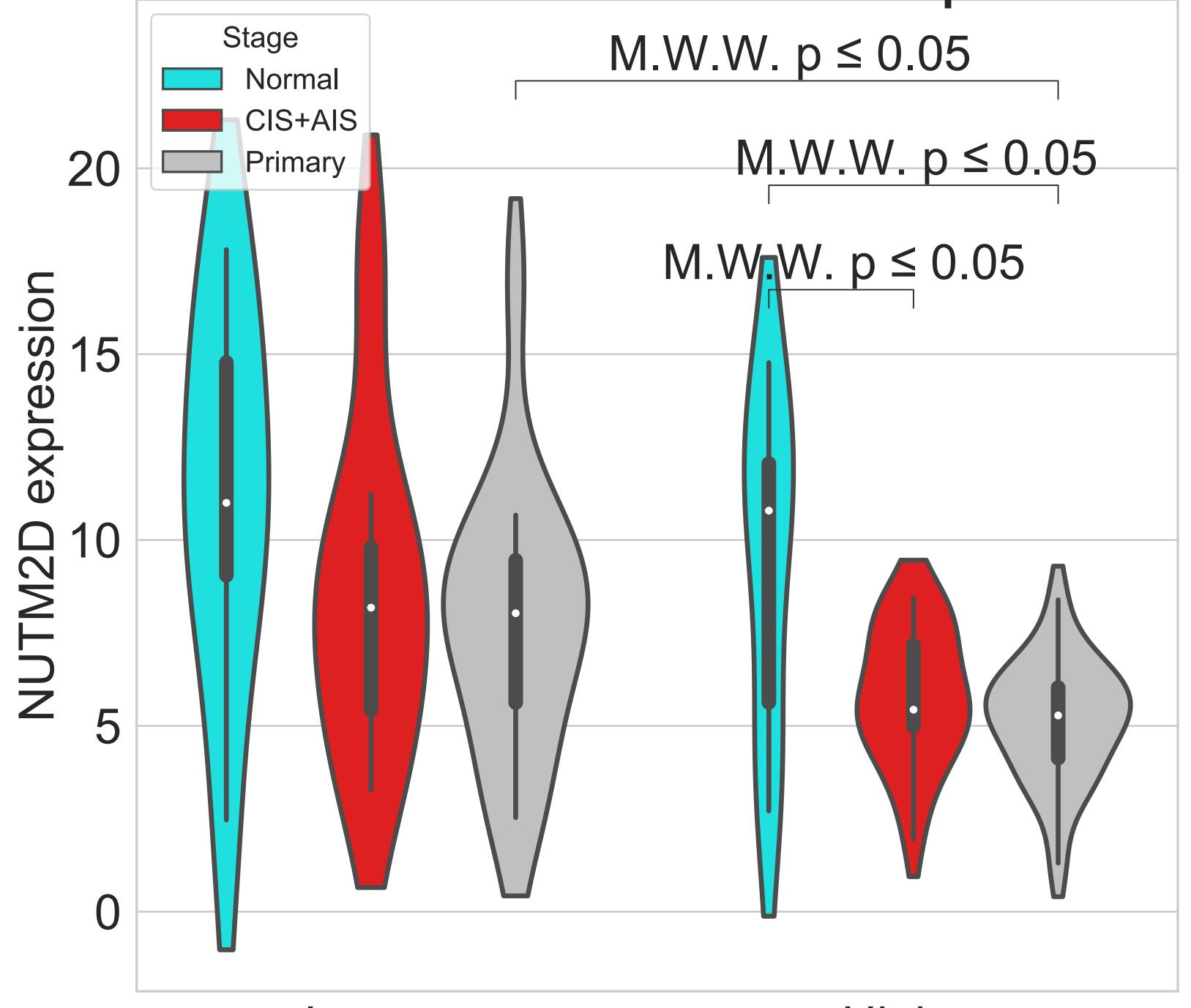
NUTM2D: Kruskal-Wallis p=0.003



Lower
Higher
Mutation Shared Proportion