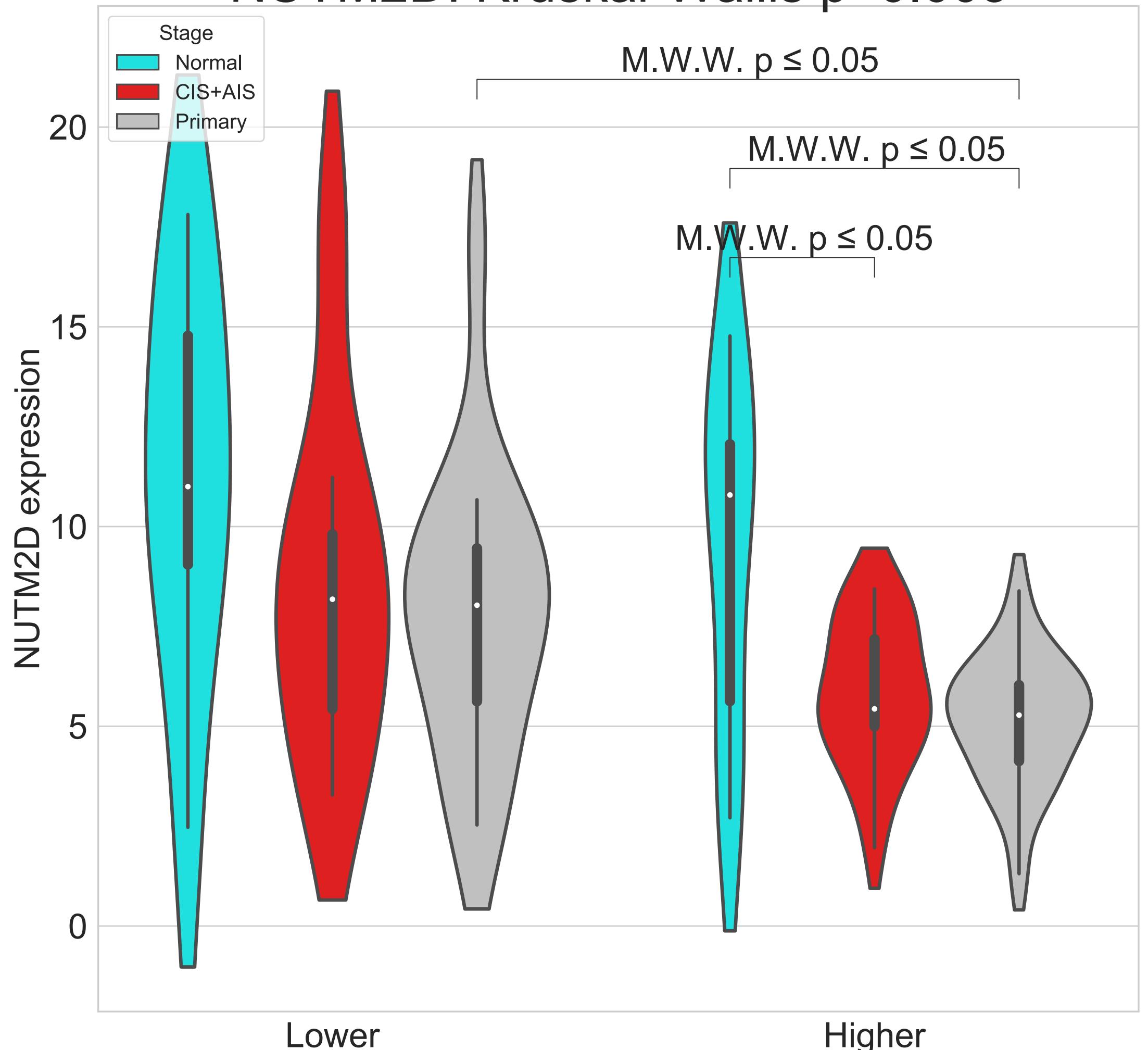
NUTM2D: Kruskal-Wallis p=0.003



wer
Mutation Shared Proportion