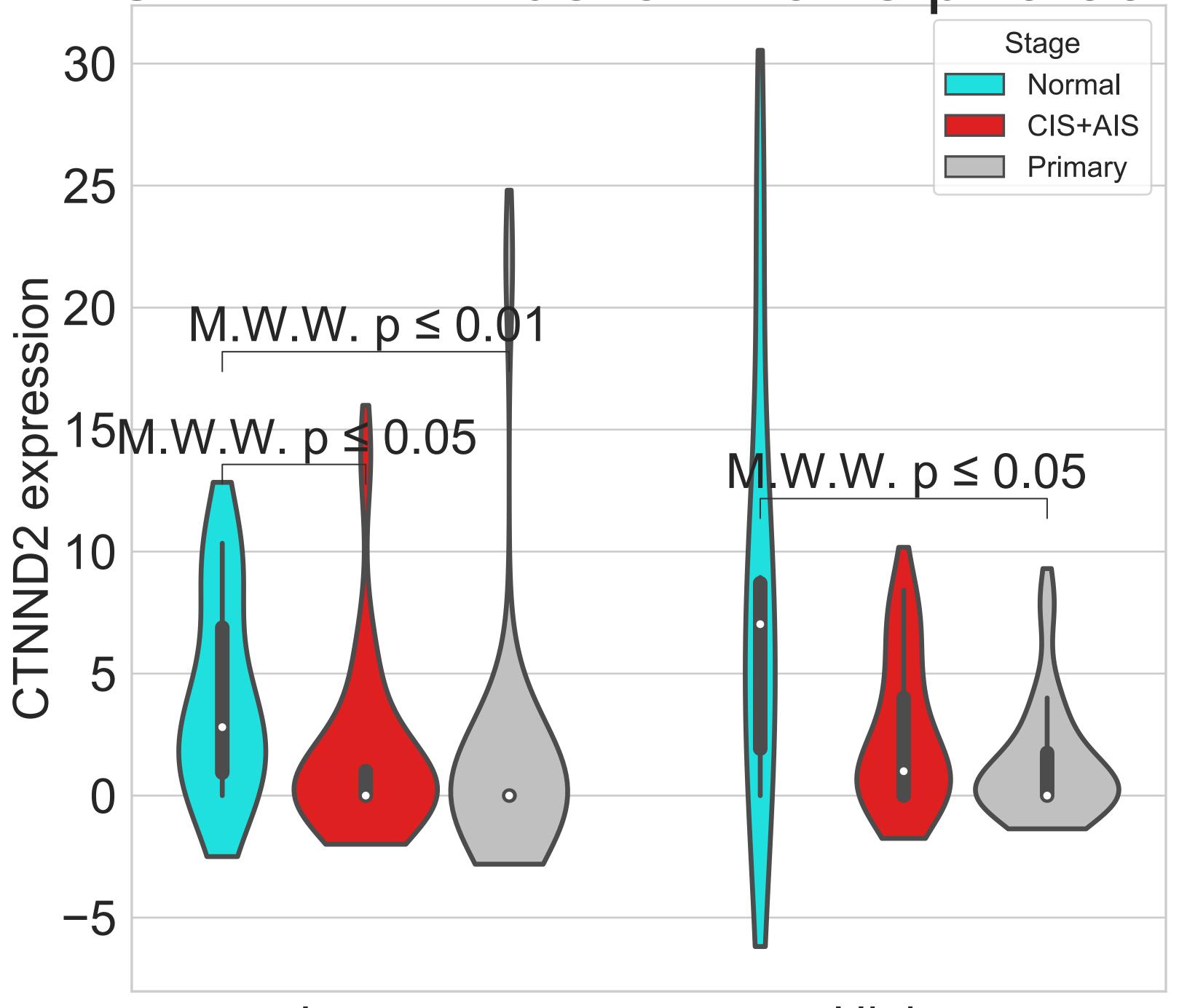
CTNND2: Kruskal-Wallis p=0.004



Lower
Higher
Mutation Shared Proportion