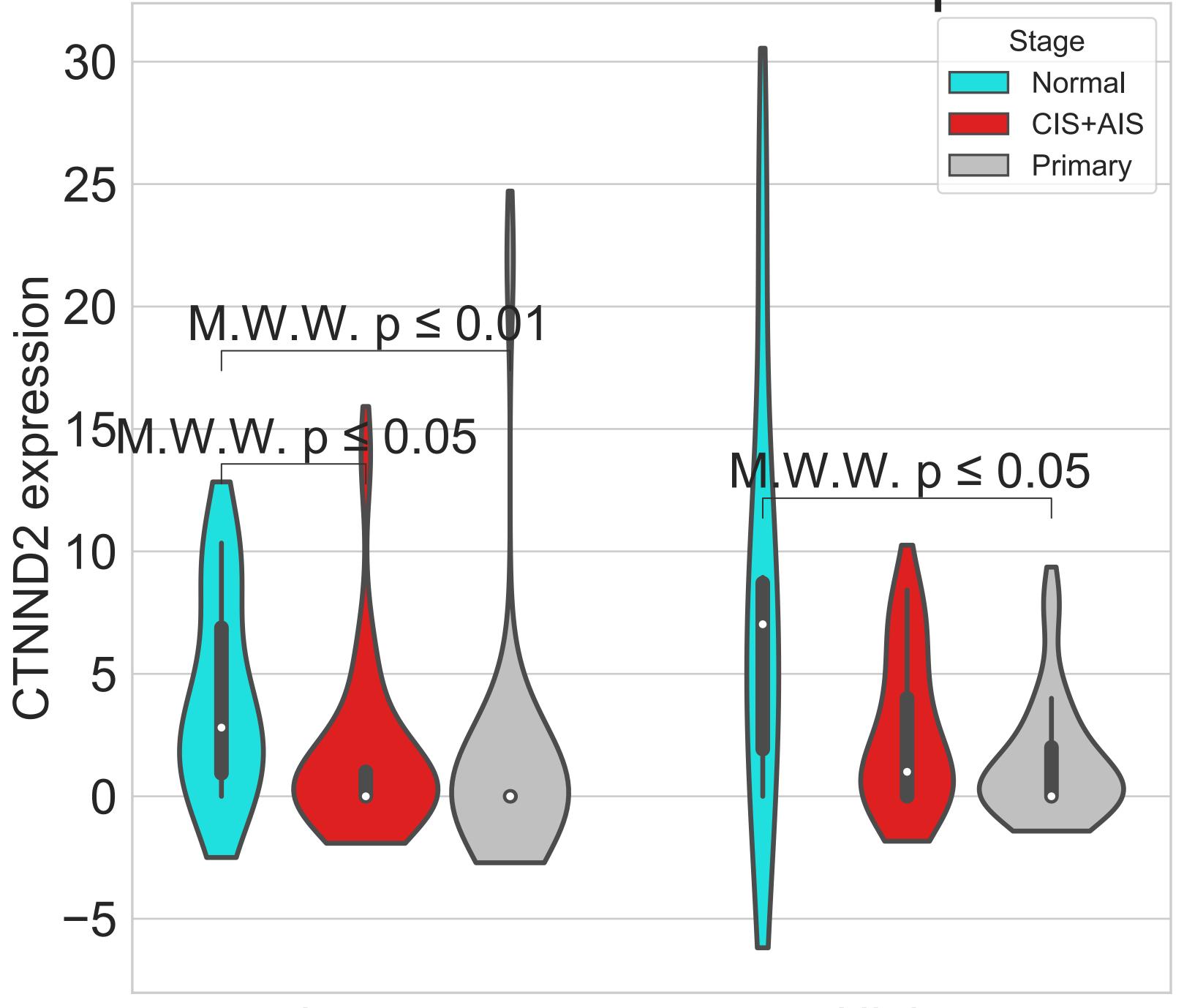
CTNND2: Kruskal-Wallis p=0.004



Lower Higher Mutation Shared Proportion (Union)