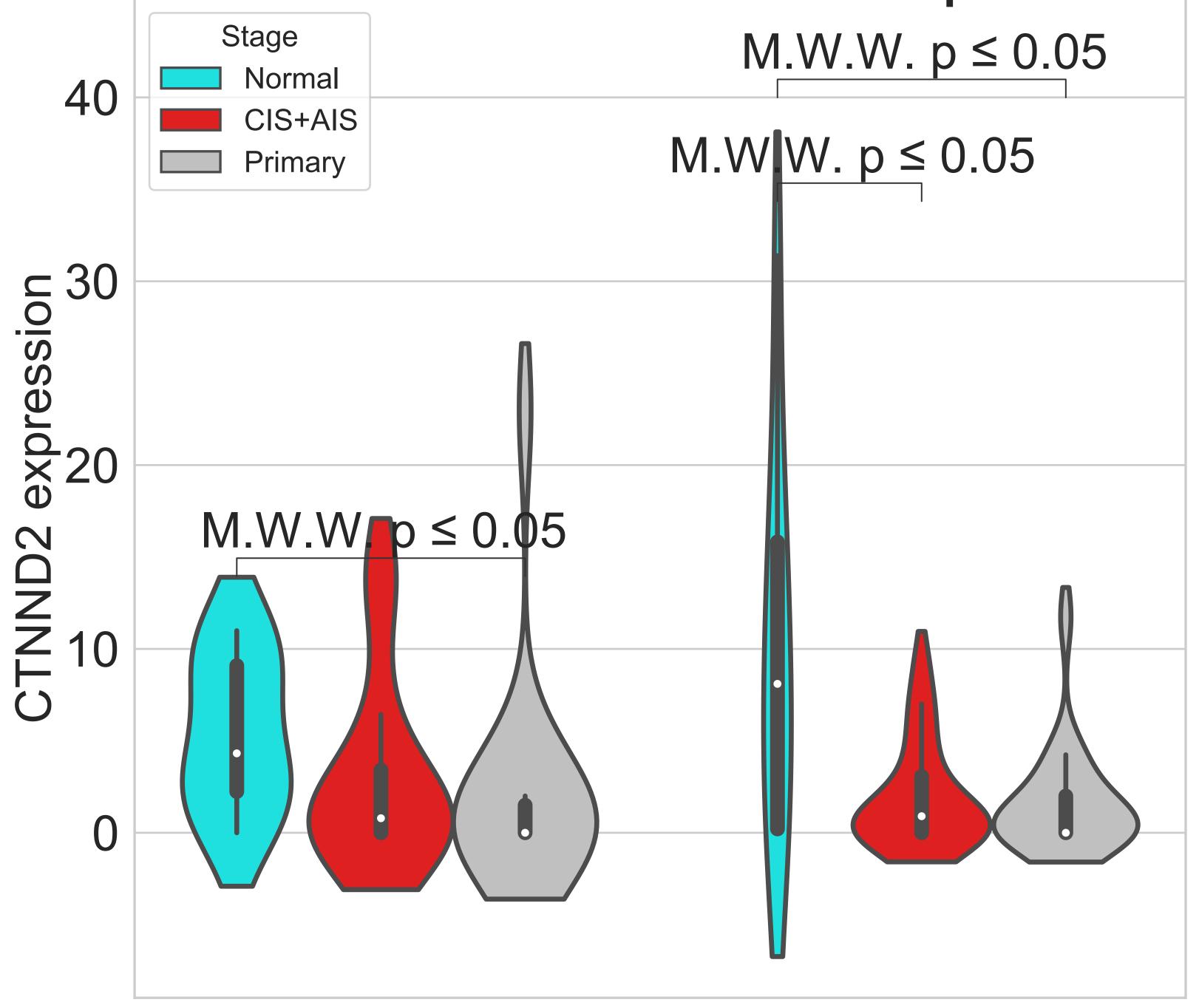
CTNND2: Kruskal-Wallis p=0.012



Lower Higher Mutation Shared Proportion (SYN)