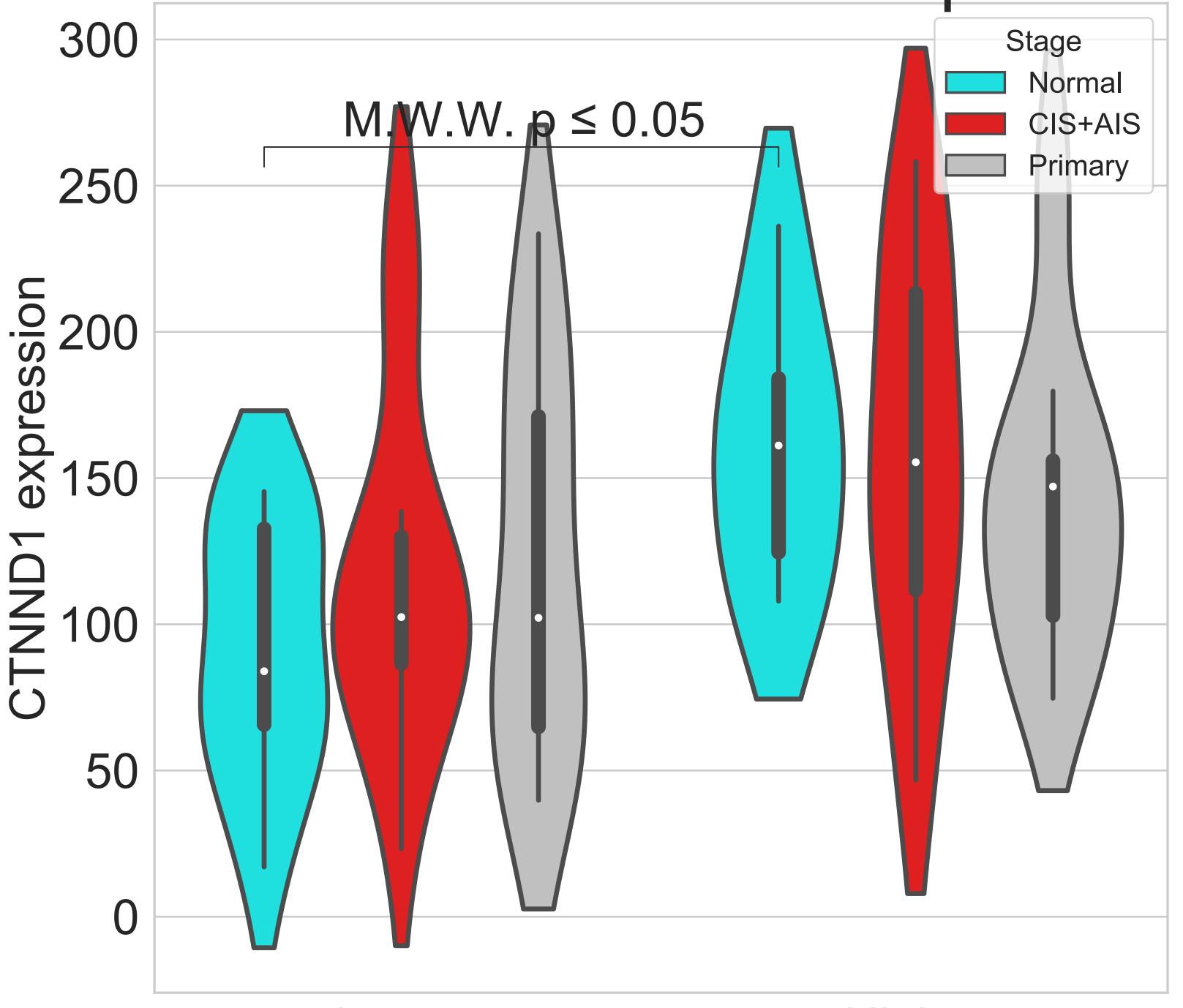
CTNND1: Kruskal-Wallis p=0.044



Lower Higher Mutation Shared Proportion (SYN)