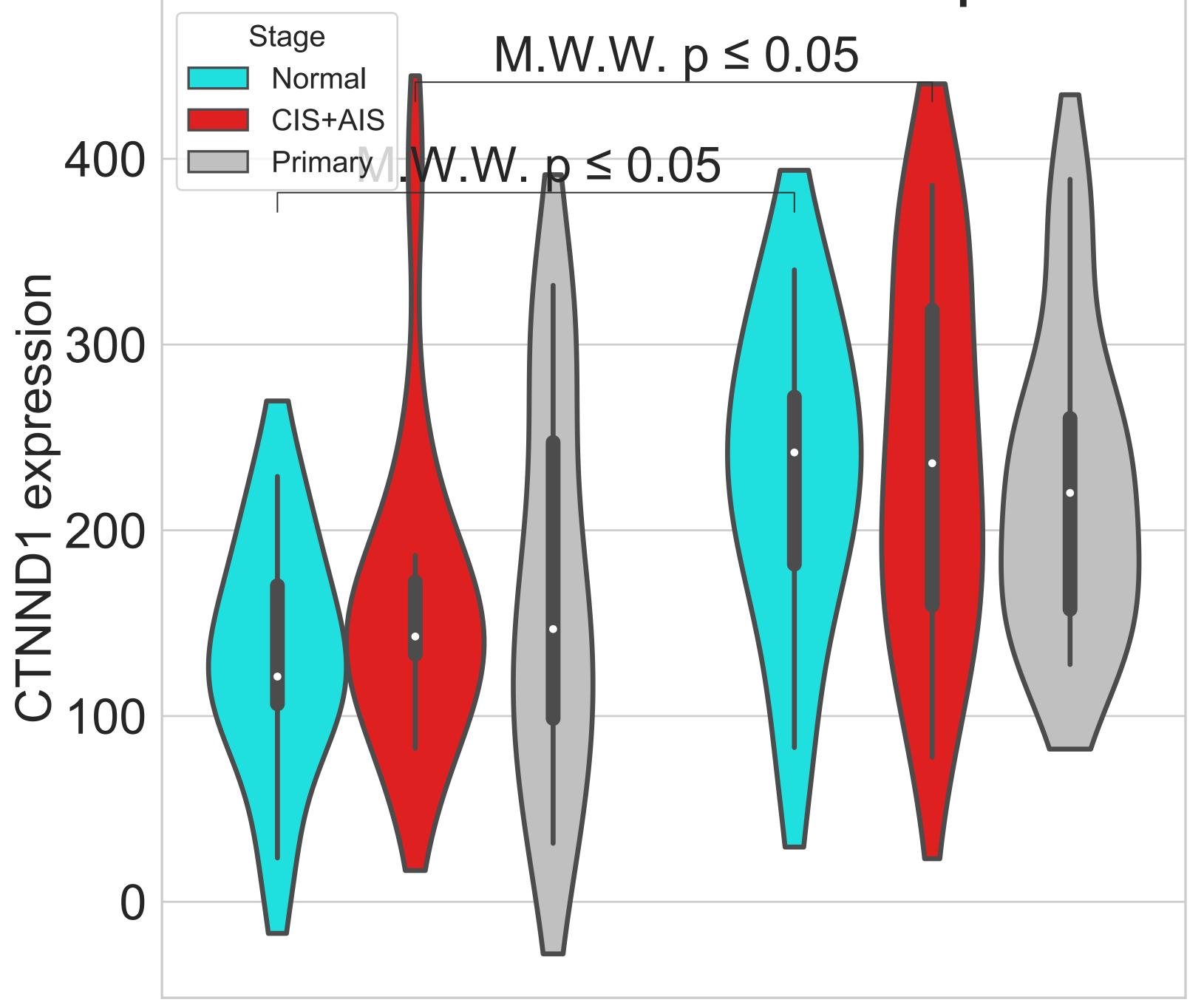
CTNND1: Kruskal-Wallis p=0.009



Lower Higher Mutation Shared Proportion (Union & SYN)