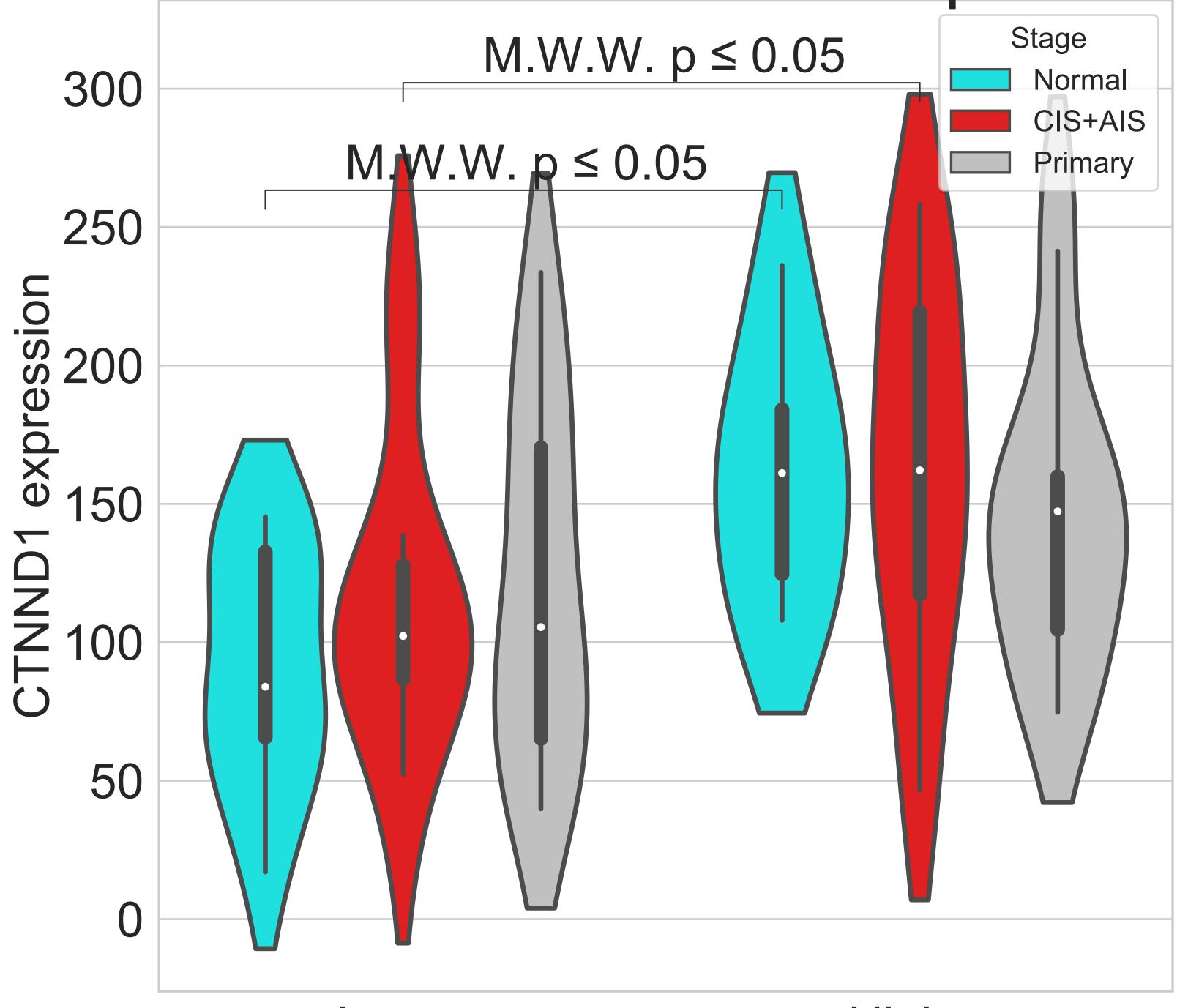
CTNND1: Kruskal-Wallis p=0.028



Lower Higher Mutation Shared Proportion