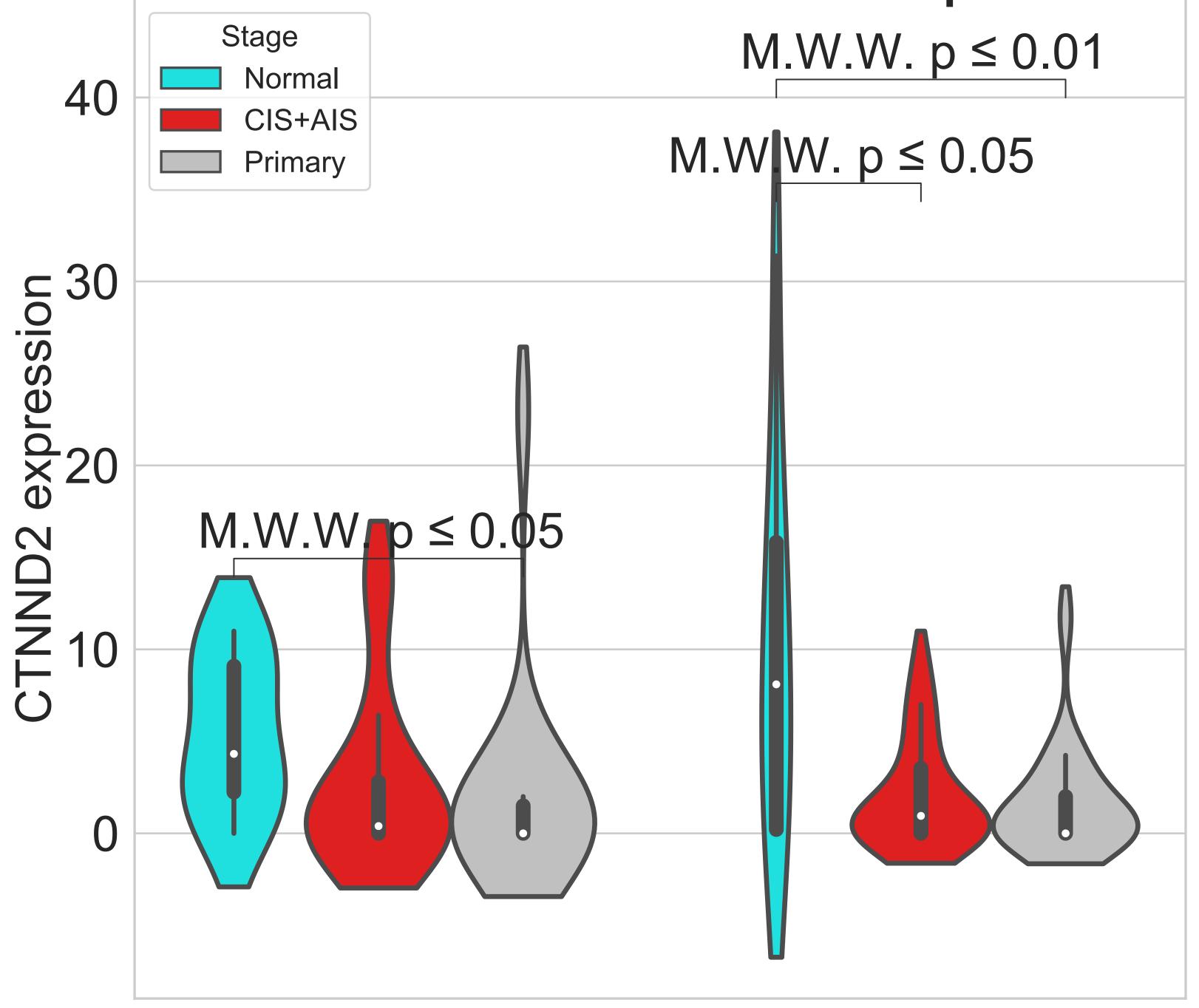
CTNND2: Kruskal-Wallis p=0.013



Lower

Mutation Shared Proportion