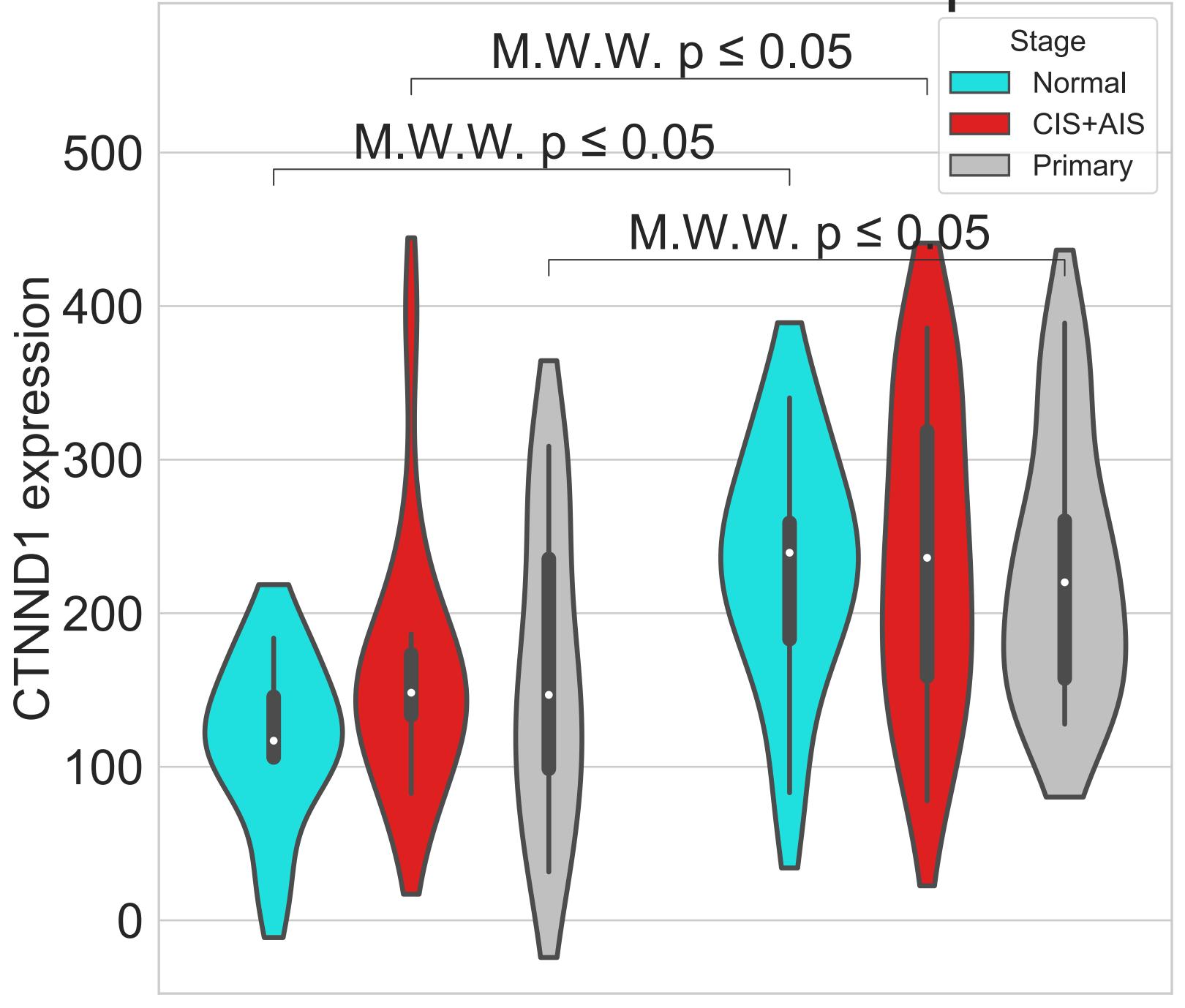
CTNND1: Kruskal-Wallis p=0.005



Lower

Mutation Shared Proportion