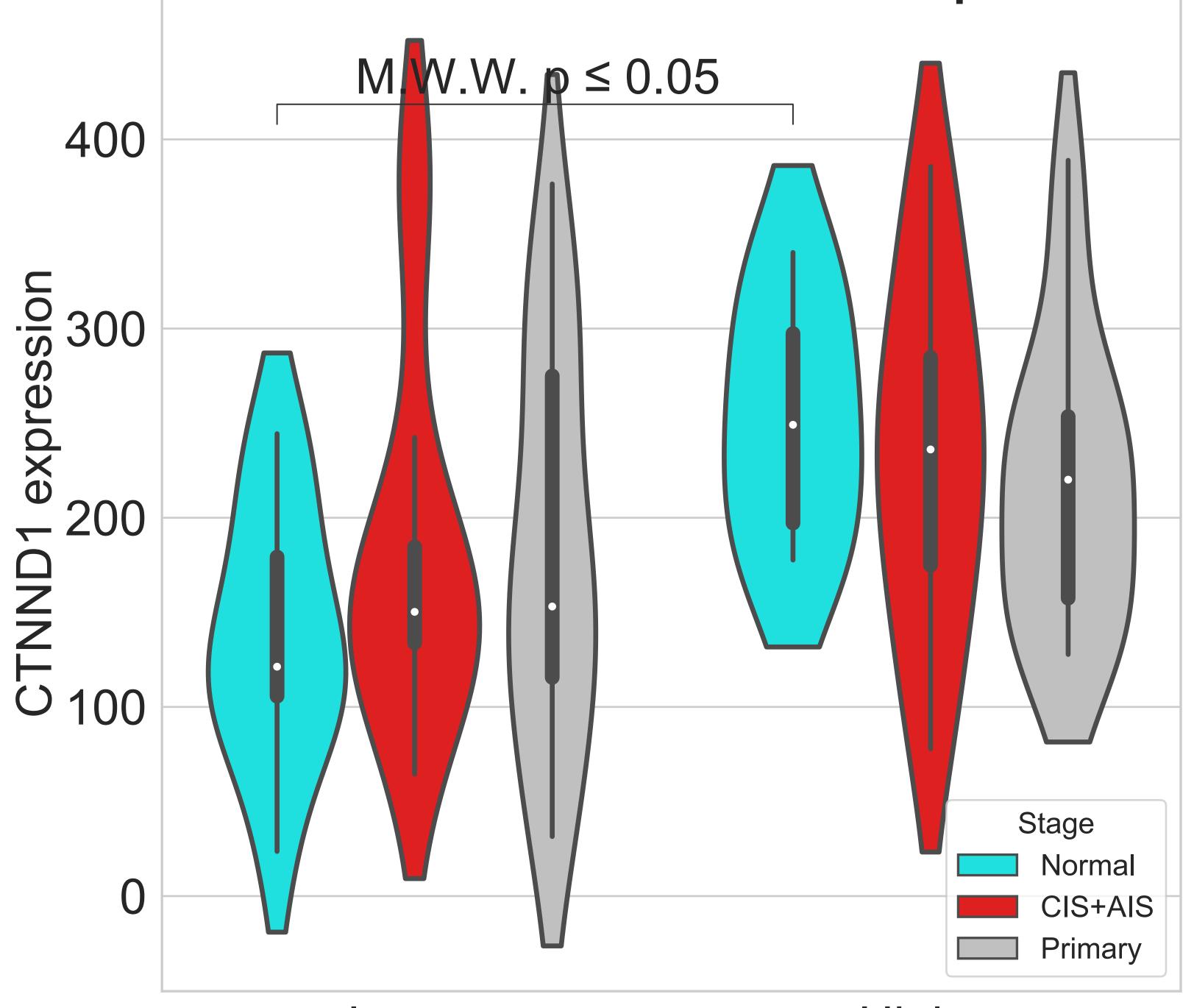
CTNND1: Kruskal-Wallis p=0.048



Lower

Mutation Shared Proportion