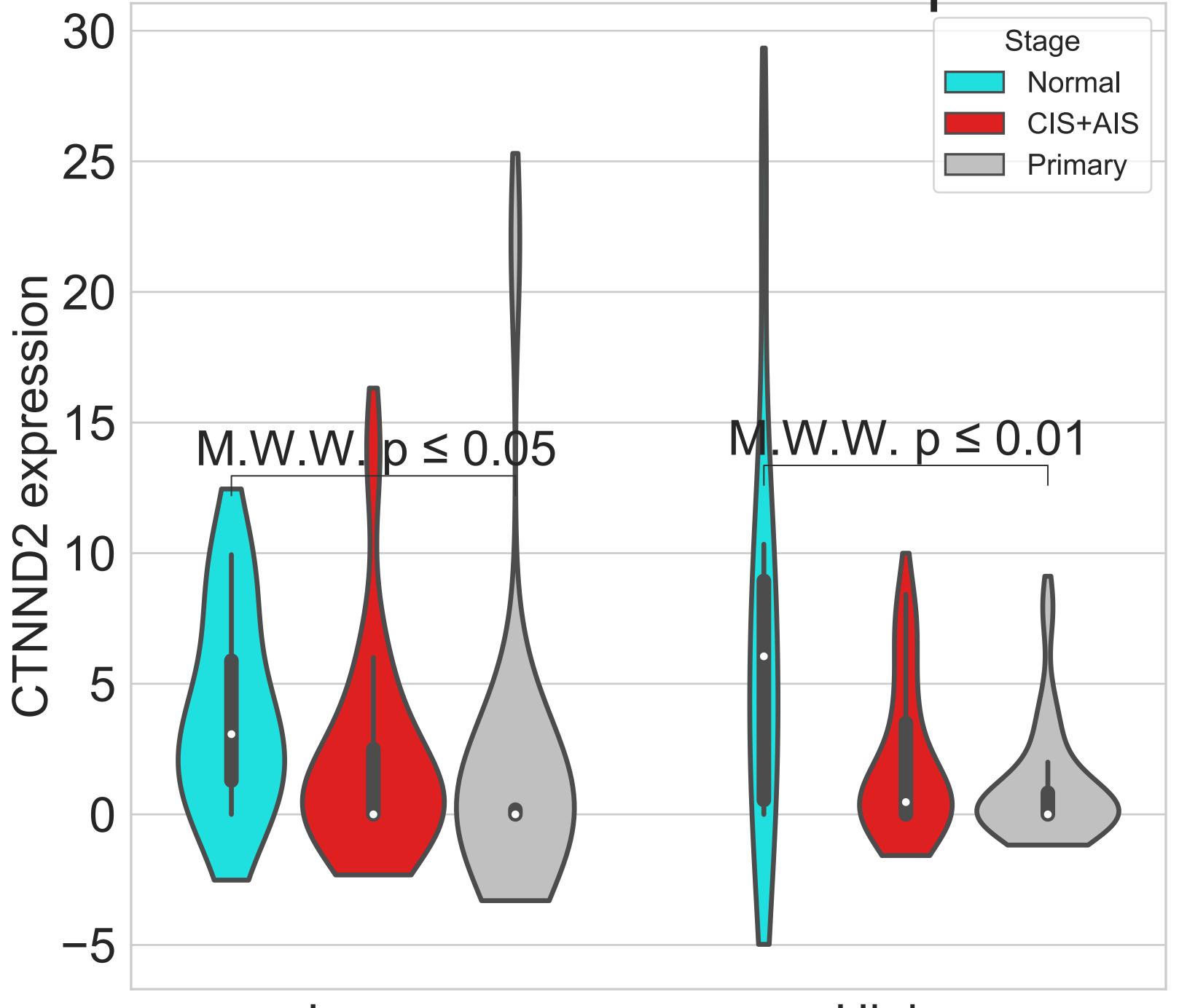
CTNND2: Kruskal-Wallis p=0.016



Lower

Higher

Mutation Shared Proportion