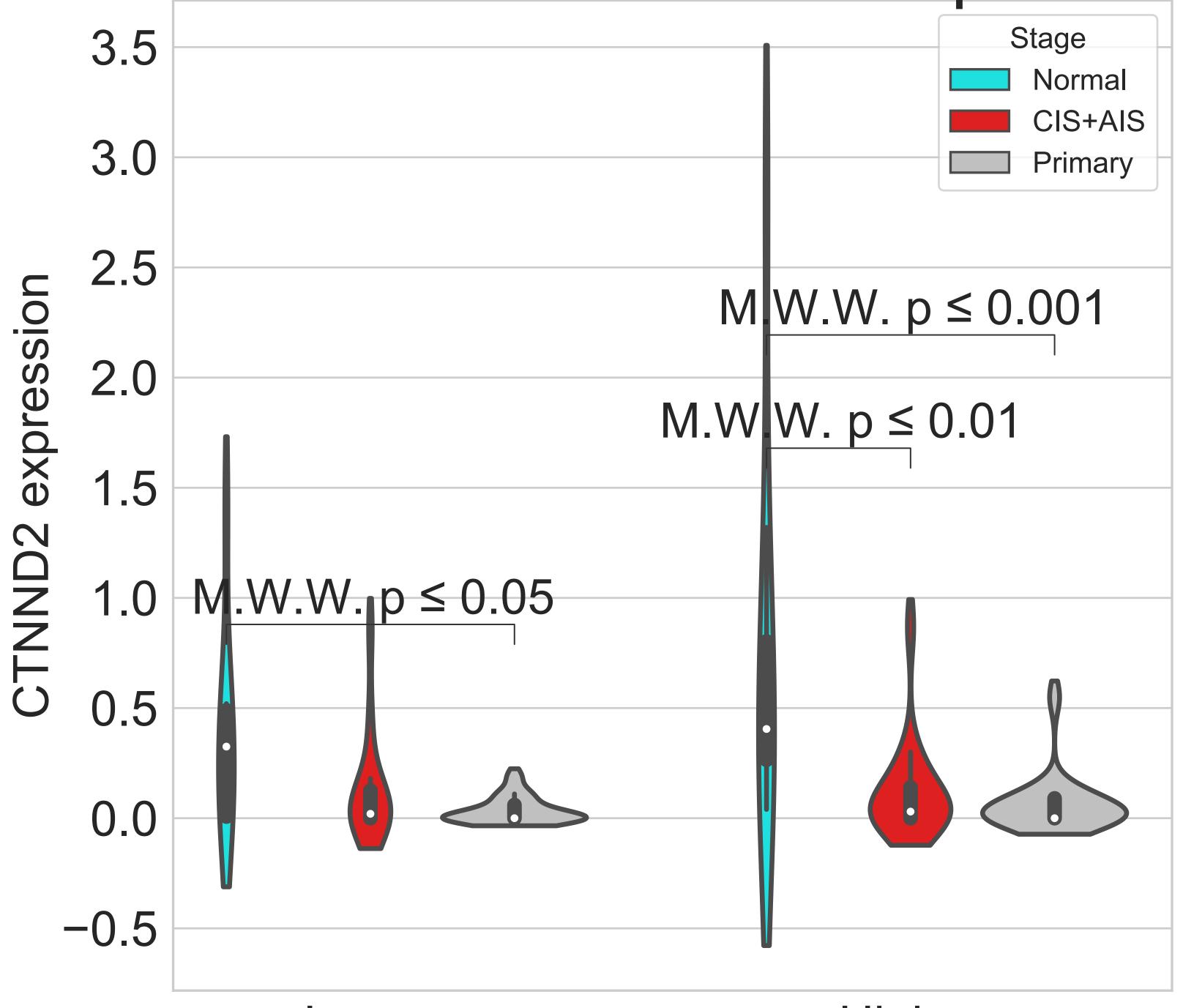
CTNND2: Kruskal-Wallis p=0.001



Lower

Higher

Mutation Shared Count (SYN)