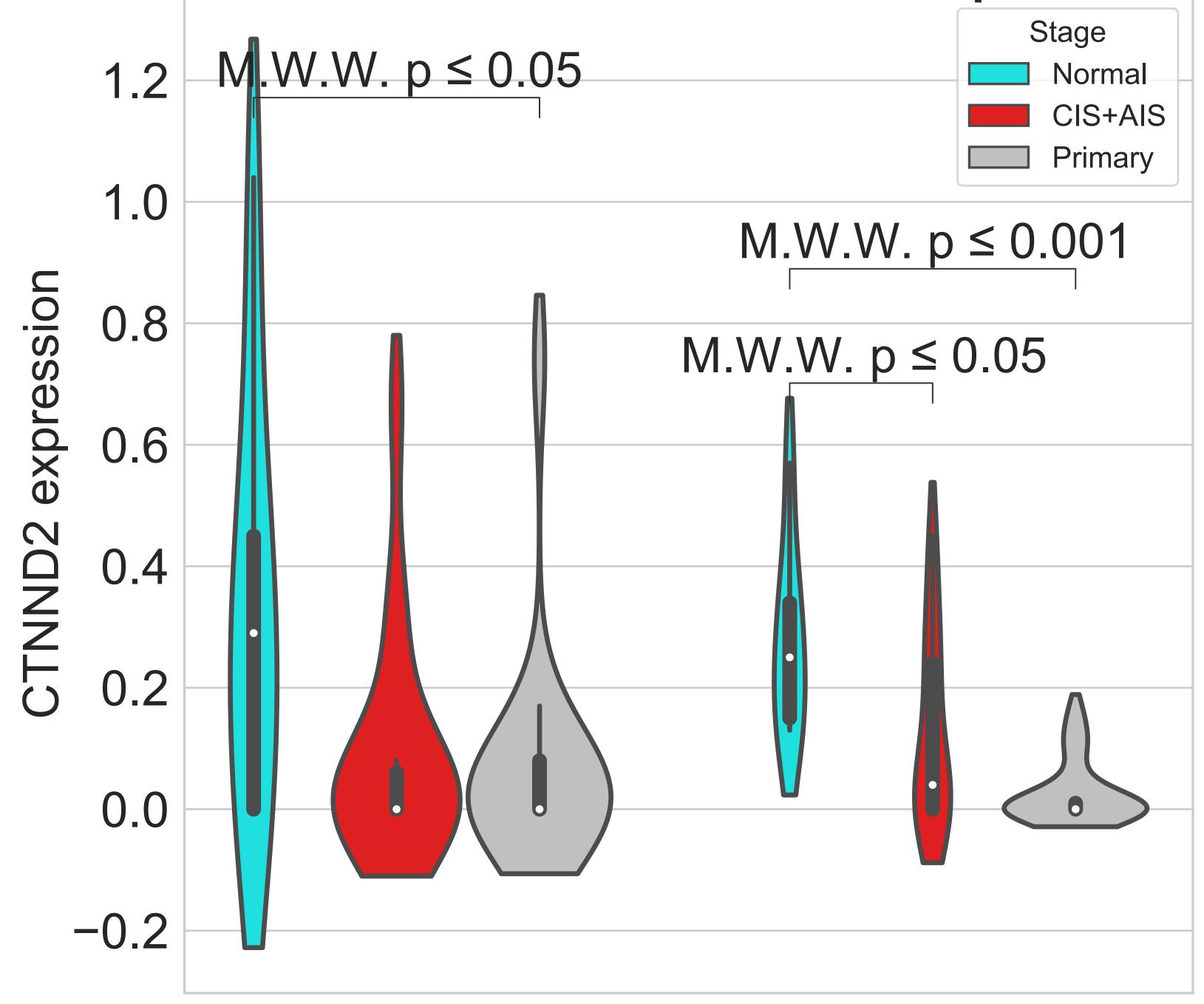
CTNND2: Kruskal-Wallis p=0.001



Lower
Higher
Mutation Shared Count (SYN)