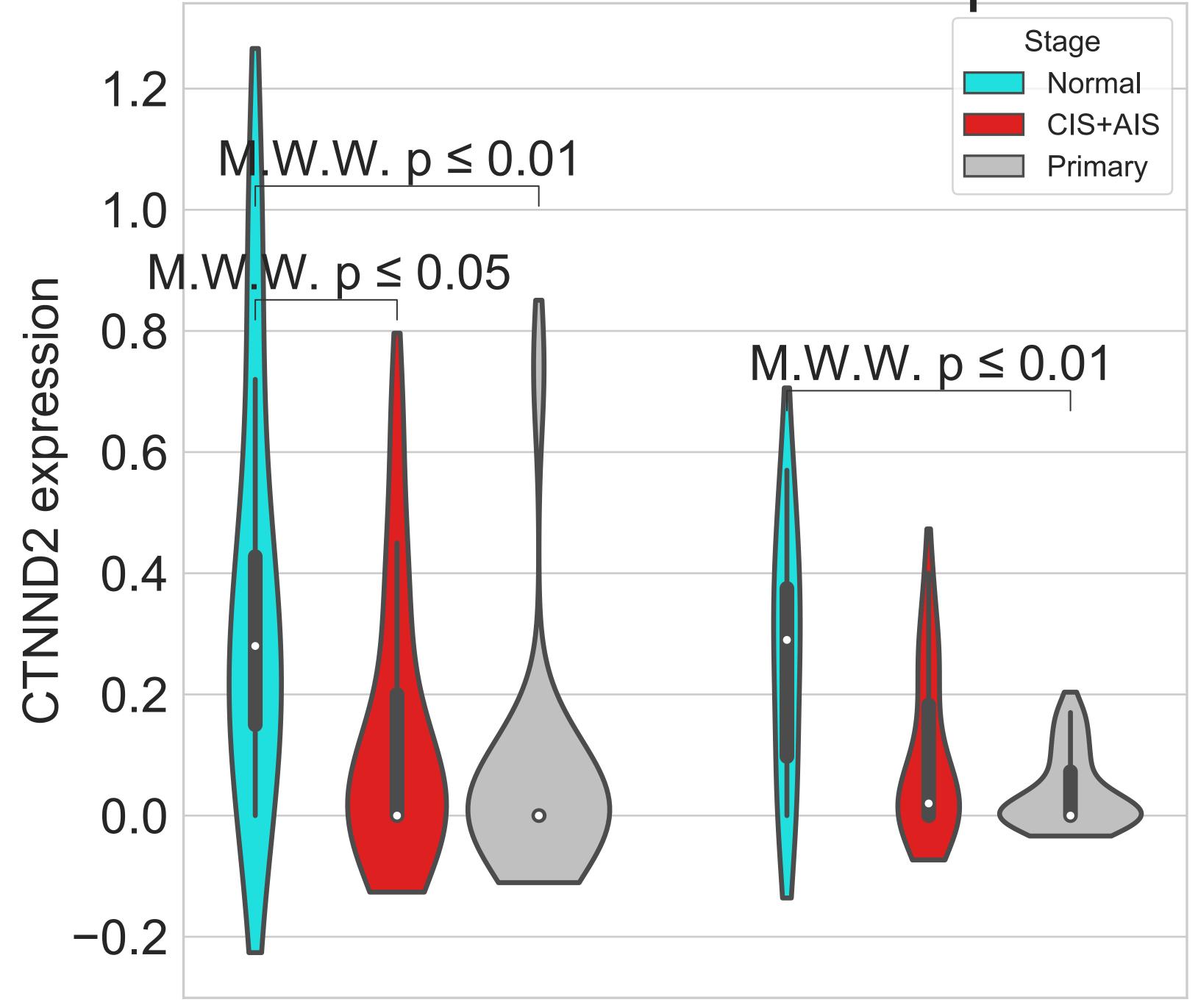
CTNND2: Kruskal-Wallis p=0.002



Lower

Mutation Shared Proportion (Union & SYN)