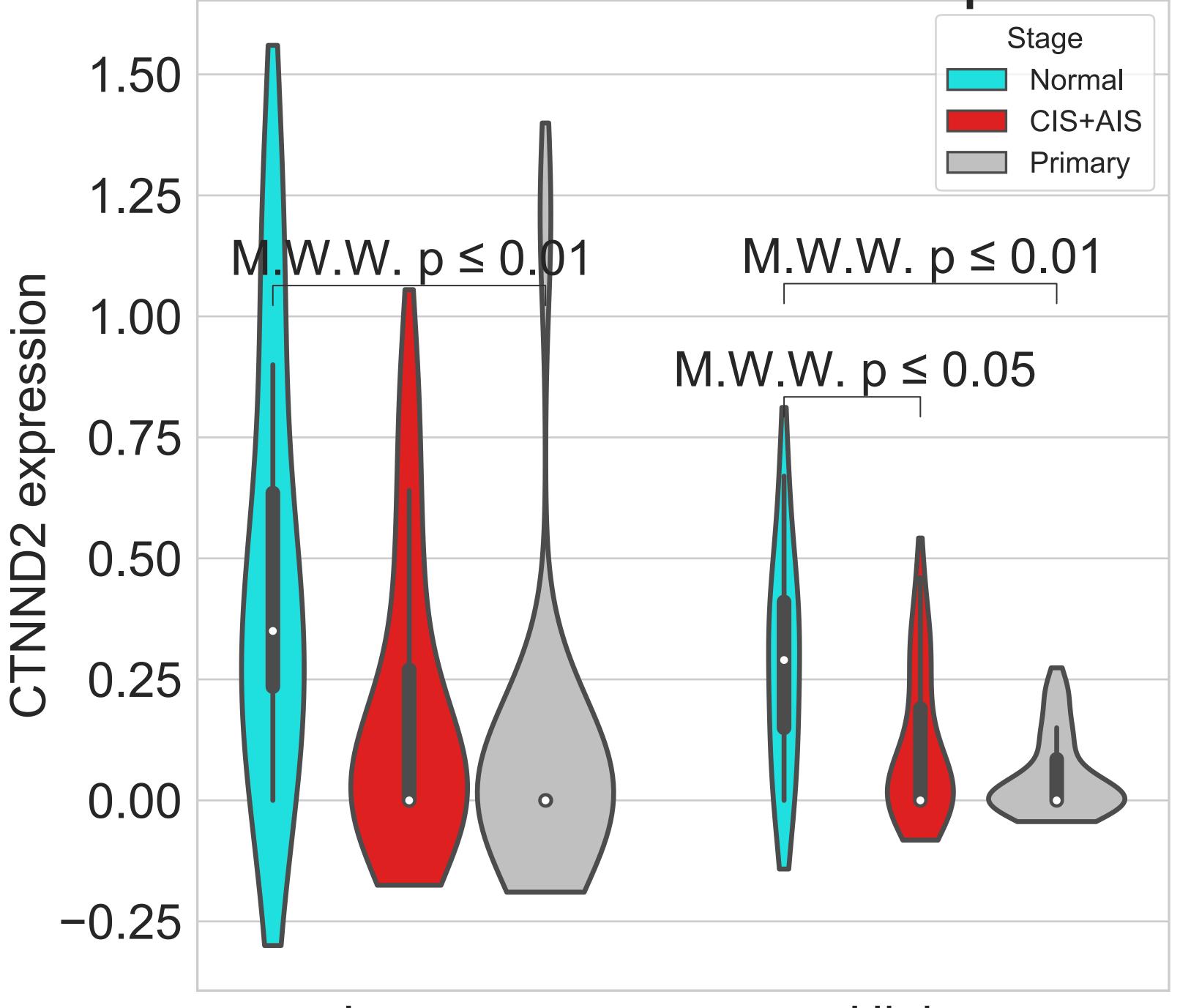
CTNND2: Kruskal-Wallis p=0.003



Lower
Mutation Shared Proportion (SYN)