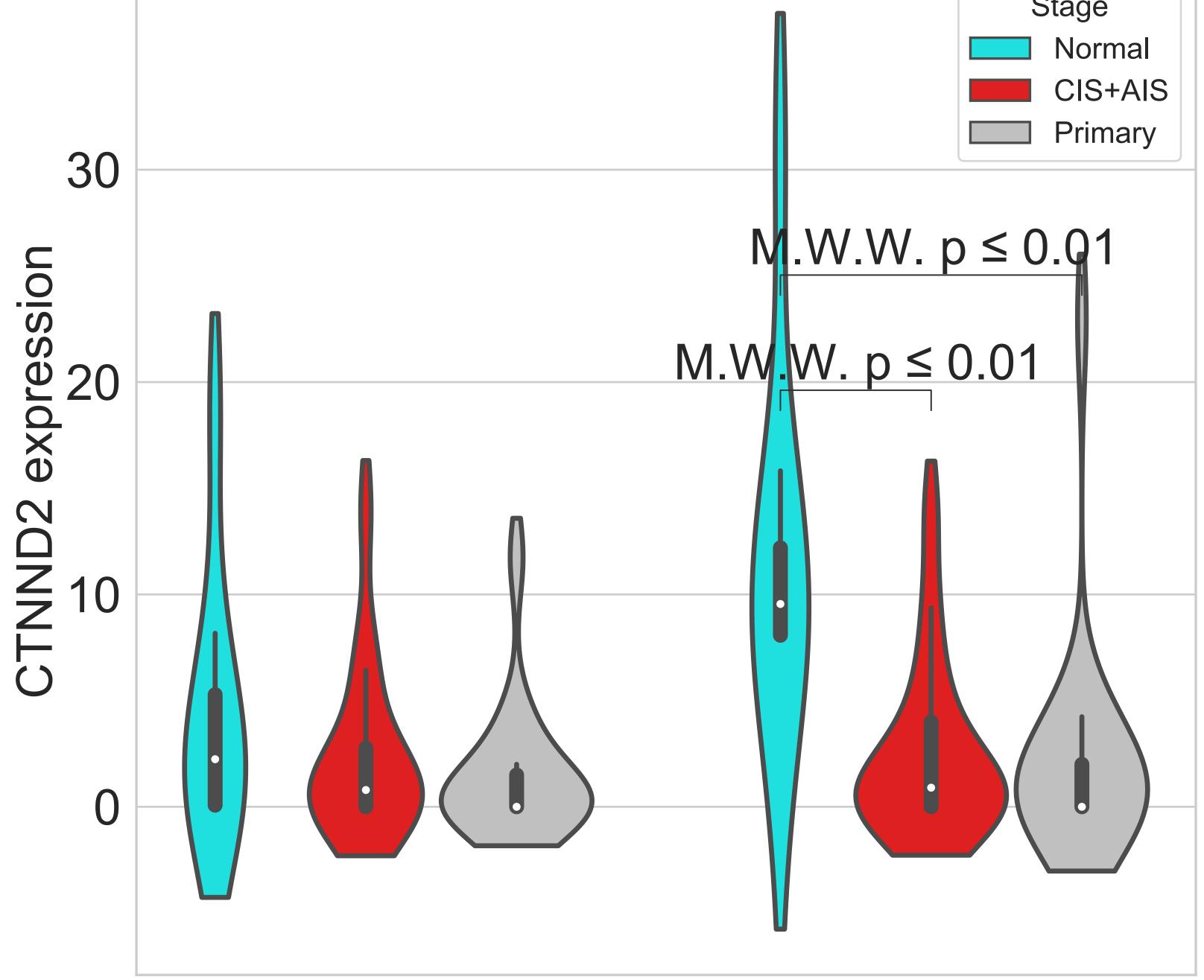
CTNND2: Kruskal-Wallis p=0.004 Stage Normal CIS+AIS Primary



Higher Lower Mutation Shared Count (SYN)