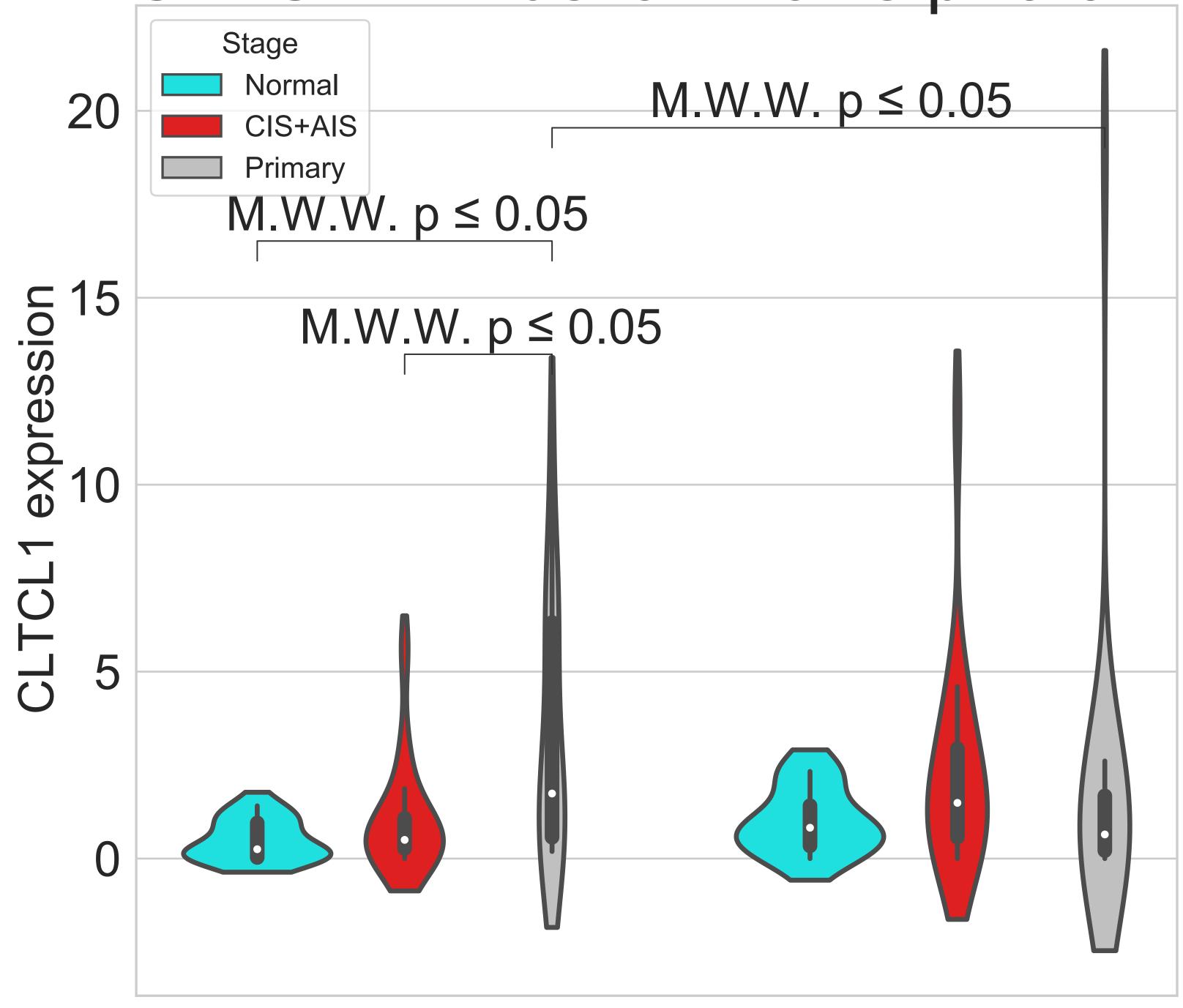
CLTCL1: Kruskal-Wallis p=0.041



Lower
Mutation Shared Count (SYN)