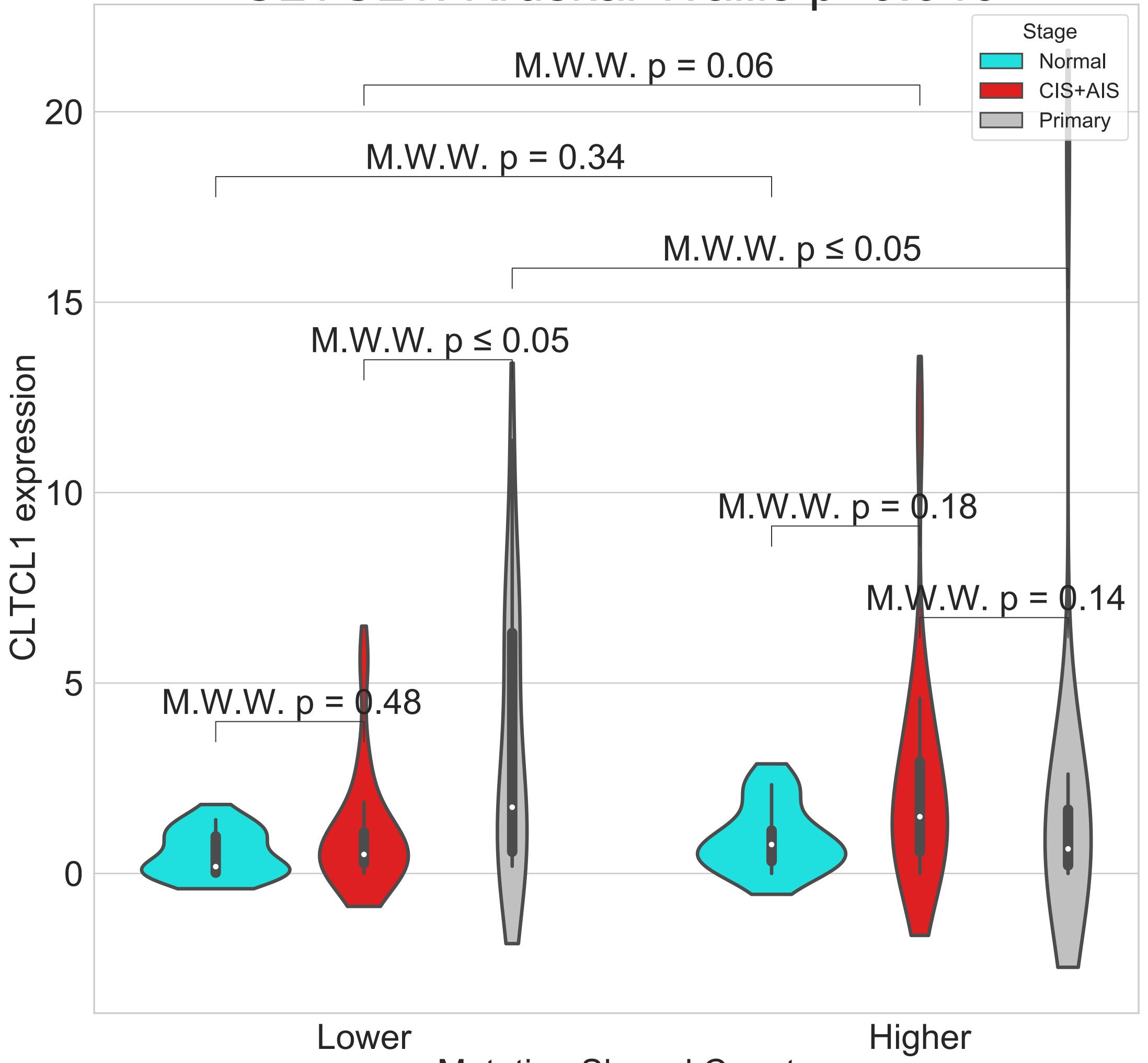
CLTCL1: Kruskal-Wallis p=0.046



Mutation Shared Count