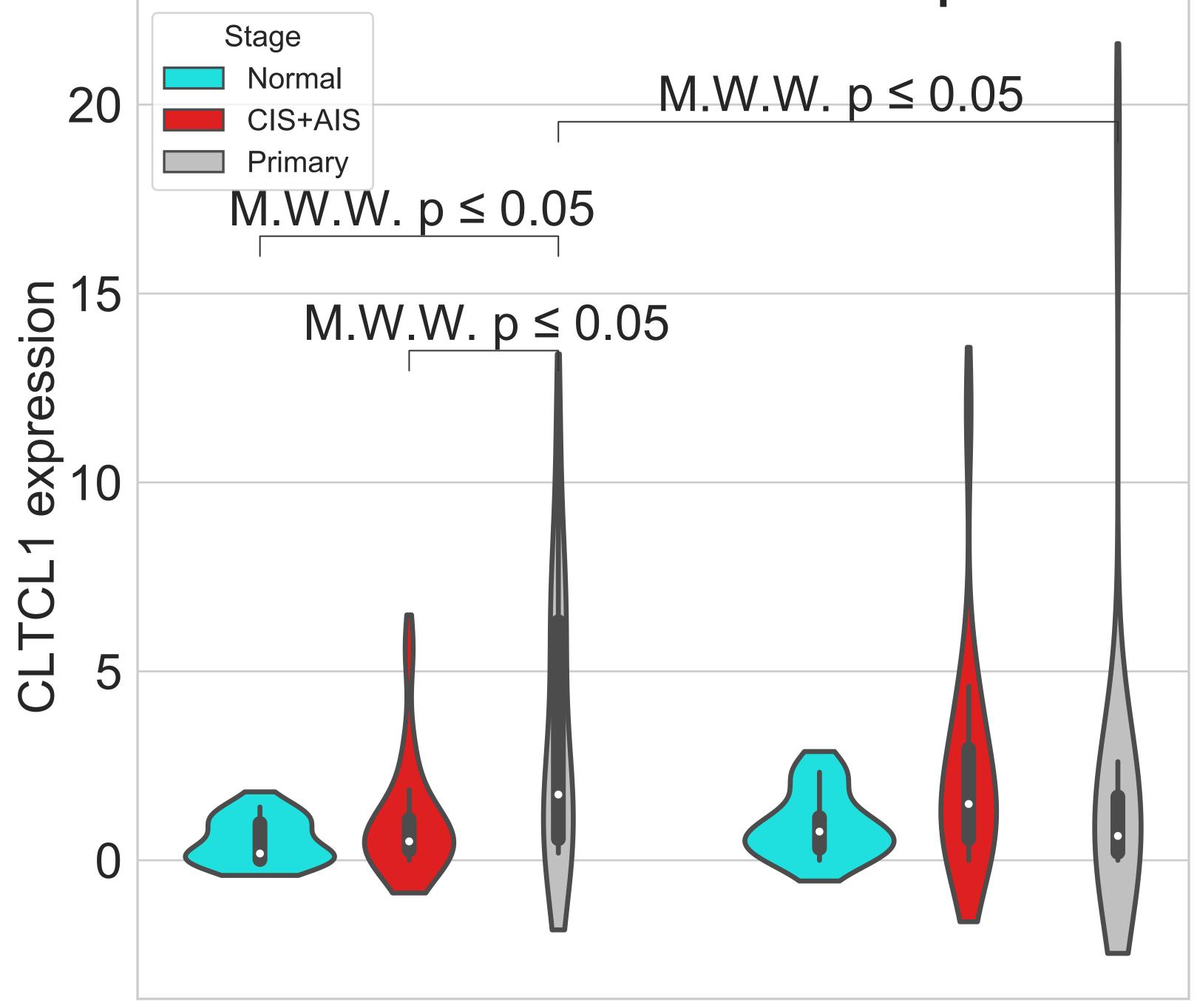
CLTCL1: Kruskal-Wallis p=0.046



Lower Higher Mutation Shared Count