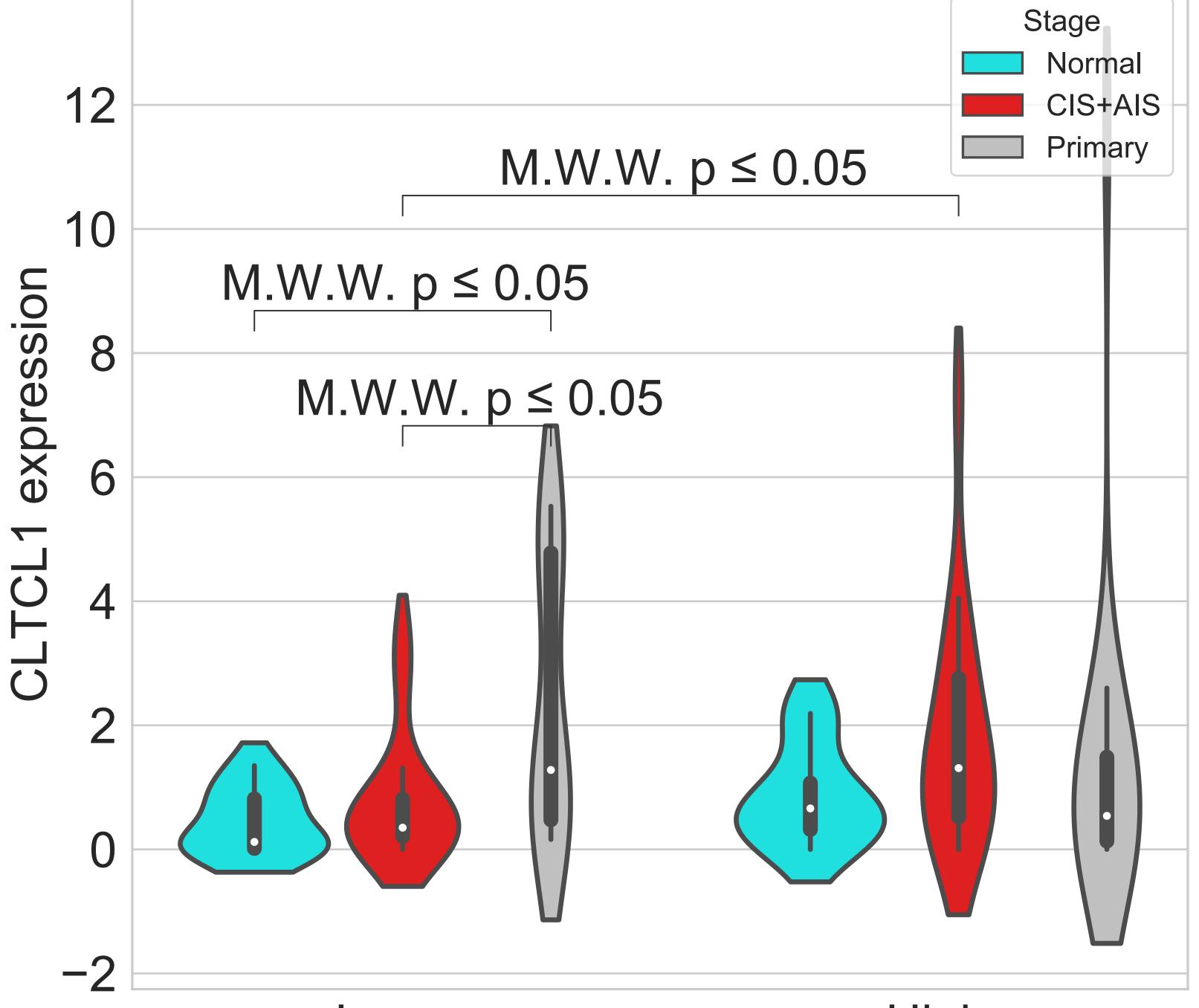
CLTCL1: Kruskal-Wallis p=0.038

Stage
Normal



Lower Higher Mutation Shared Count