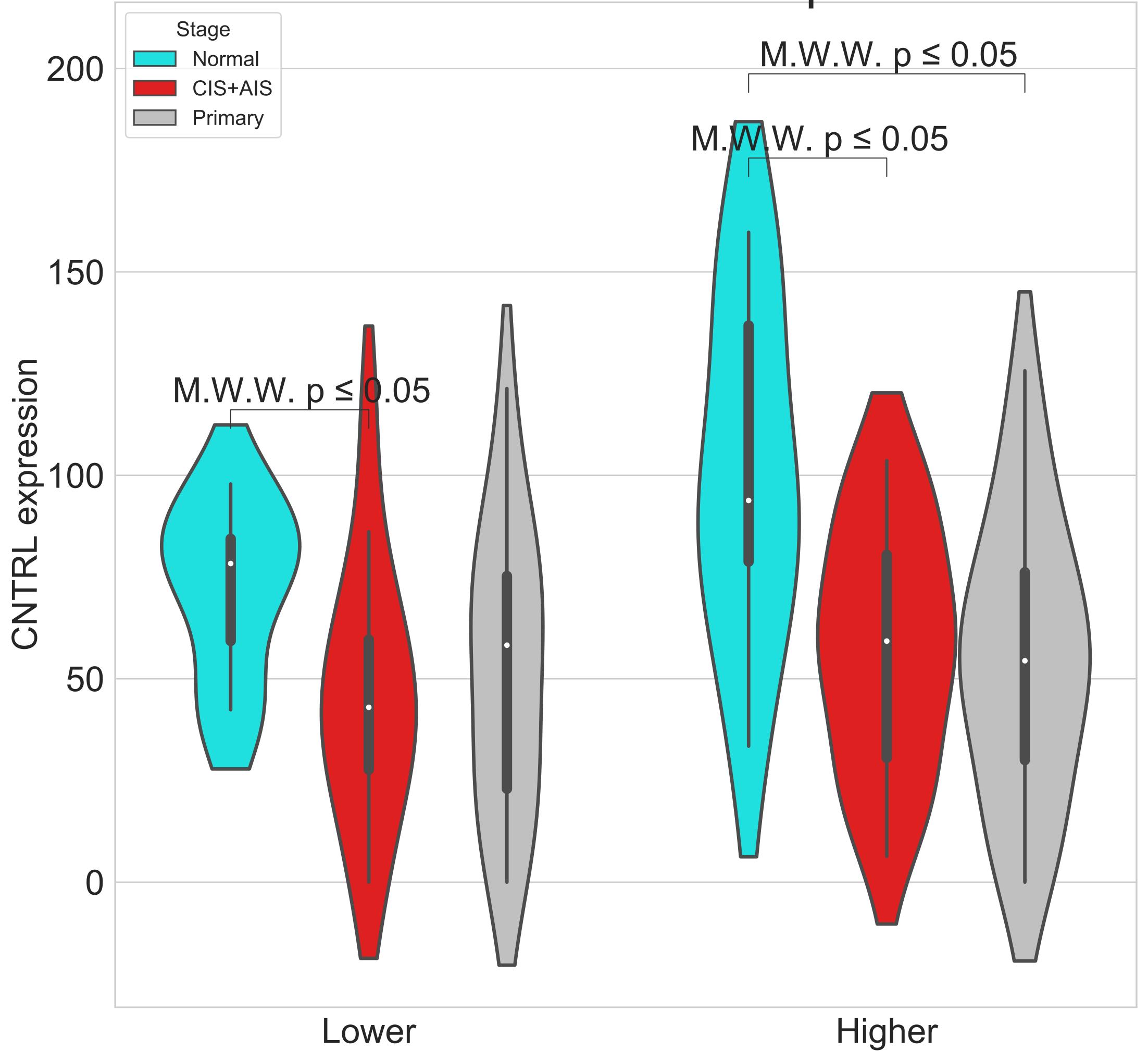
CNTRL: Kruskal-Wallis p=0.026



Mutation Shared Count per TMB