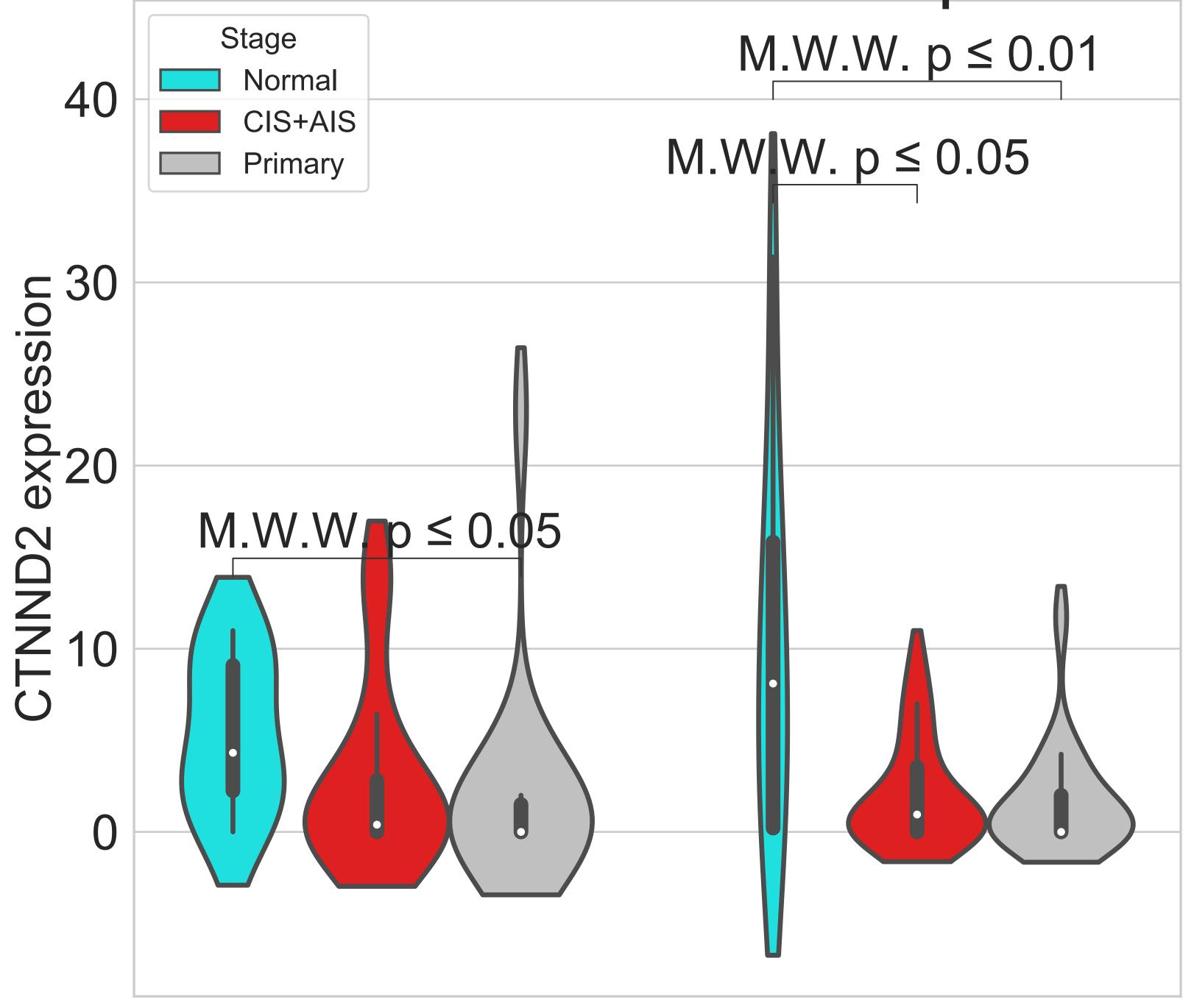
CTNND2: Kruskal-Wallis p=0.013



Lower
Mutation Shared Count per TMB