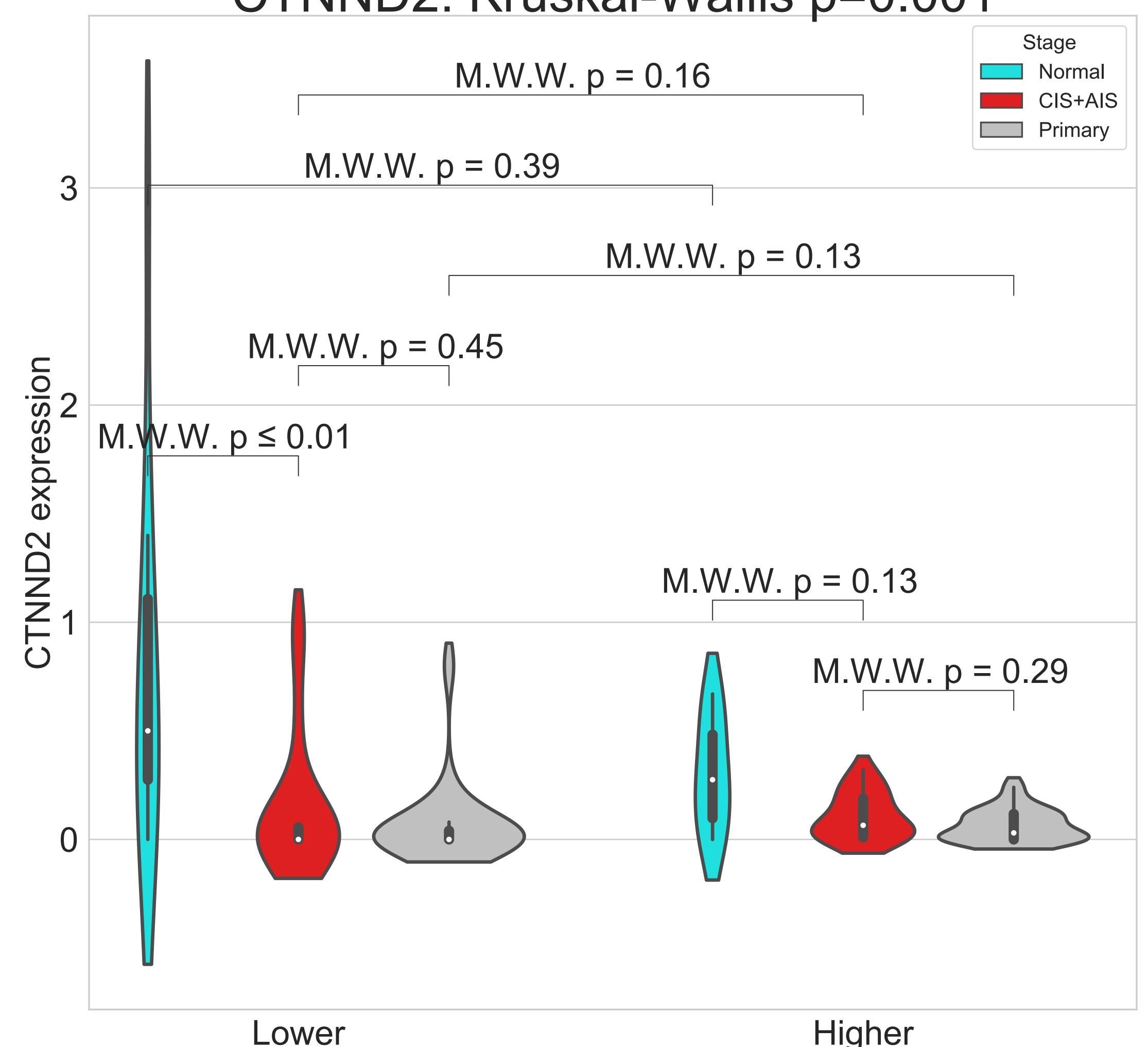
CTNND2: Kruskal-Wallis p=0.001



Higher Mutation Shared Count per TMB