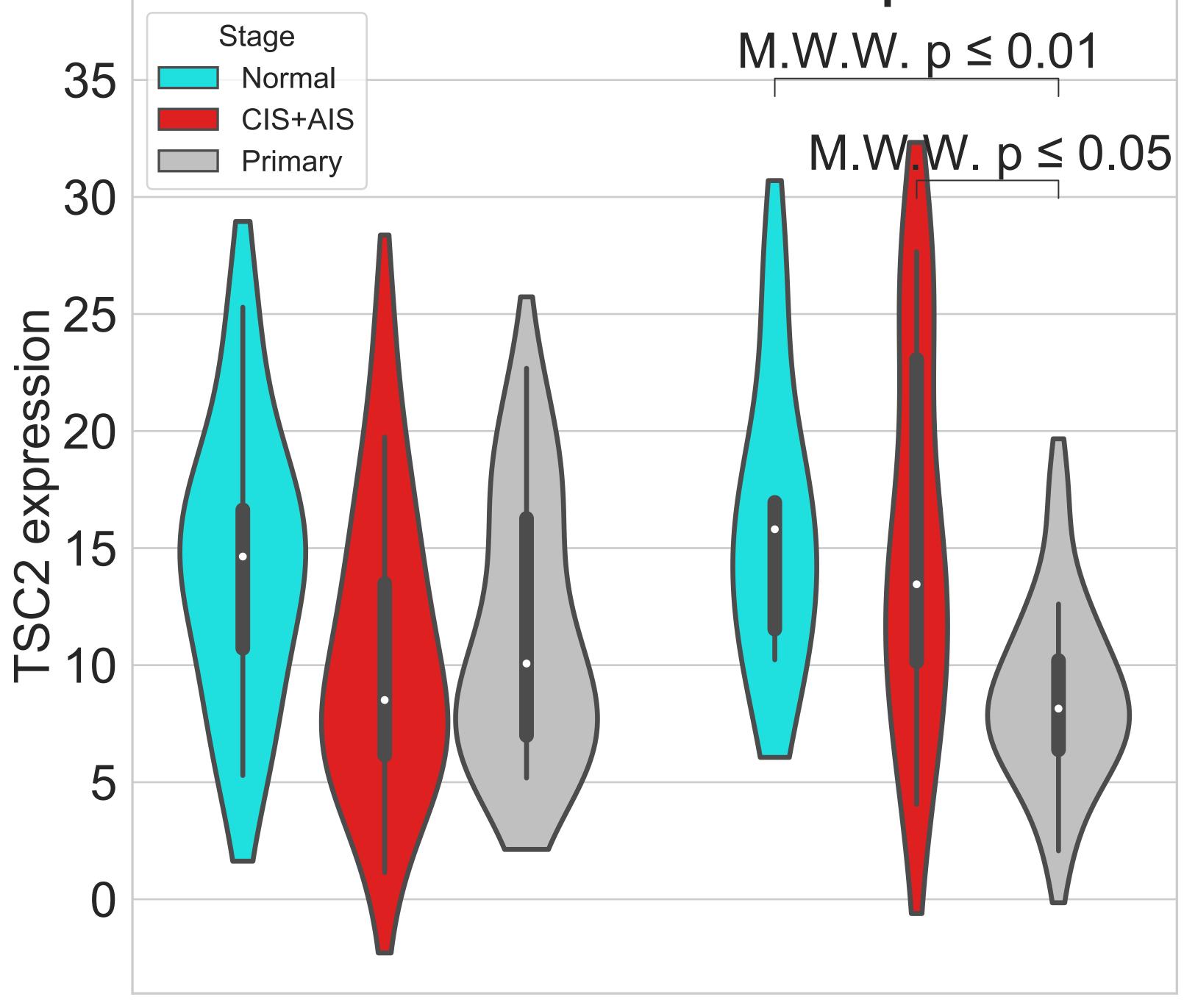
TSC2: Kruskal-Wallis p=0.035



Lower
Mutation Shared Count per TMB