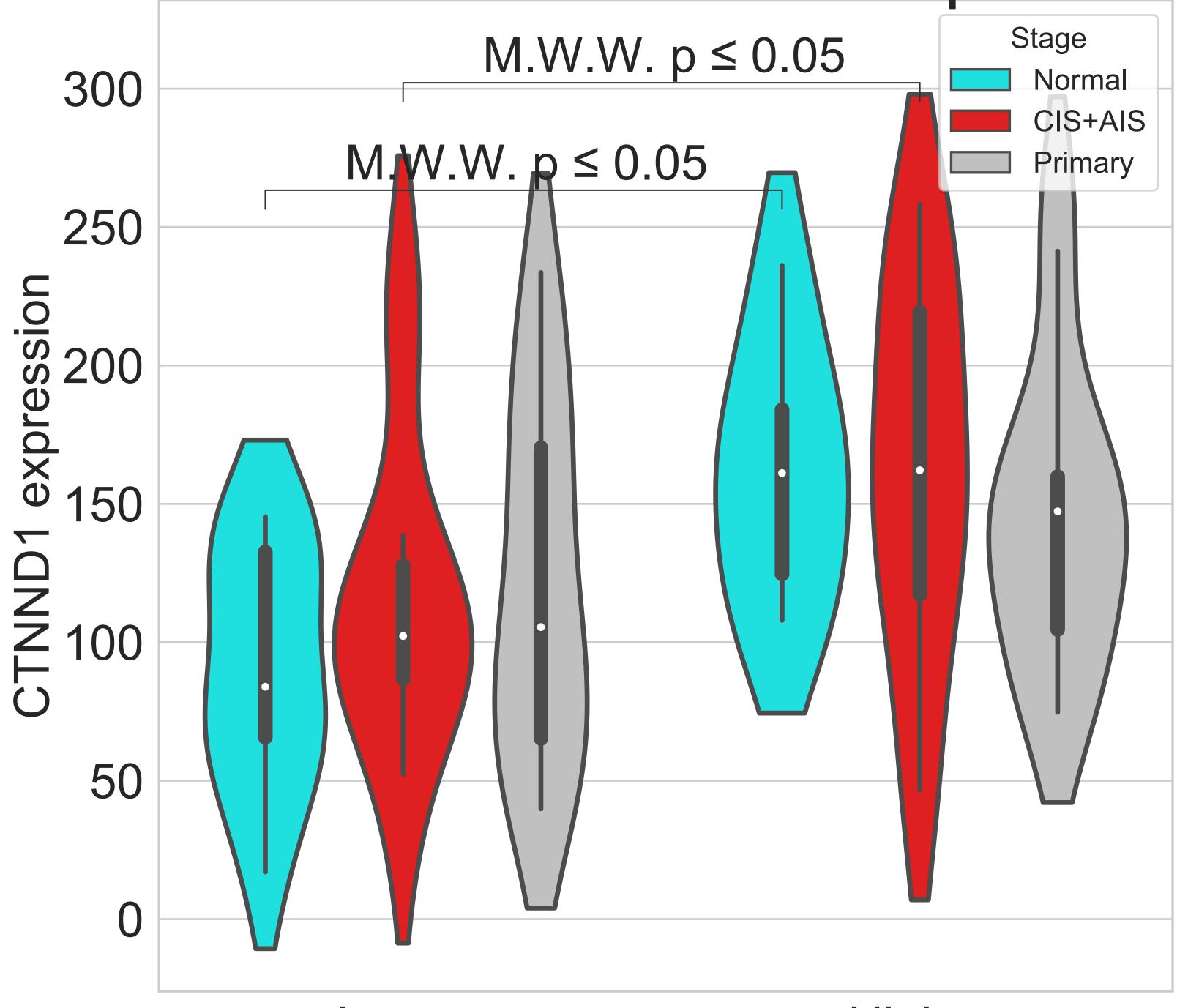
CTNND1: Kruskal-Wallis p=0.028



Lower
Mutation Shared Count per TMB