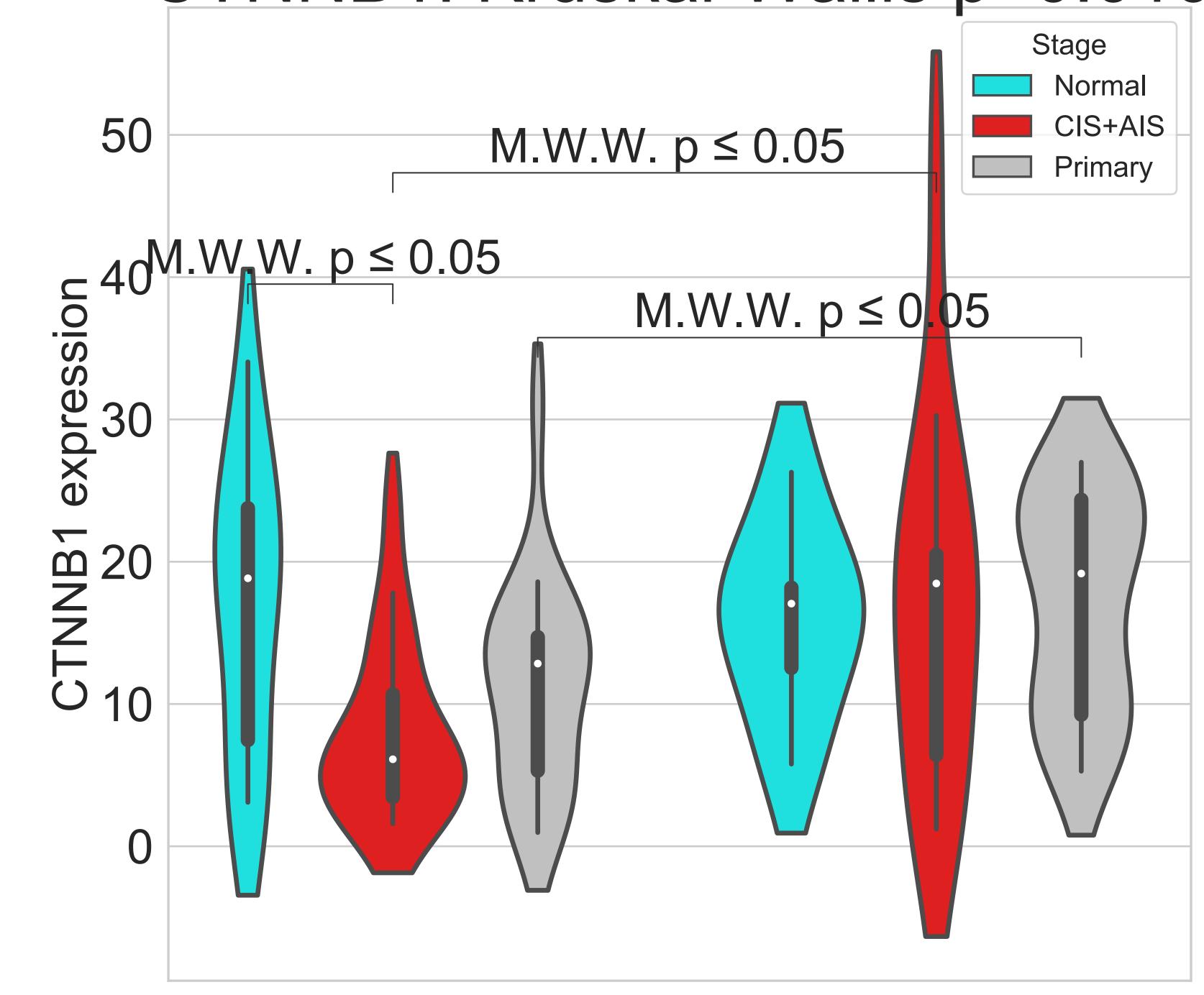
CTNNB1: Kruskal-Wallis p=0.010



Lower
Mutation Shared Count per TMB