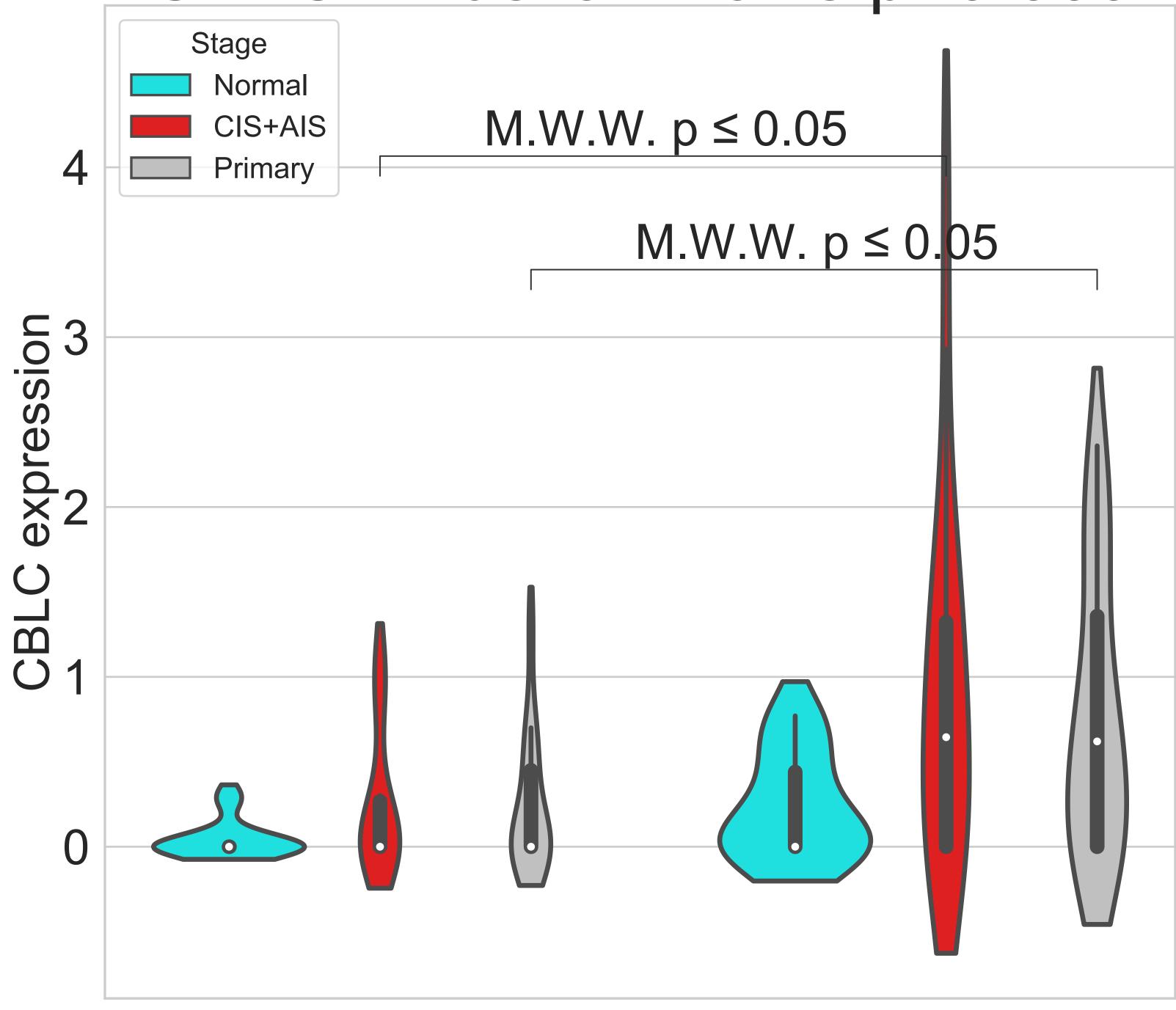
CBLC: Kruskal-Wallis p=0.009



Lower
Mutation Shared Count per TMB