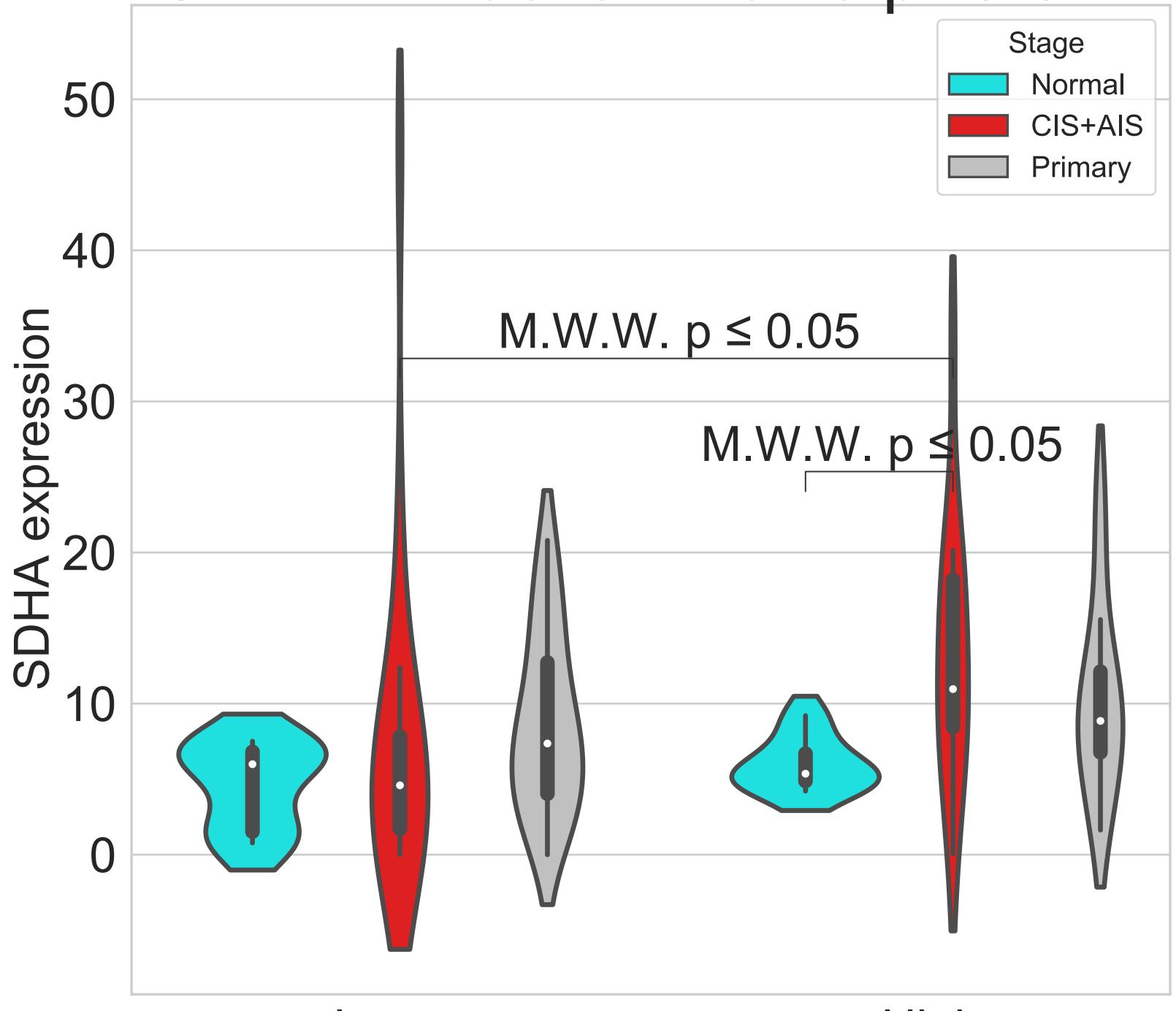
SDHA: Kruskal-Wallis p=0.012



Lower
Mutation Shared Count per TMB