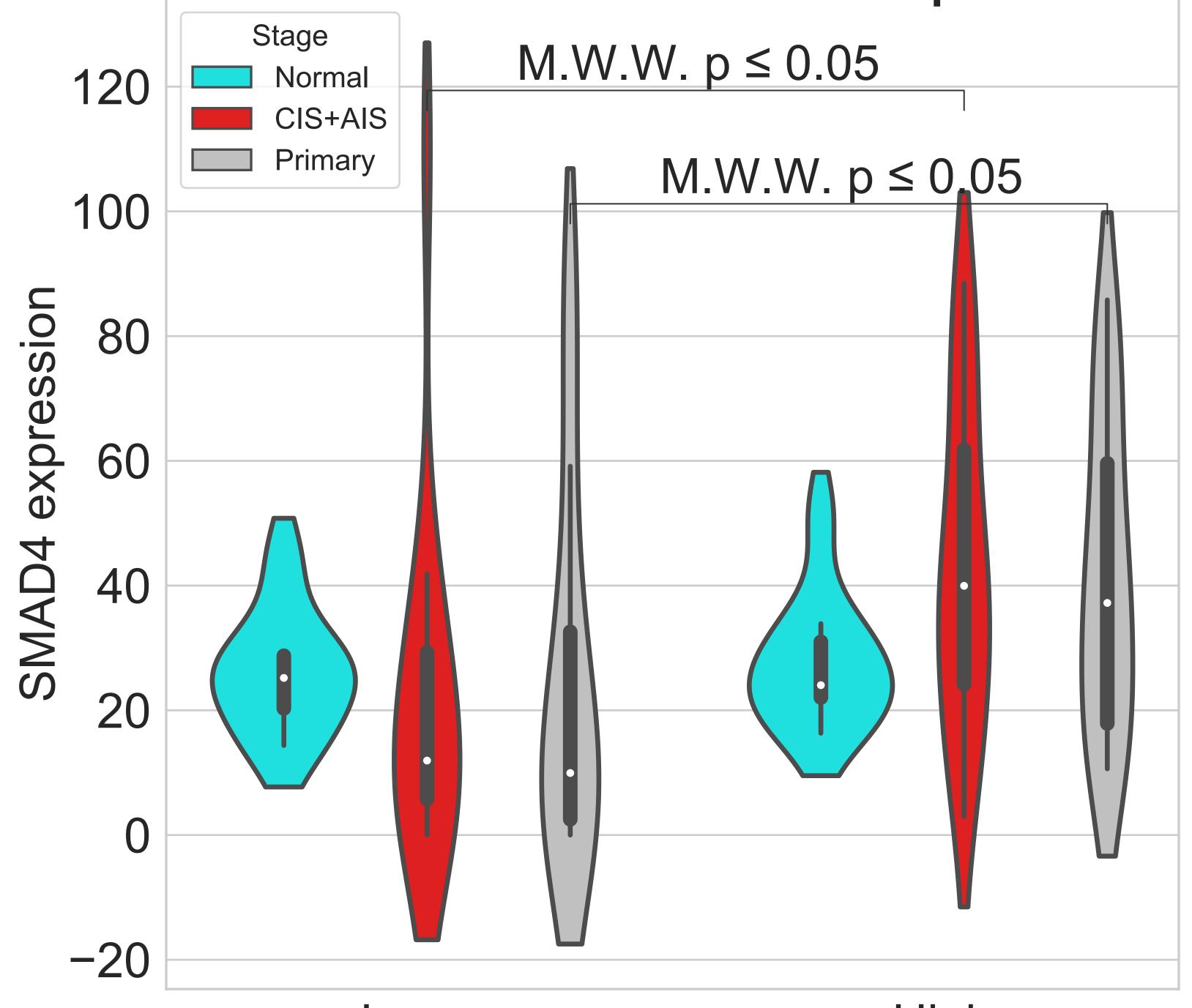
SMAD4: Kruskal-Wallis p=0.015



Lower
Mutation Shared Count per TMB