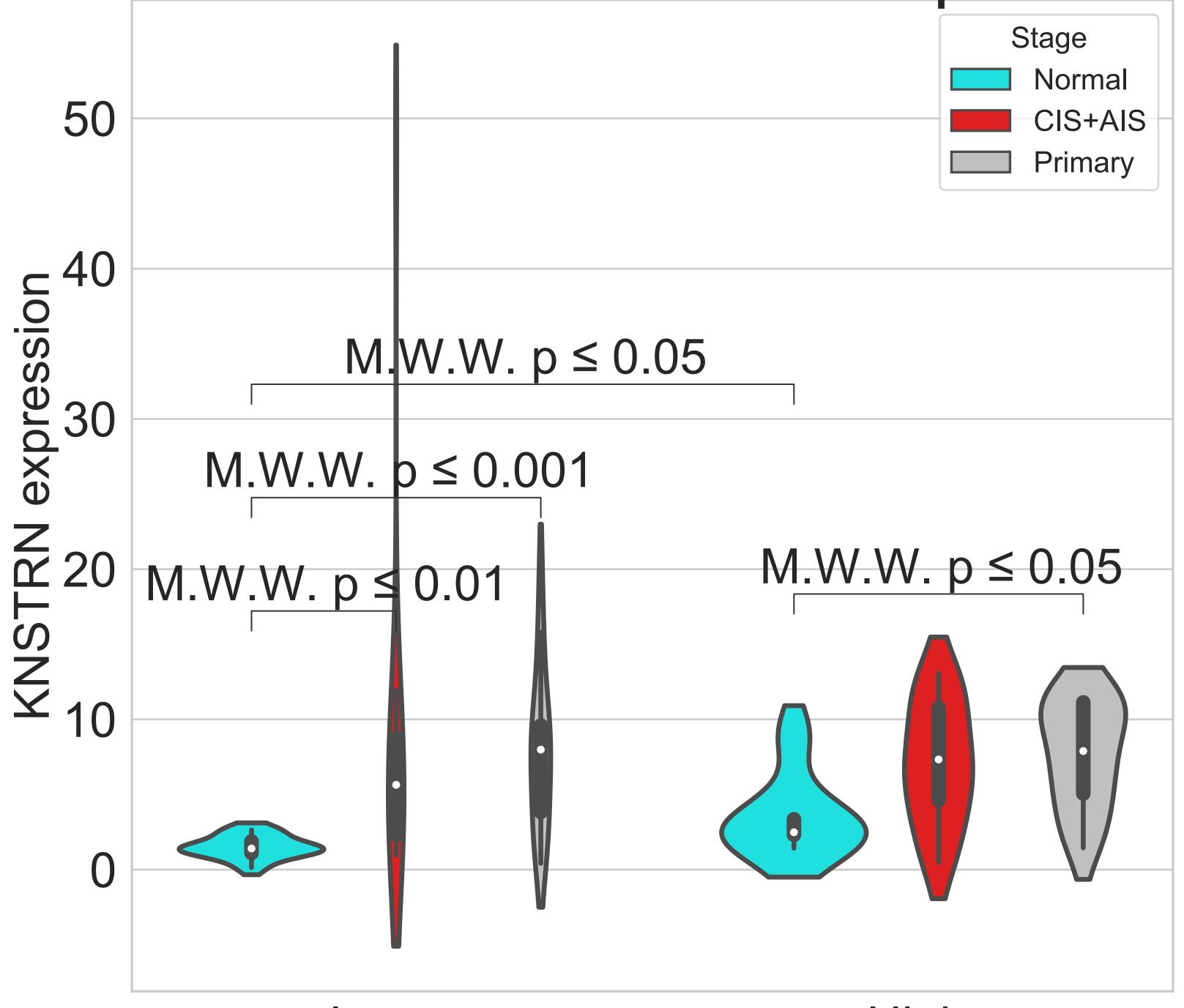
KNSTRN: Kruskal-Wallis p=0.001



Lower
Mutation Shared Count per TMB