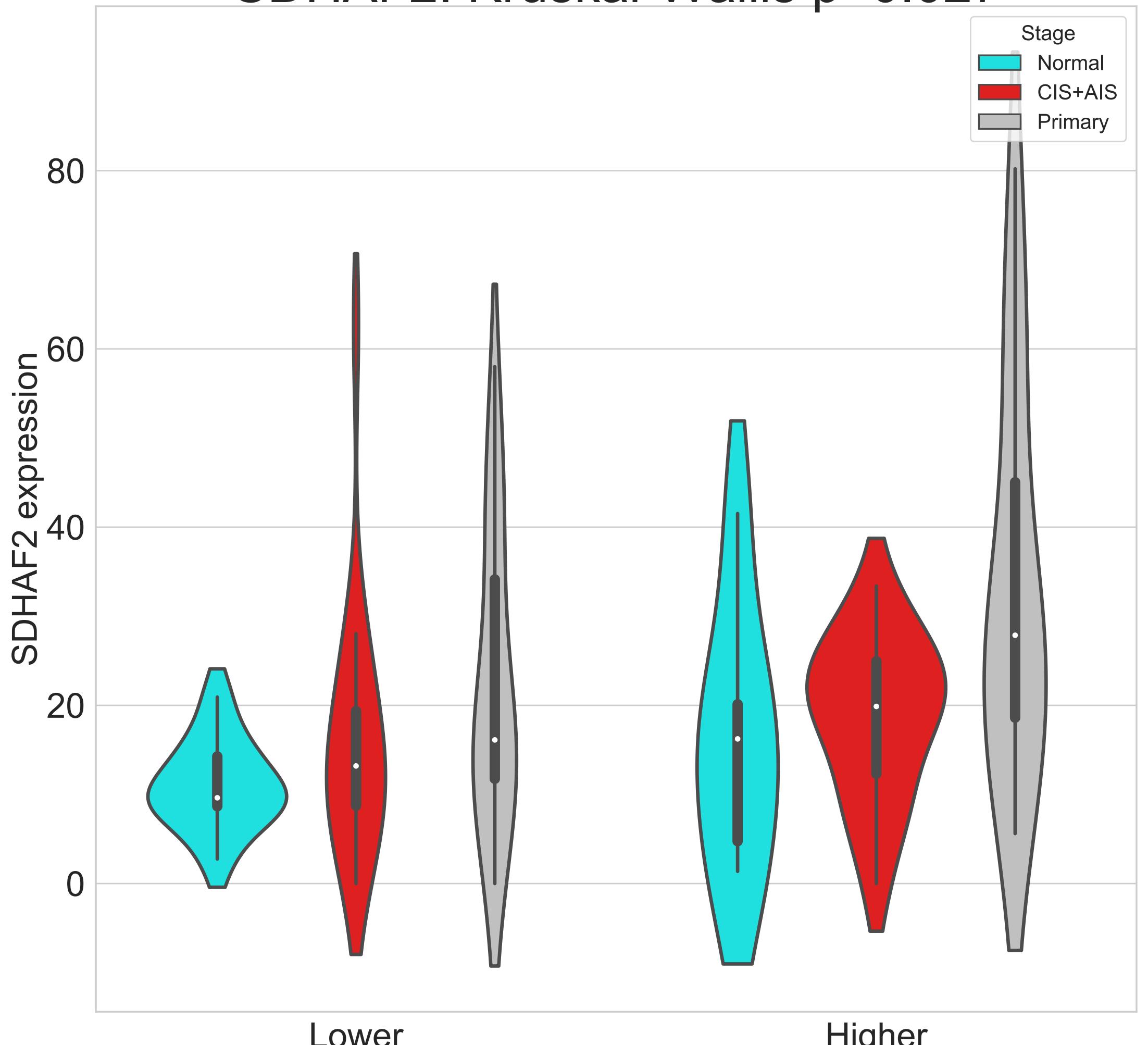
SDHAF2: Kruskal-Wallis p=0.027



Lower
Mutation Shared Count per TMB