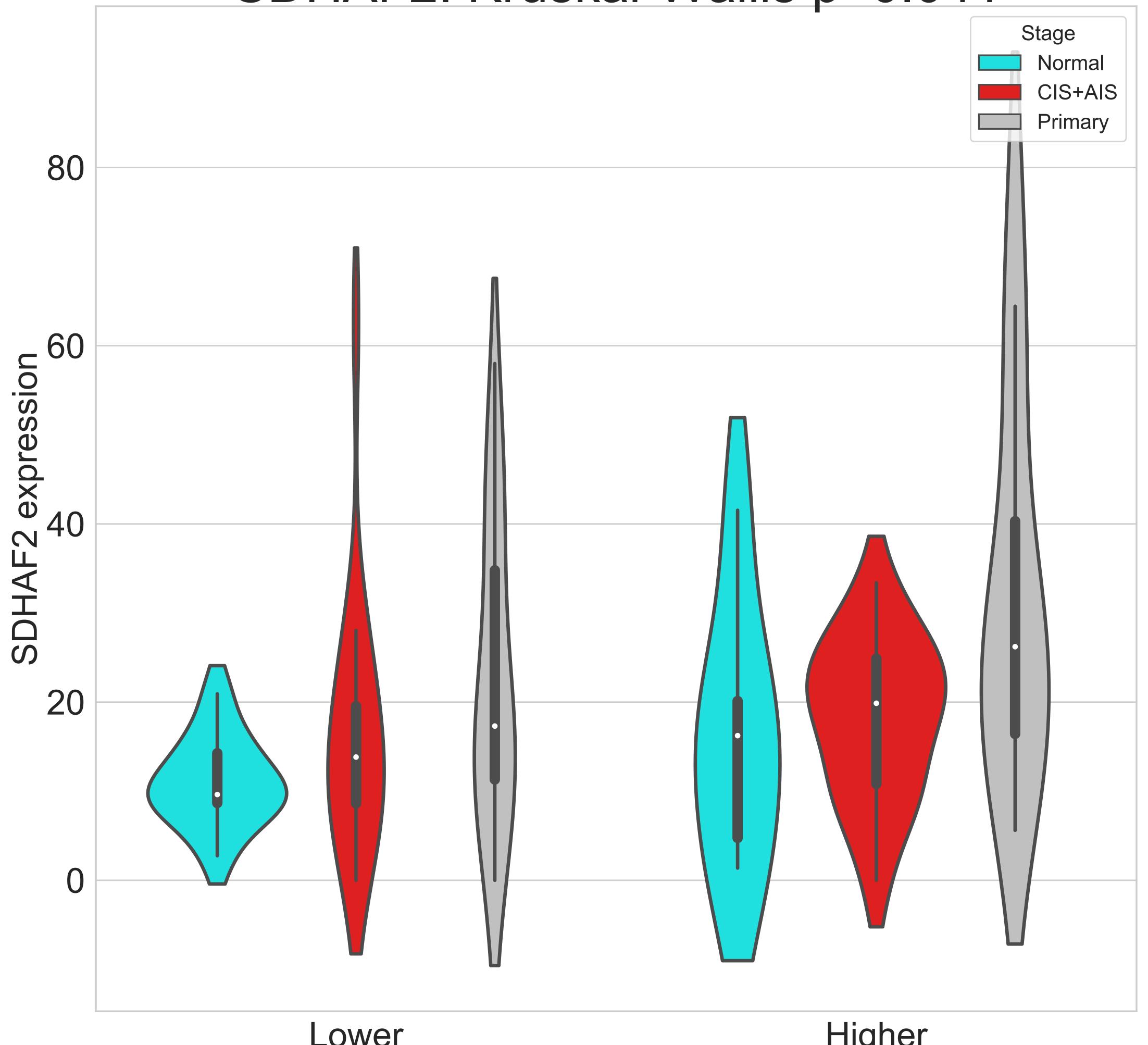
SDHAF2: Kruskal-Wallis p=0.041



Lower
Mutation Shared Count (SYN) per TMB