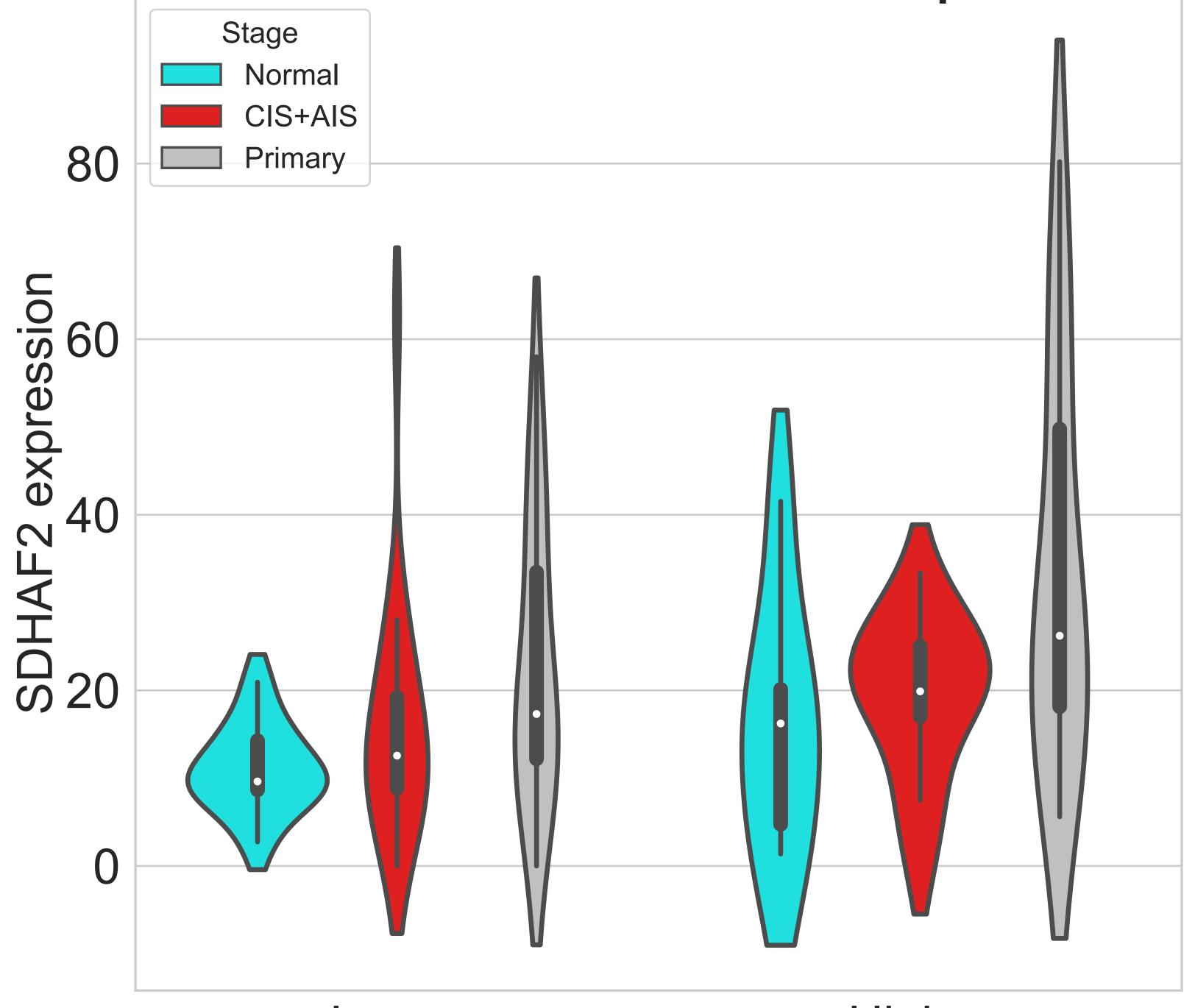
SDHAF2: Kruskal-Wallis p=0.030



Lower Higher Mutation Shared Proportion (Union)