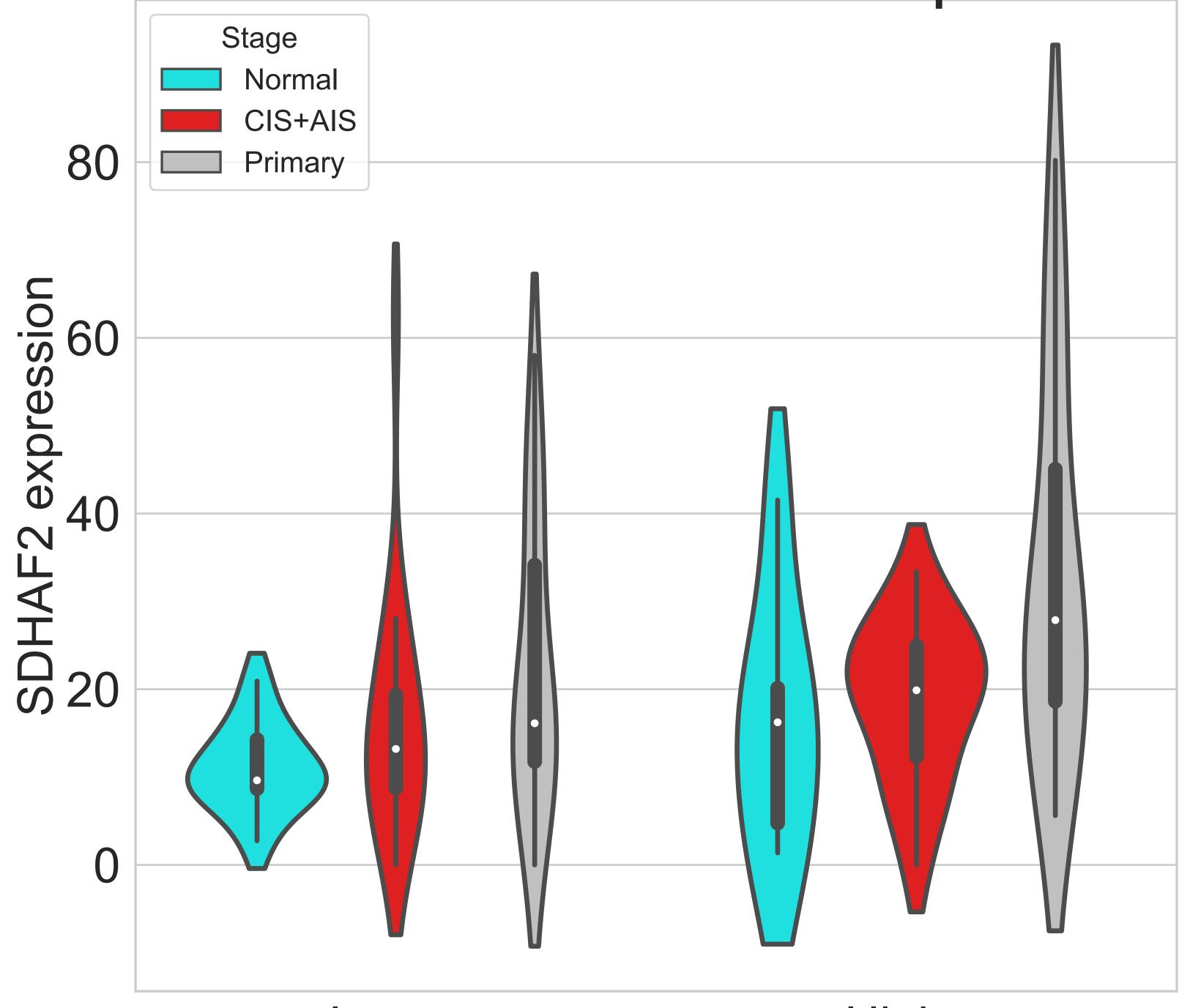
SDHAF2: Kruskal-Wallis p=0.027



Lower Higher Mutation Shared Proportion