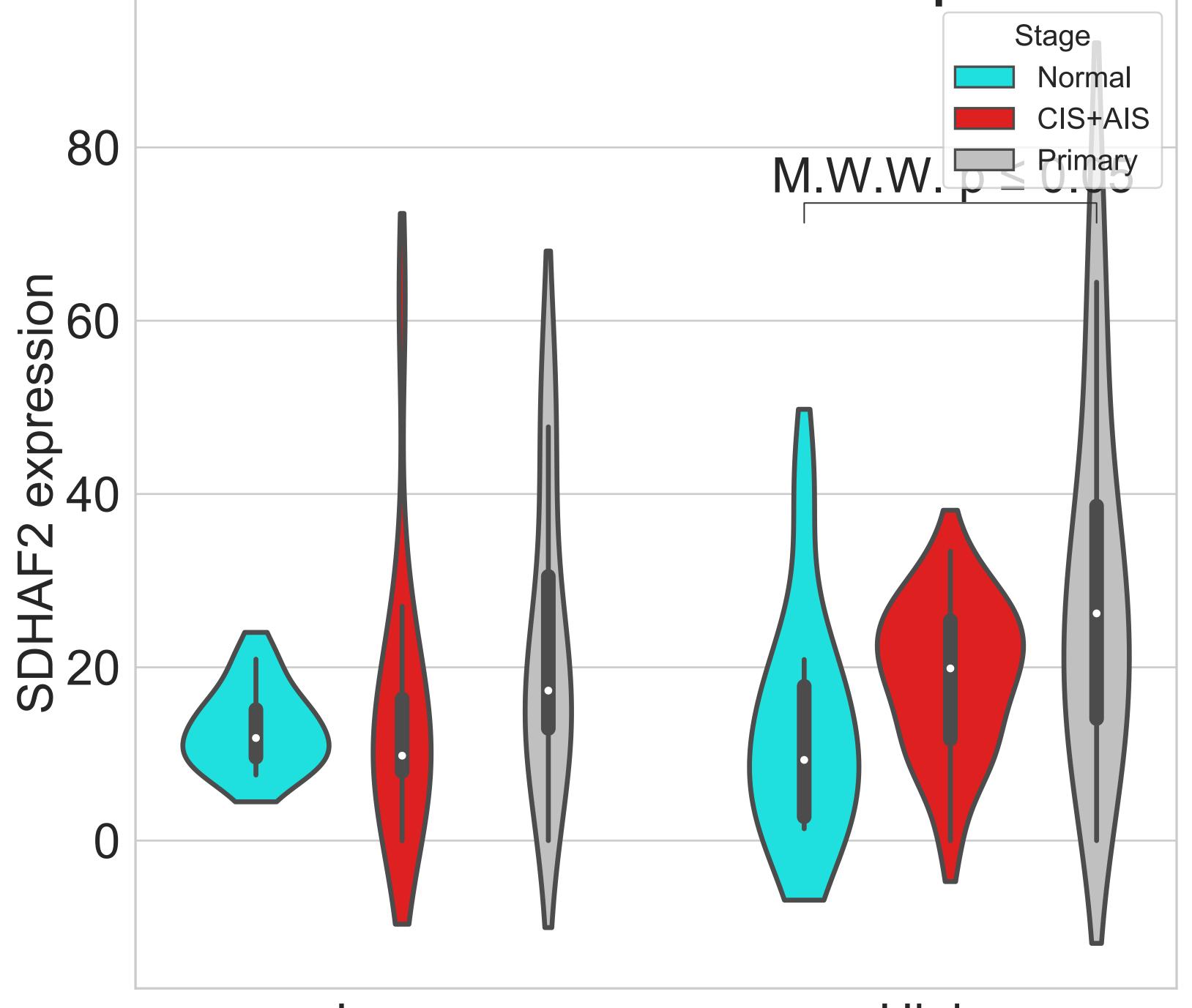
SDHAF2: Kruskal-Wallis p=0.032



Lower Higher Mutation Shared Proportion (SYN)