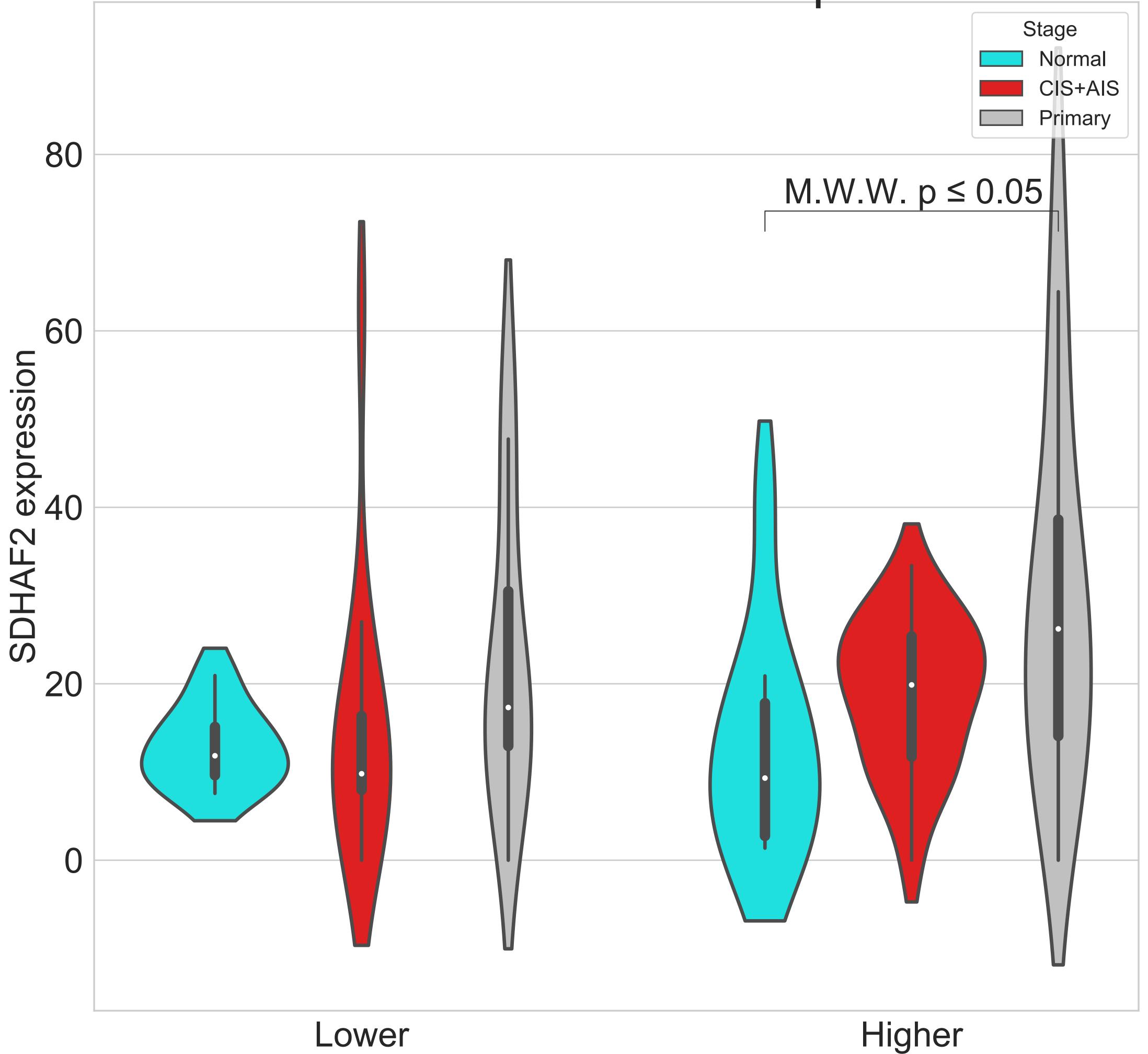
SDHAF2: Kruskal-Wallis p=0.032



Mutation Shared Proportion (SYN)