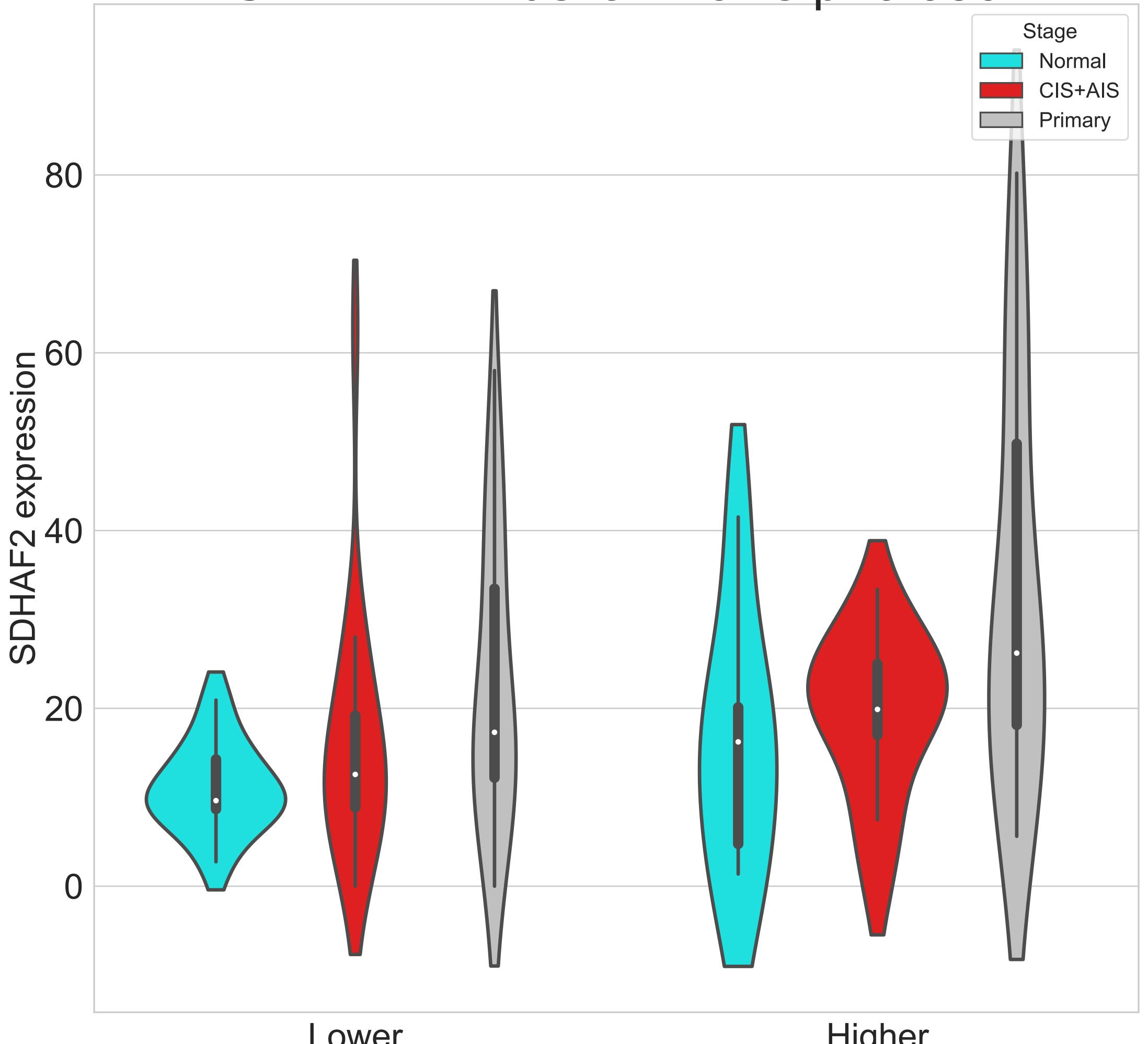
SDHAF2: Kruskal-Wallis p=0.030



Lower
Mutation Shared Proportion (Union)