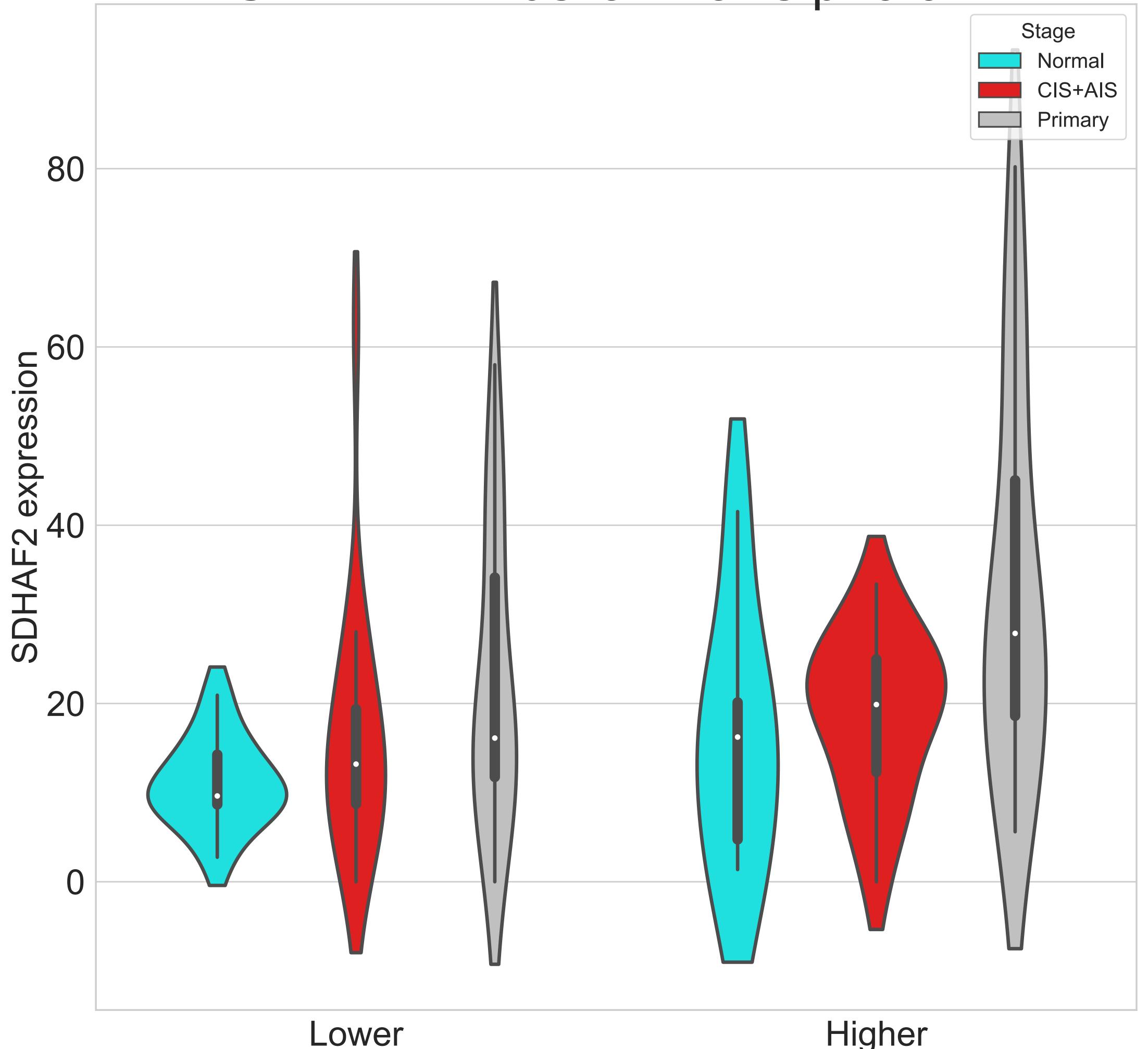
SDHAF2: Kruskal-Wallis p=0.027



ver
Mutation Shared Proportion