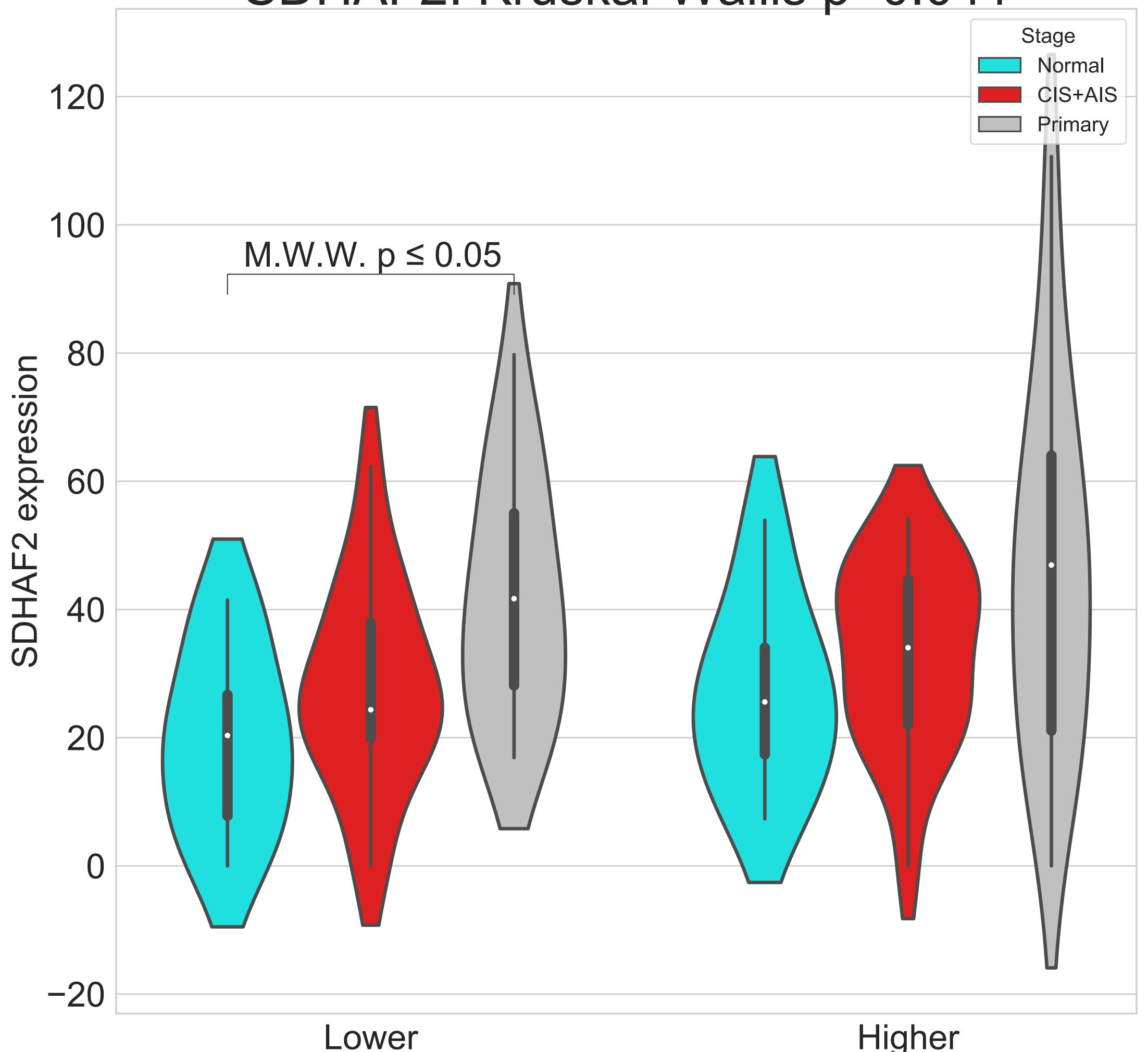
SDHAF2: Kruskal-Wallis p=0.041



ower Mutation Shared Count (SYN)