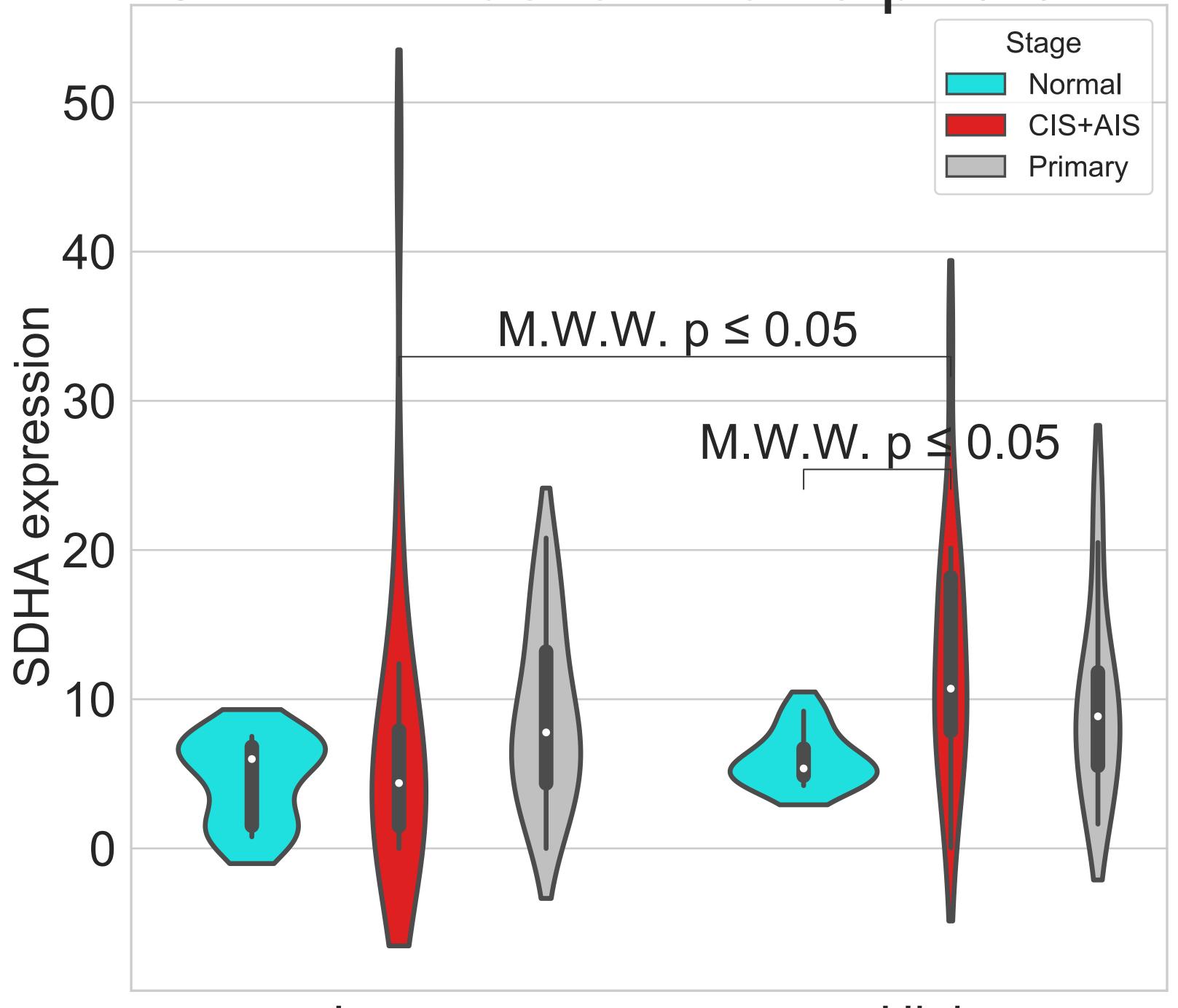
SDHA: Kruskal-Wallis p=0.014



Lower Higher Mutation Shared Proportion (SYN)