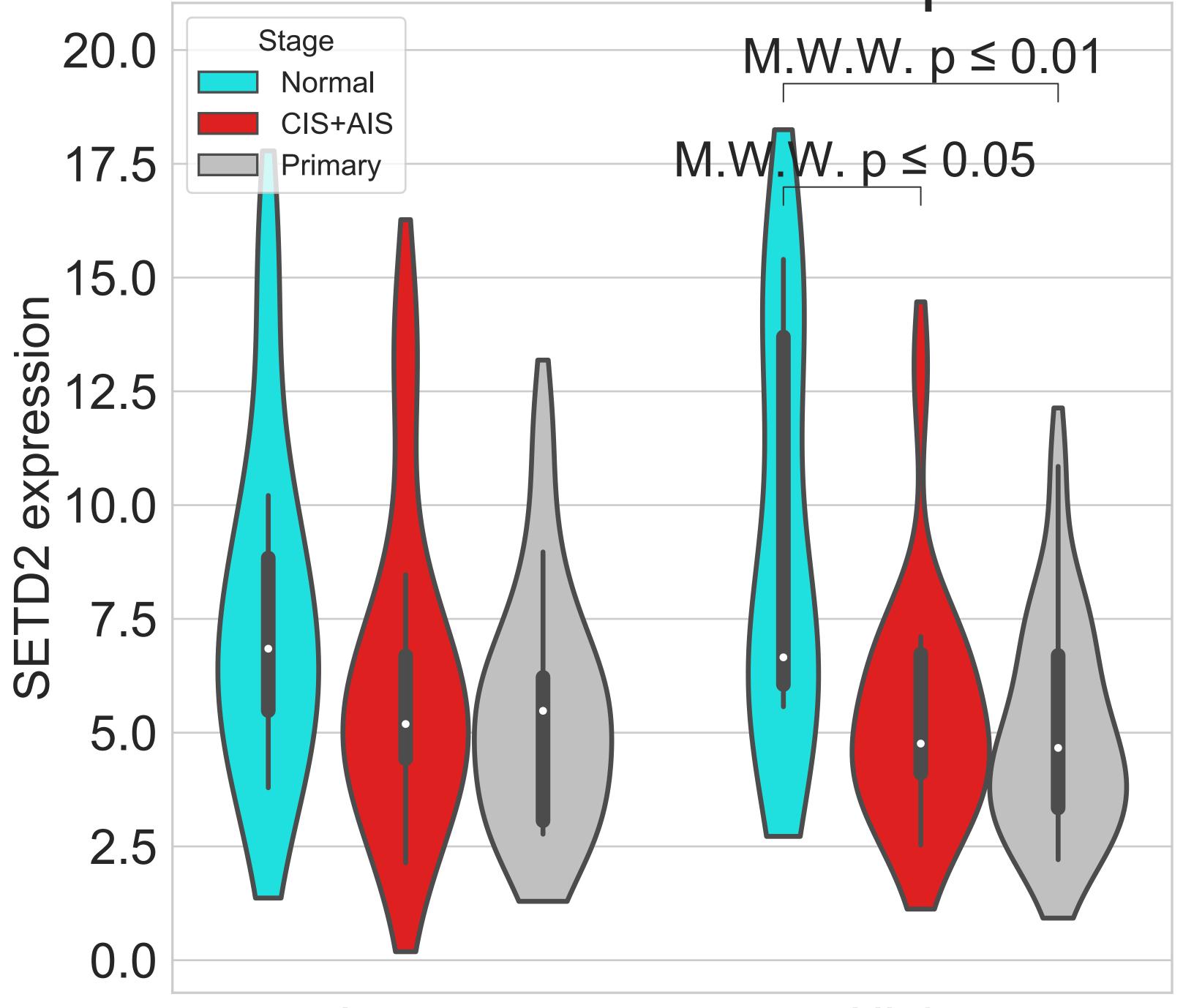
SETD2: Kruskal-Wallis p=0.037



Lower Higher Mutation Shared Count (SYN)