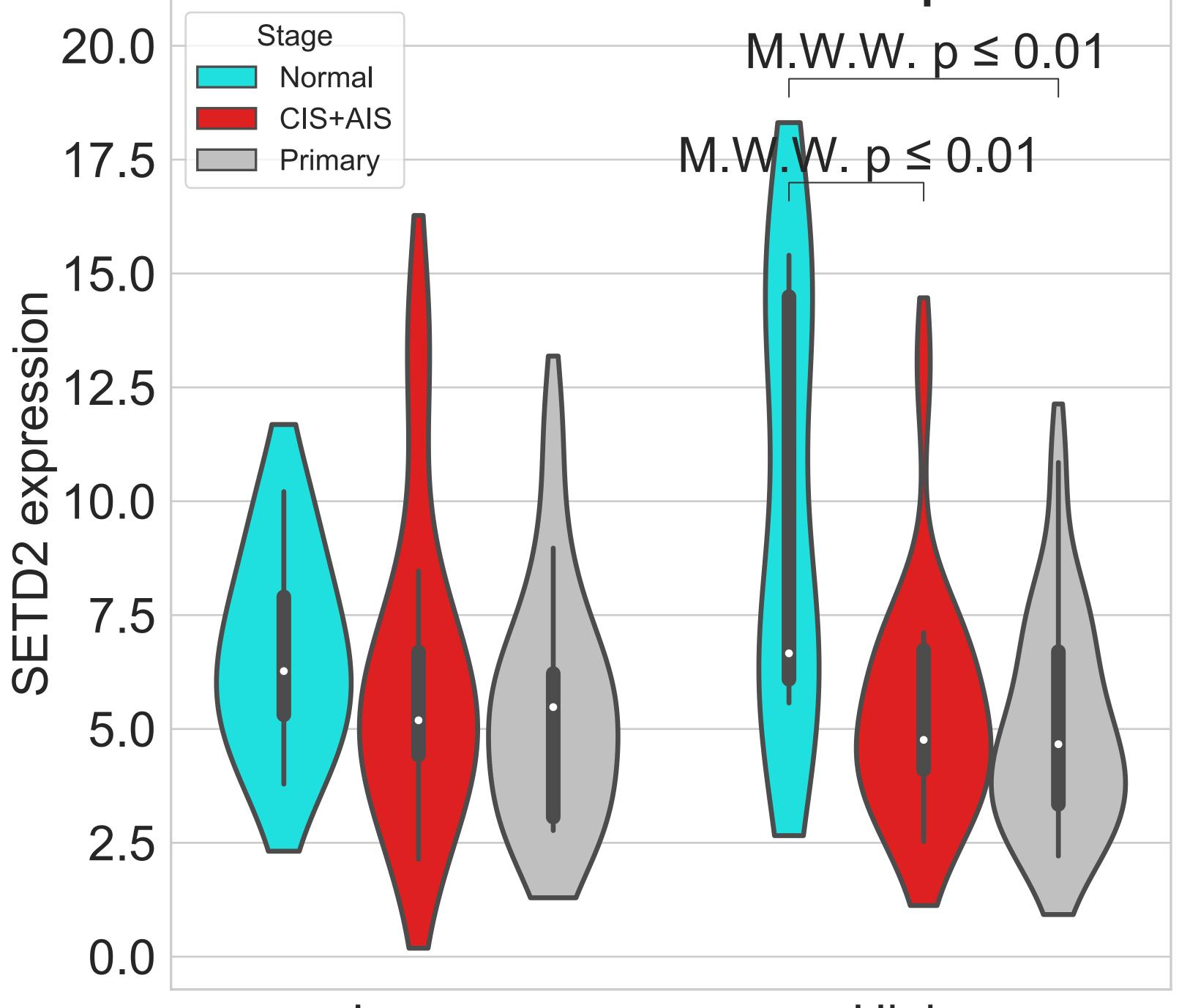
SETD2: Kruskal-Wallis p=0.027



Lower Higher Mutation Shared Count