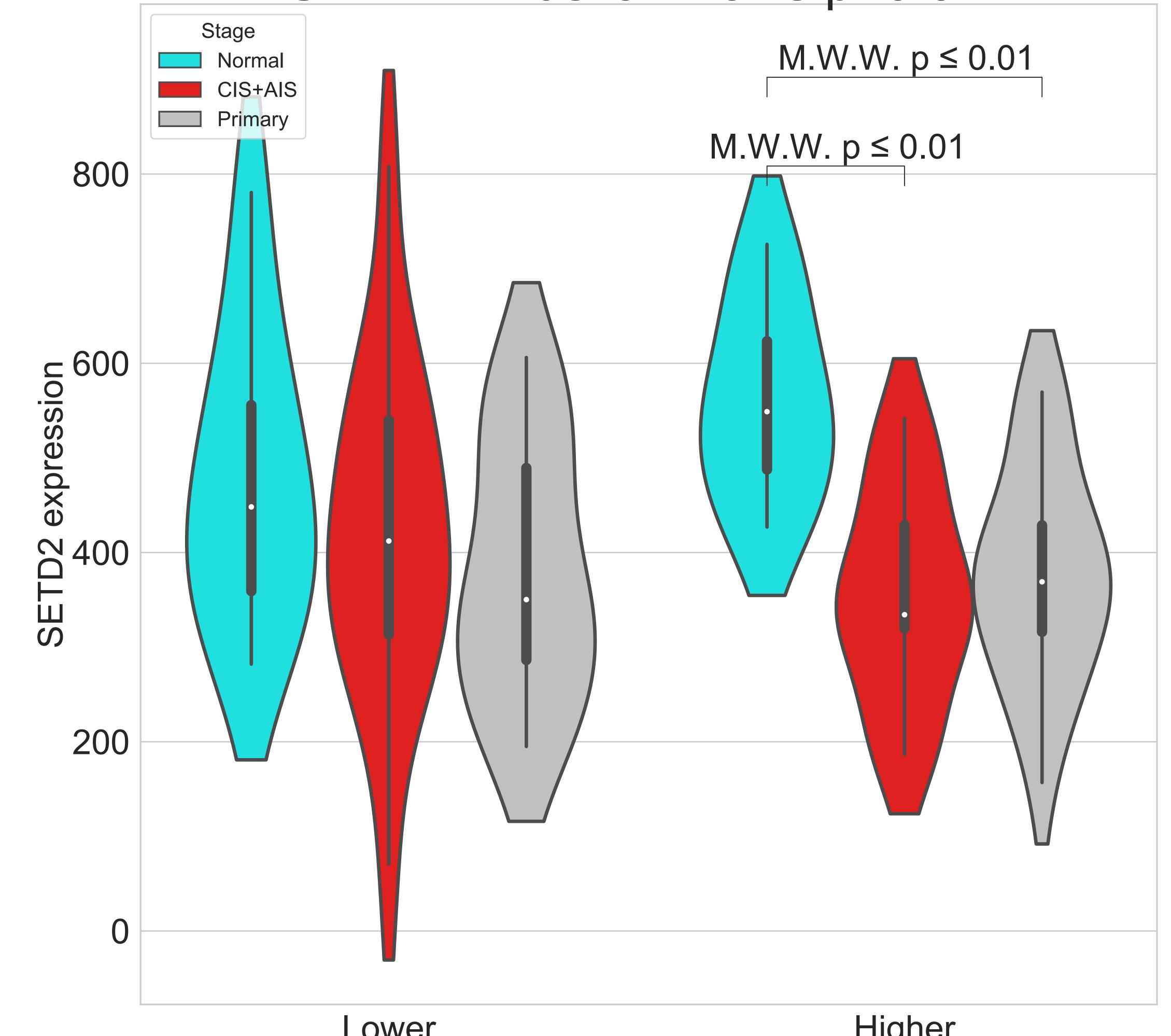
SETD2: Kruskal-Wallis p=0.024



Lower
Mutation Shared Count (SYN)