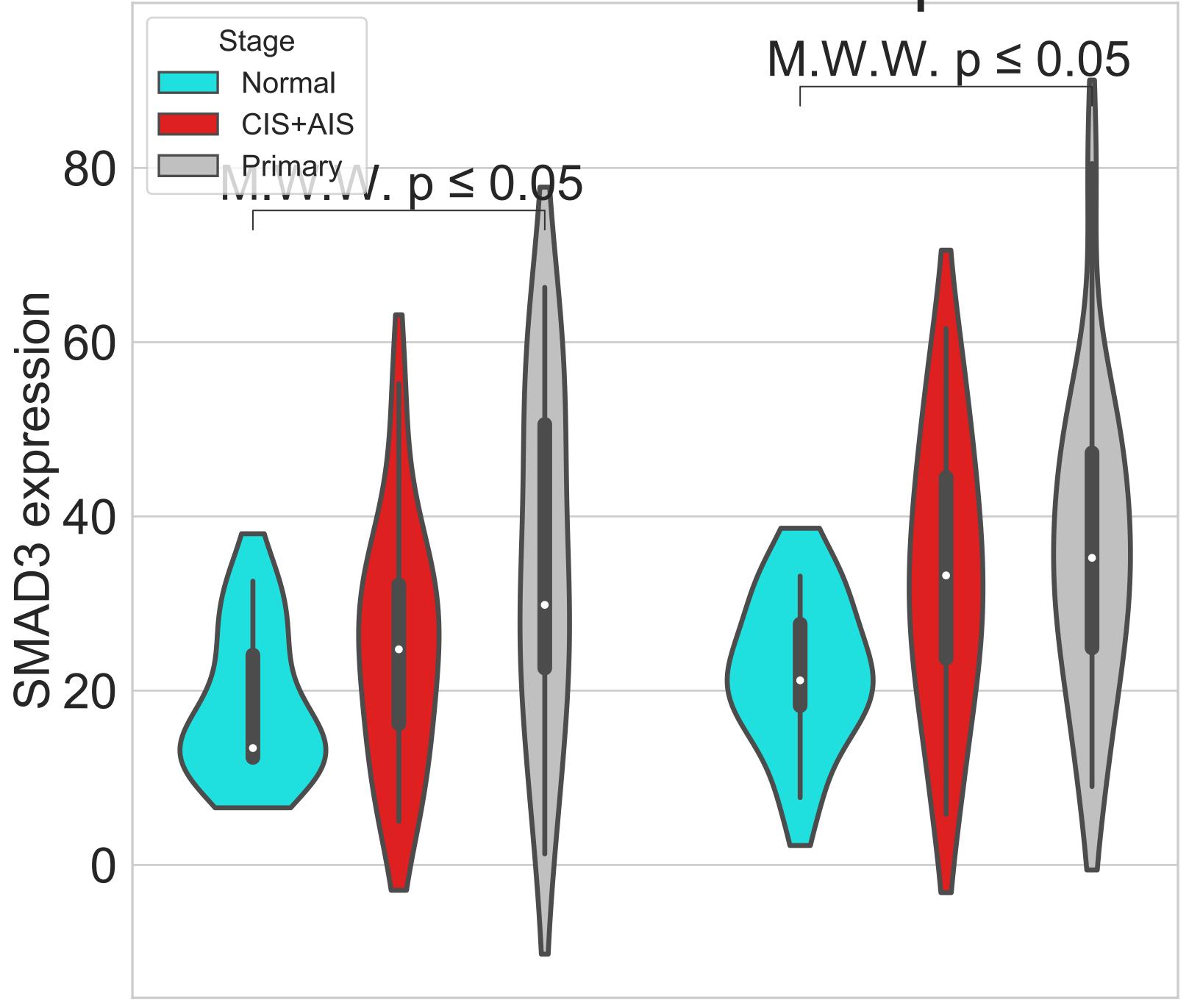
SMAD3: Kruskal-Wallis p=0.029



Lower
Mutation Shared Count (SYN)