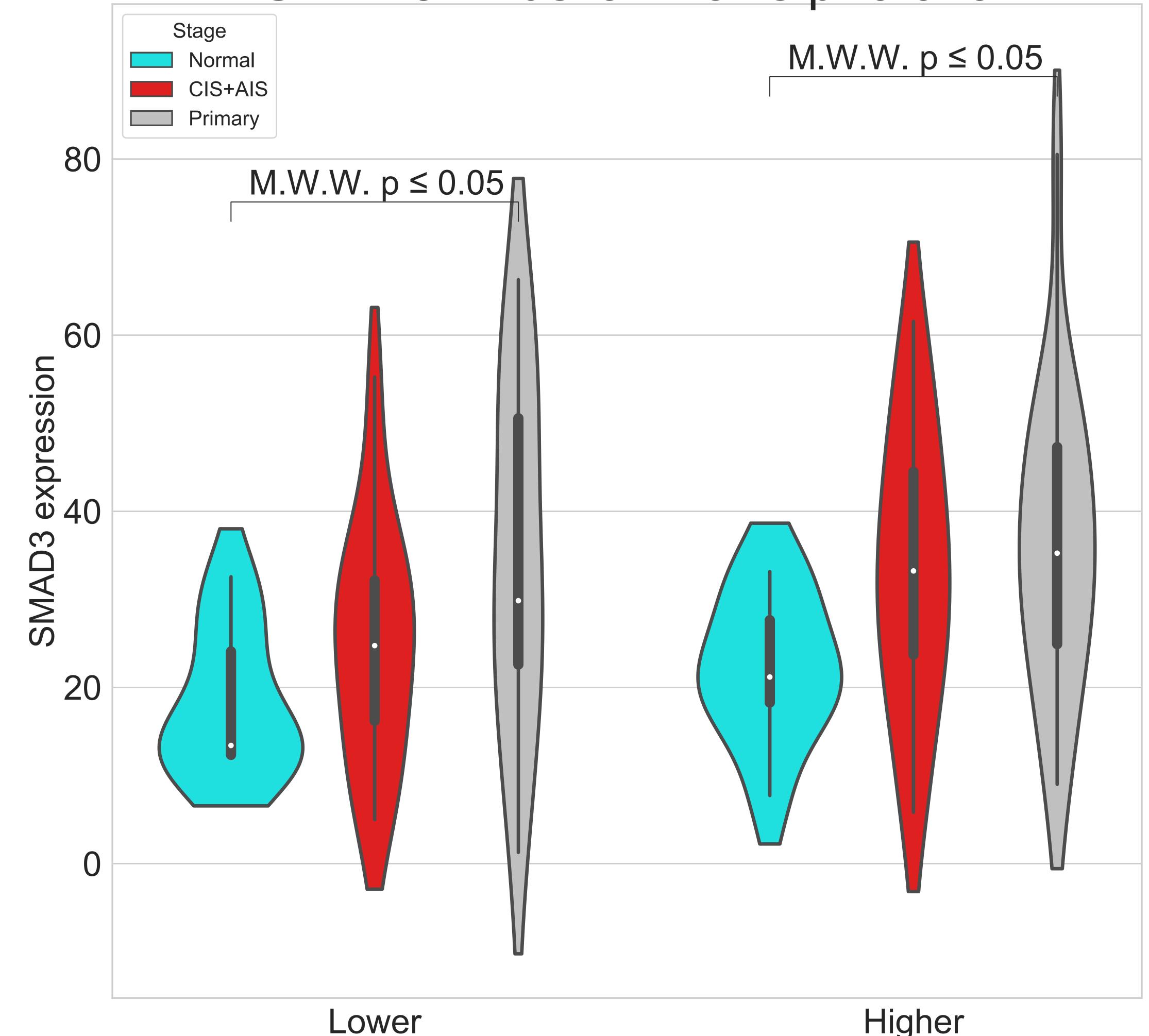
SMAD3: Kruskal-Wallis p=0.029



wer
Mutation Shared Count (SYN)