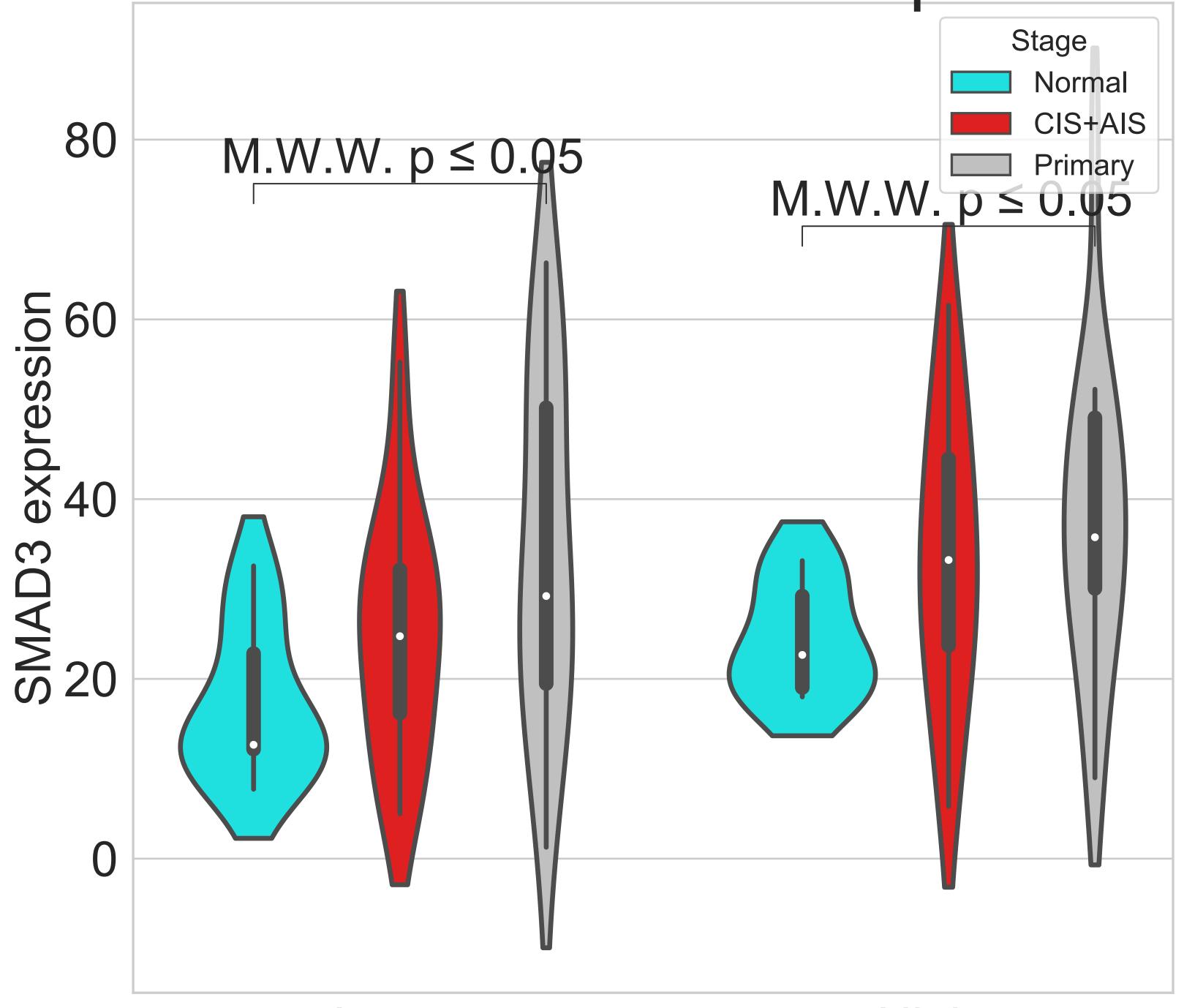
SMAD3: Kruskal-Wallis p=0.018



Lower
Mutation Shared Count (SYN)