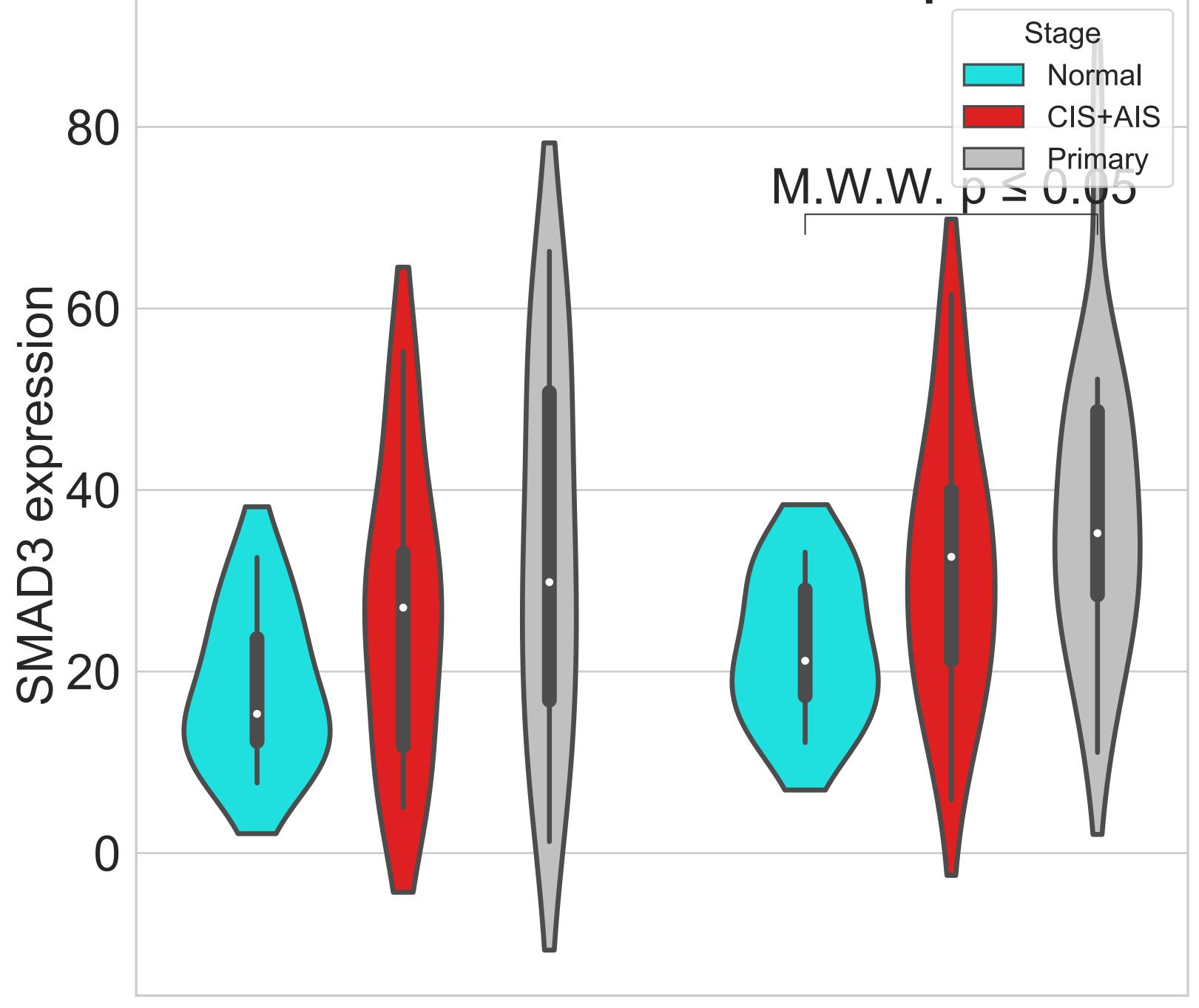
SMAD3: Kruskal-Wallis p=0.037



Lower Higher Mutation Shared Proportion (Union & SYN)