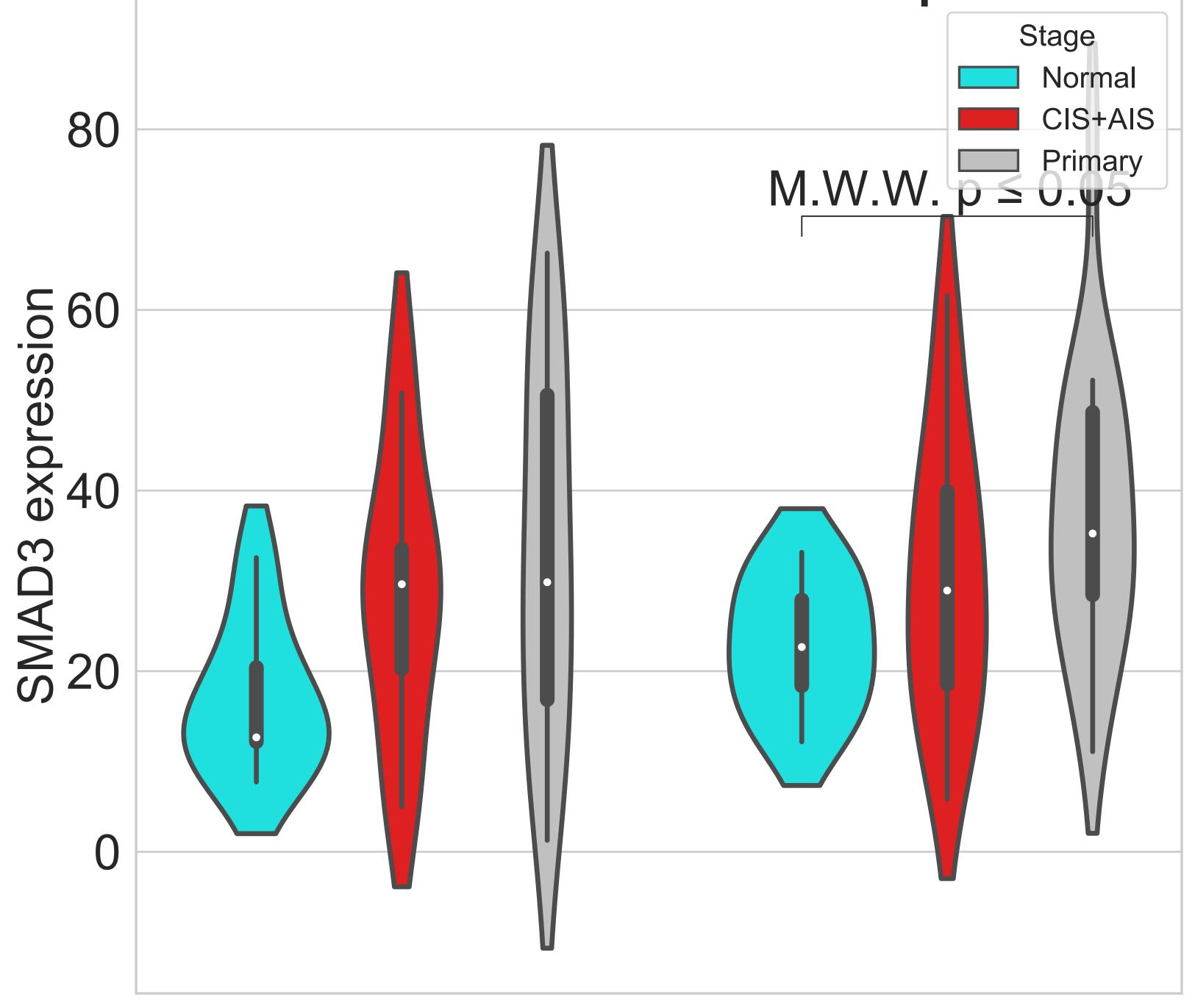
SMAD3: Kruskal-Wallis p=0.045



Lower Higher Mutation Shared Proportion