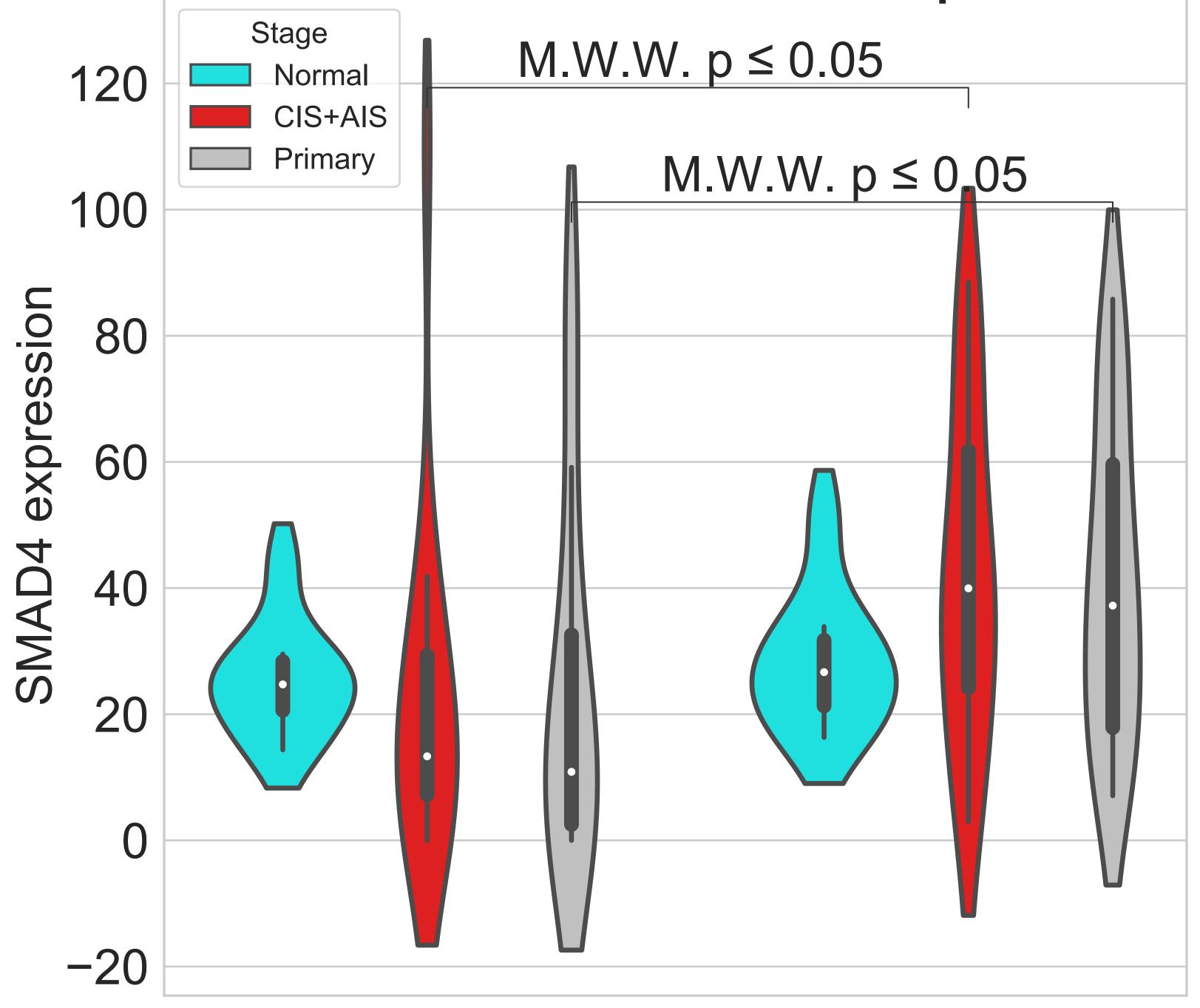
SMAD4: Kruskal-Wallis p=0.025



Lower Higher Mutation Shared Proportion (Union & SYN)