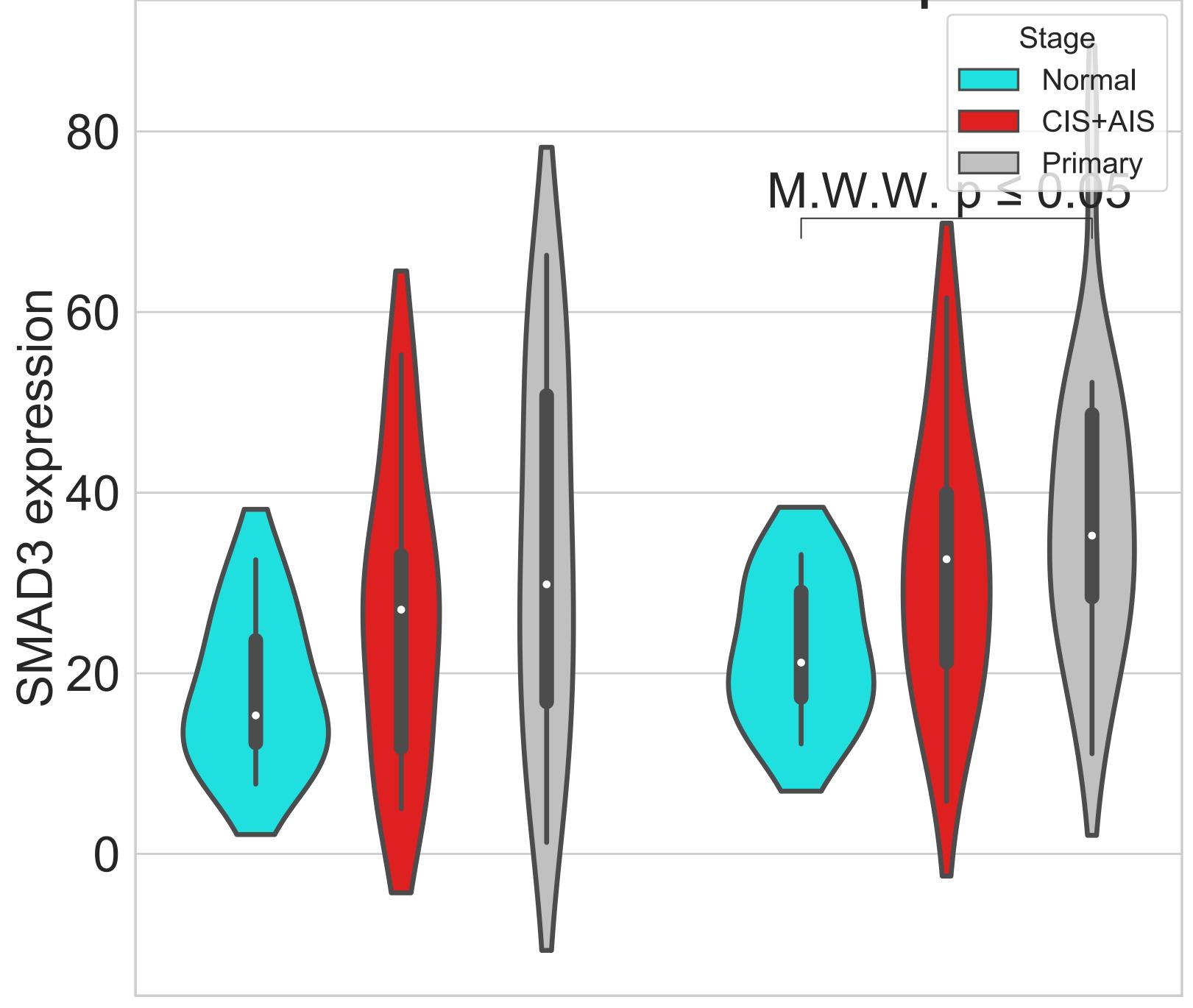
SMAD3: Kruskal-Wallis p=0.037



Lower
Mutation Shared Proportion (Union)