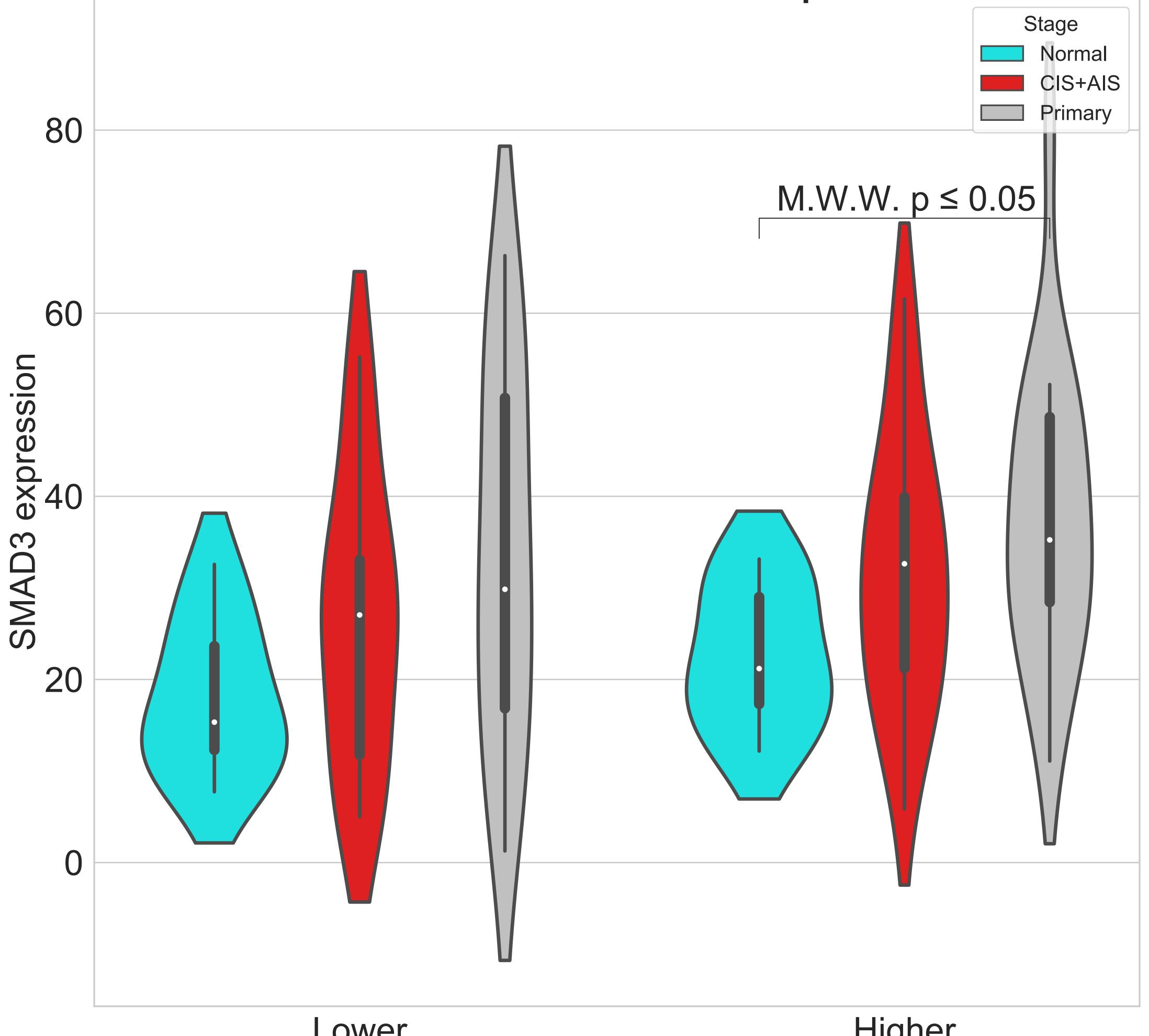
SMAD3: Kruskal-Wallis p=0.037



Higher Lower Mutation Shared Proportion (Union)