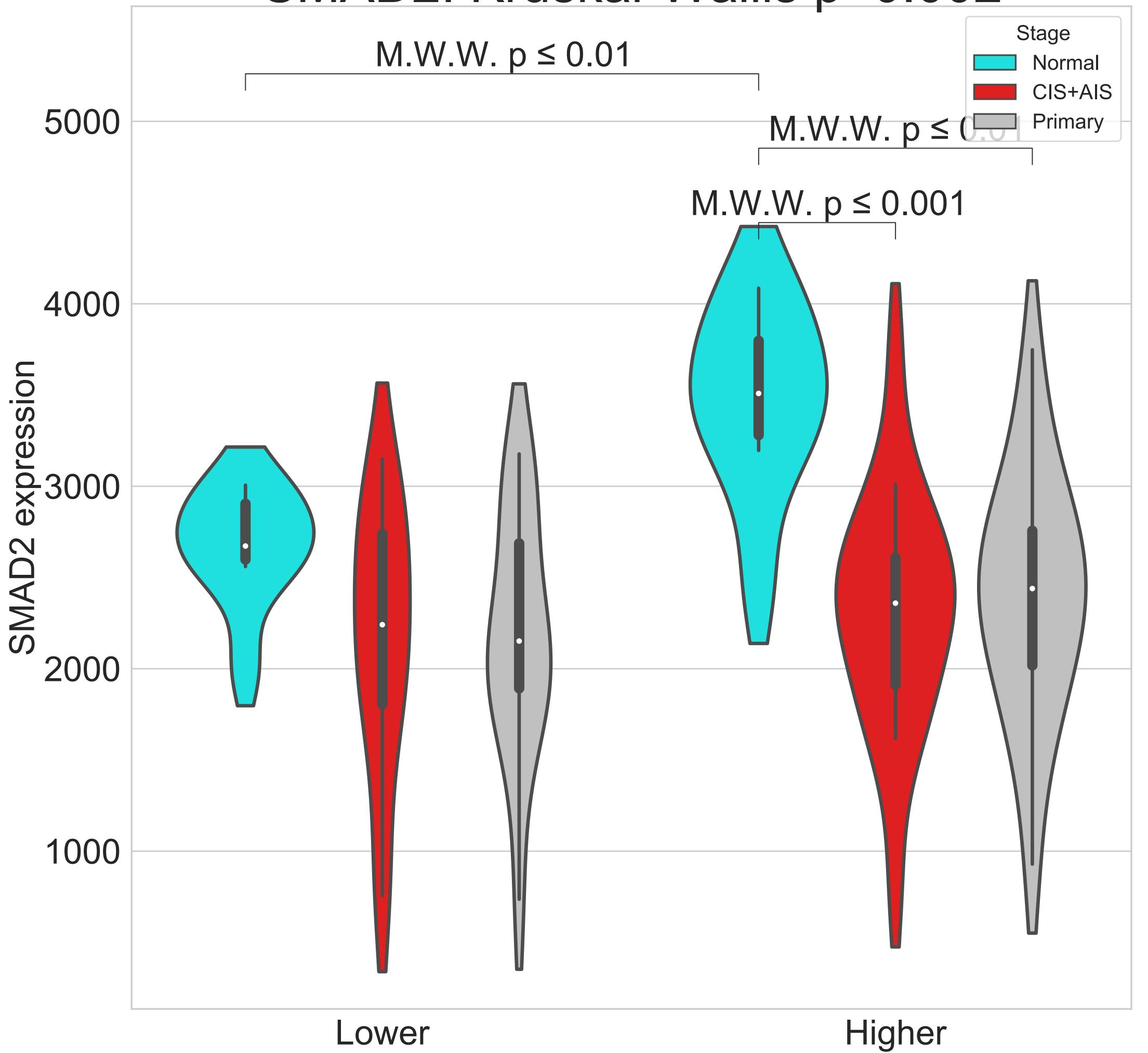
SMAD2: Kruskal-Wallis p=0.002



Mutation Shared Proportion (Union)