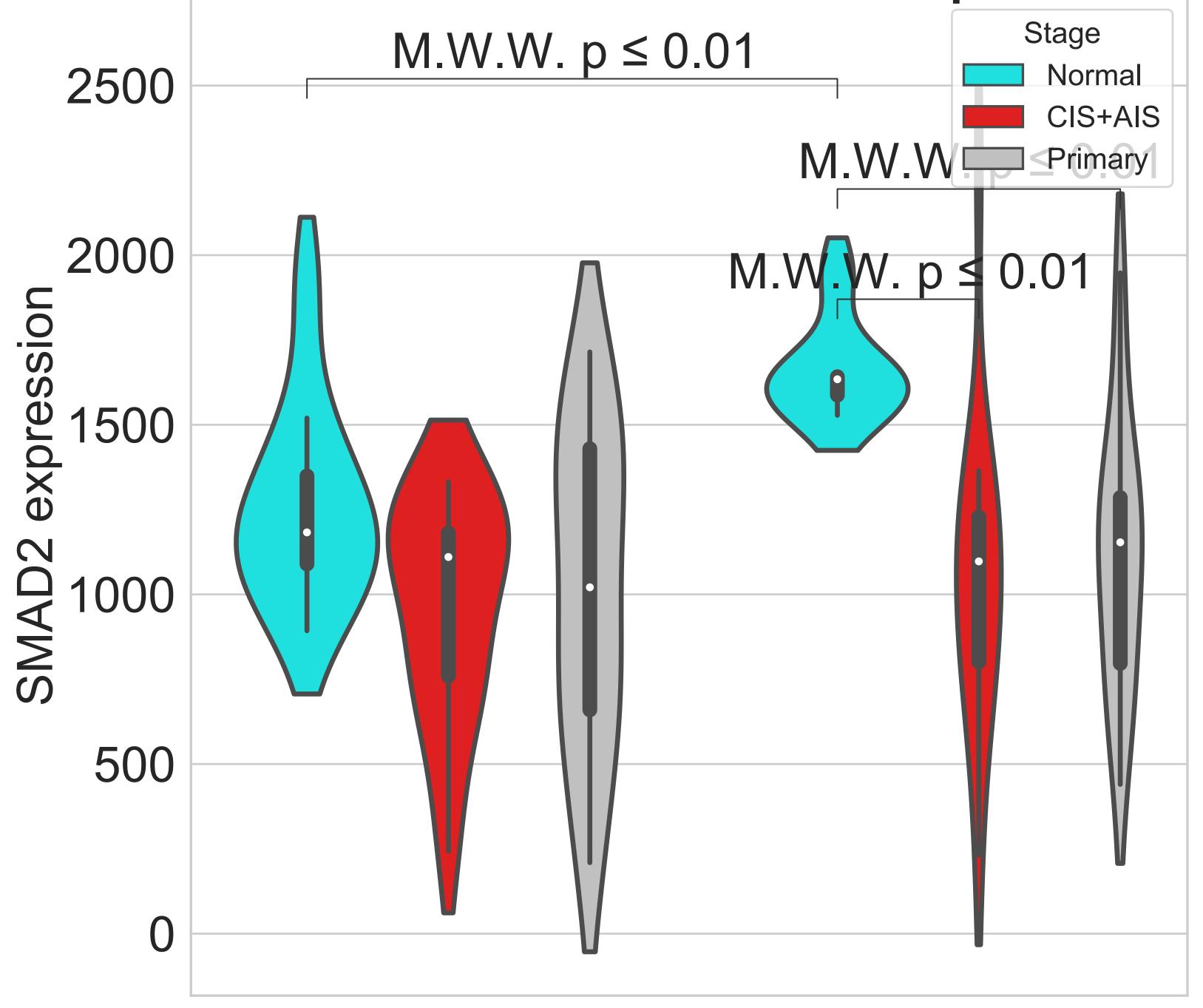
SMAD2: Kruskal-Wallis p=0.010



Lower
Mutation Shared Proportion (SYN)