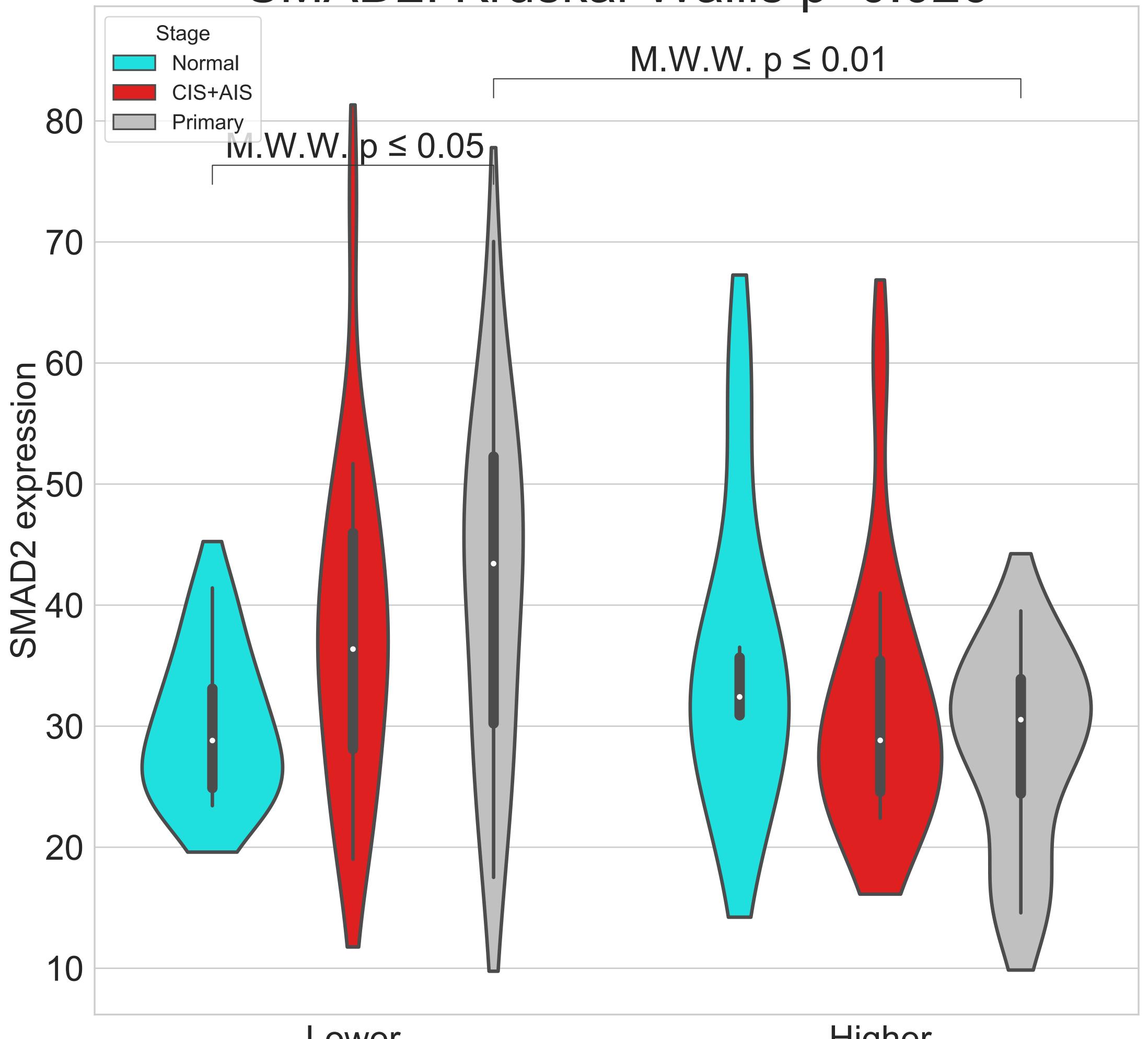
SMAD2: Kruskal-Wallis p=0.026



Lower
Mutation Shared Proportion (Union)