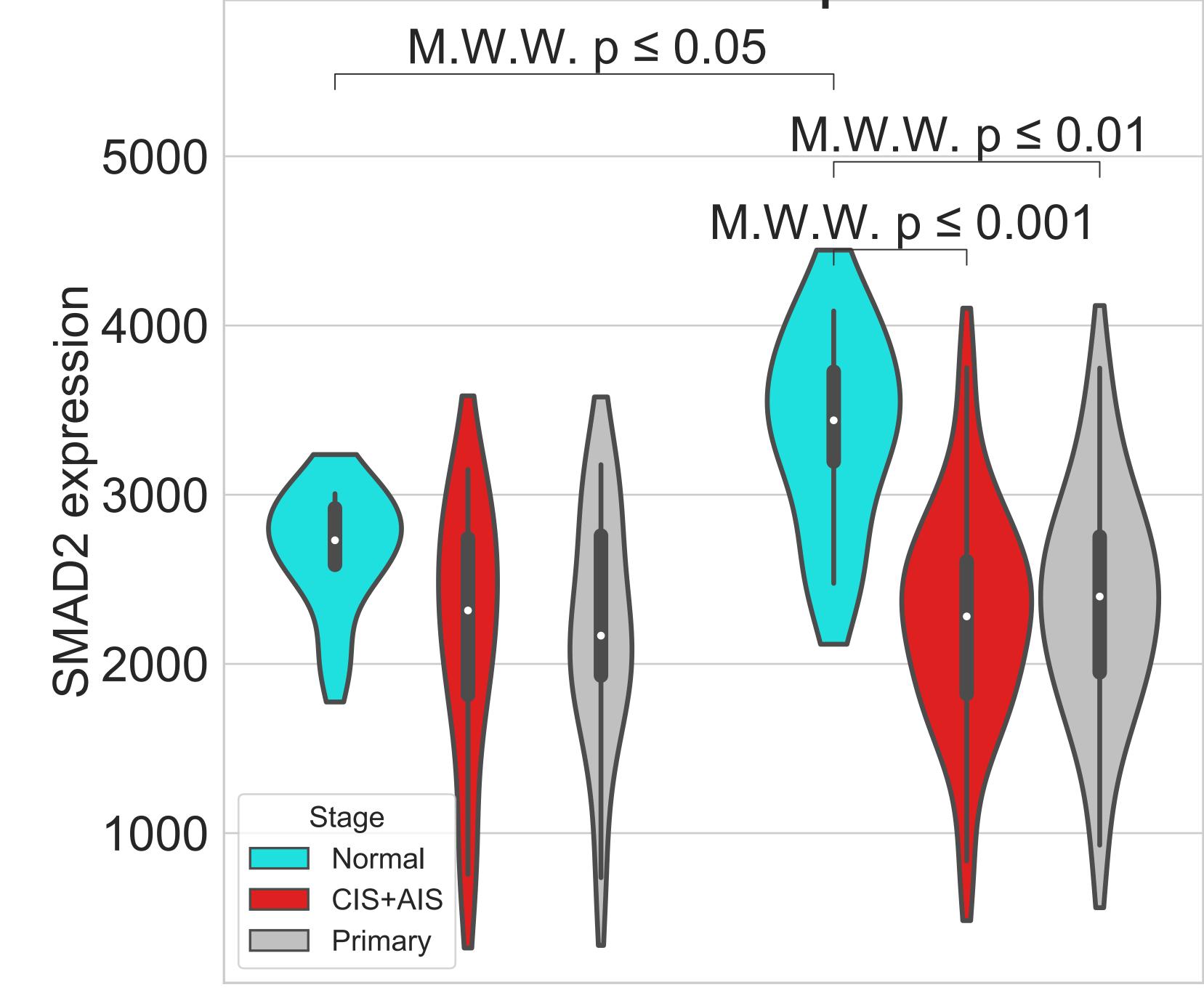
SMAD2: K.W. p=0.003



Lower Higher Mutation Shared Proportion (SYN)