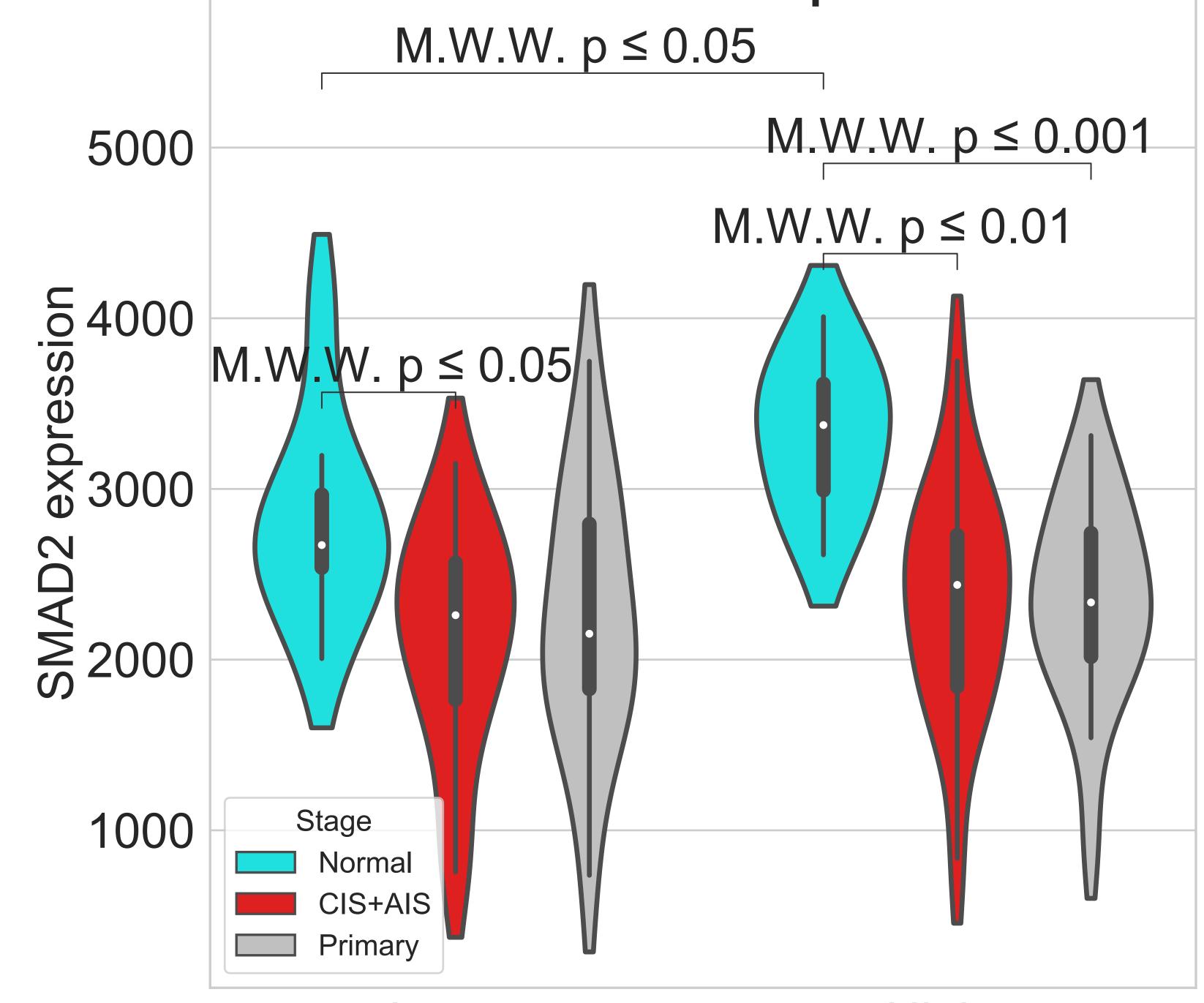
SMAD2: K.W. p=0.003



Lower
Mutation Shared Count (SYN)