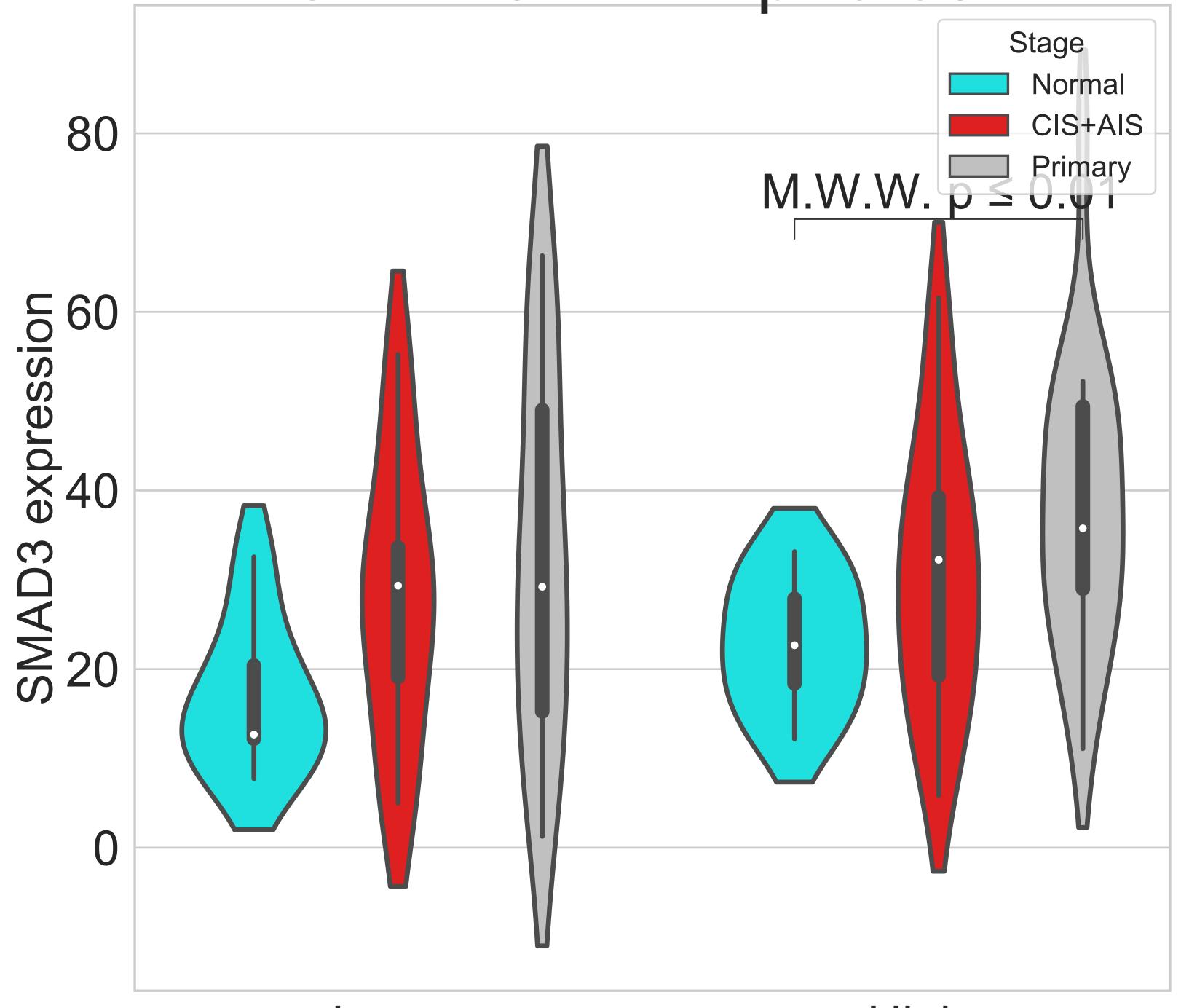
SMAD3: K.W. p=0.034



Lower Higher Mutation Shared Count (SYN) per TMB