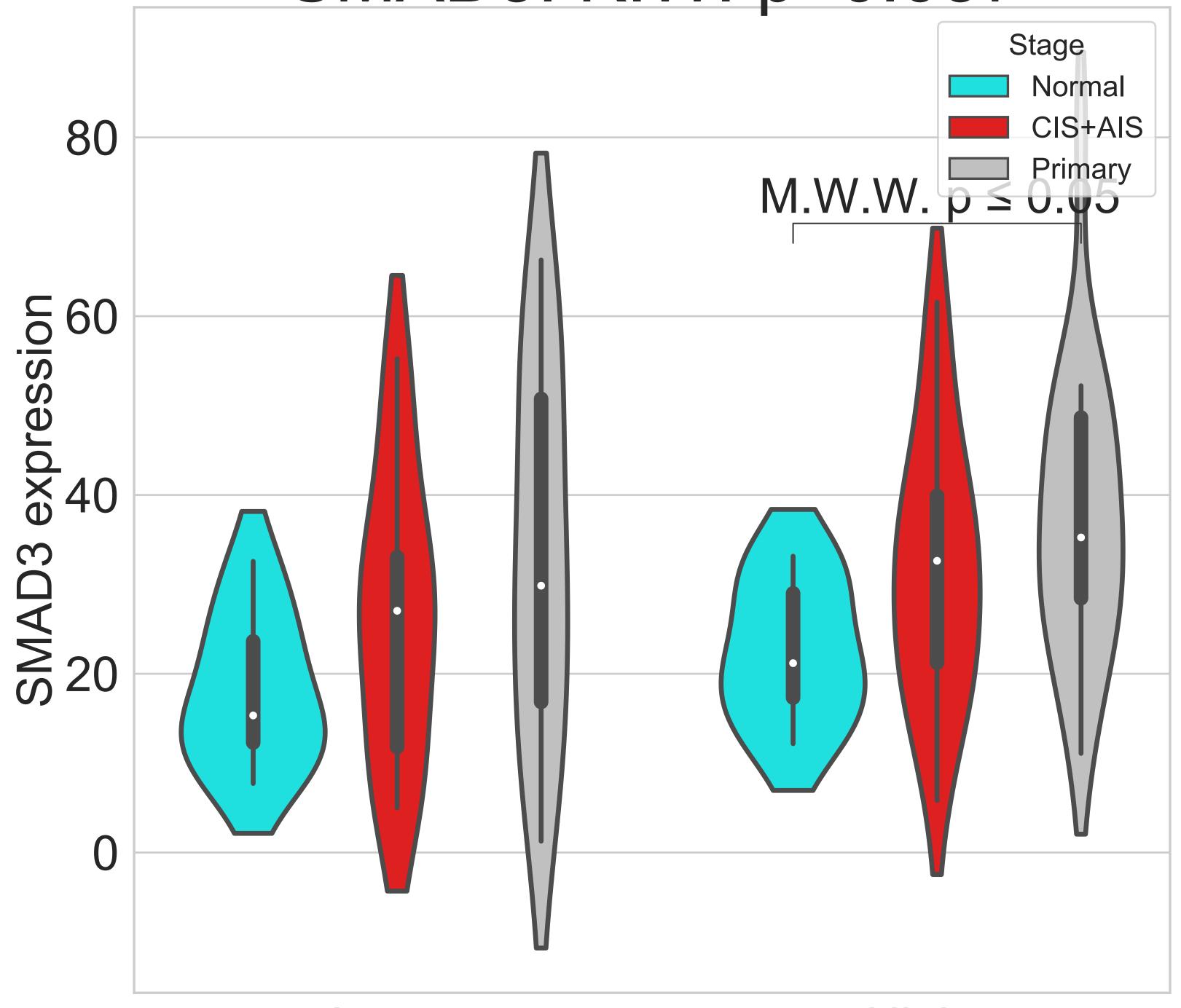
SMAD3: K.W. p=0.037



Lower Higher Mutation Shared Proportion (Union & SYN)