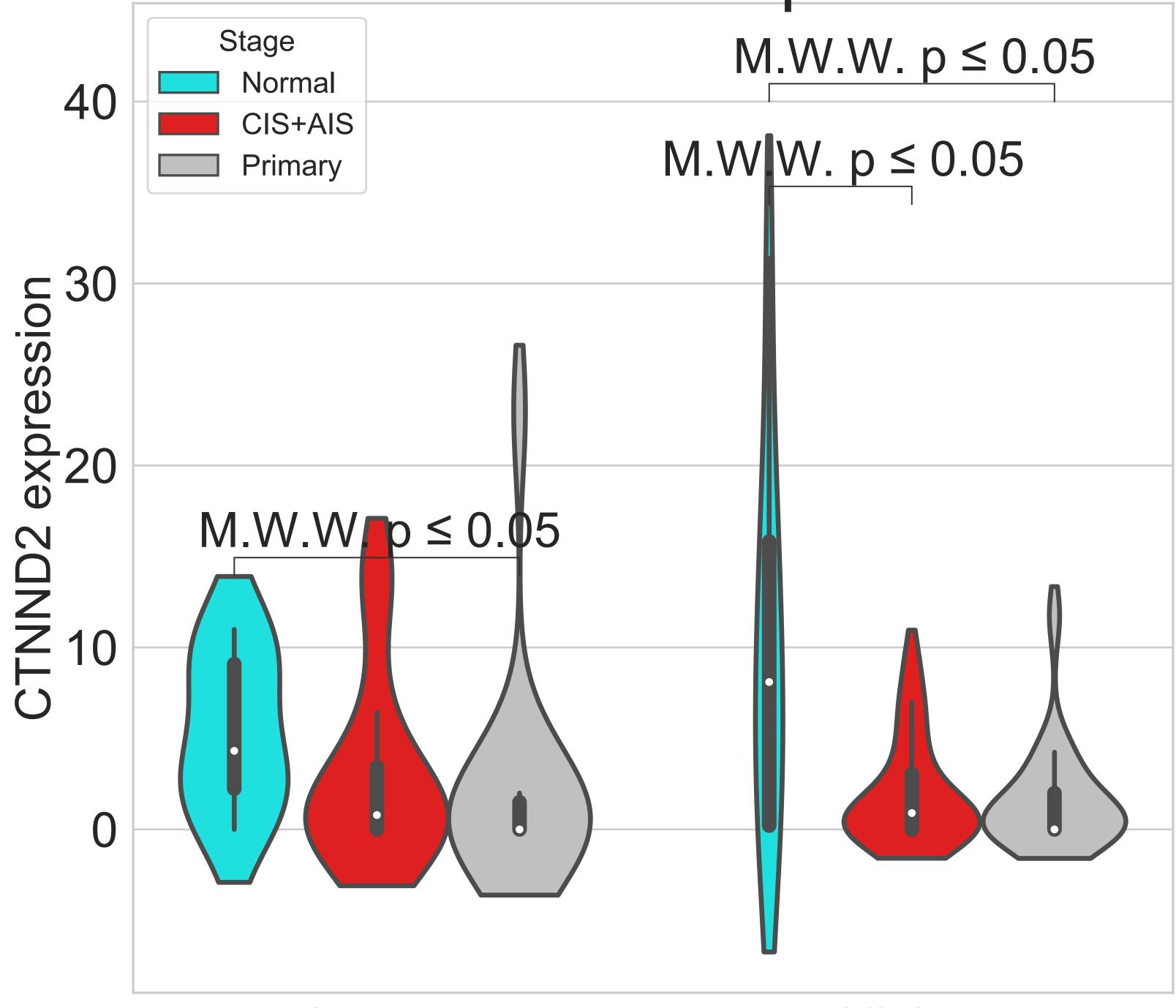
CTNND2: K.W. p=0.012



Lower Higher Mutation Shared Proportion (SYN)