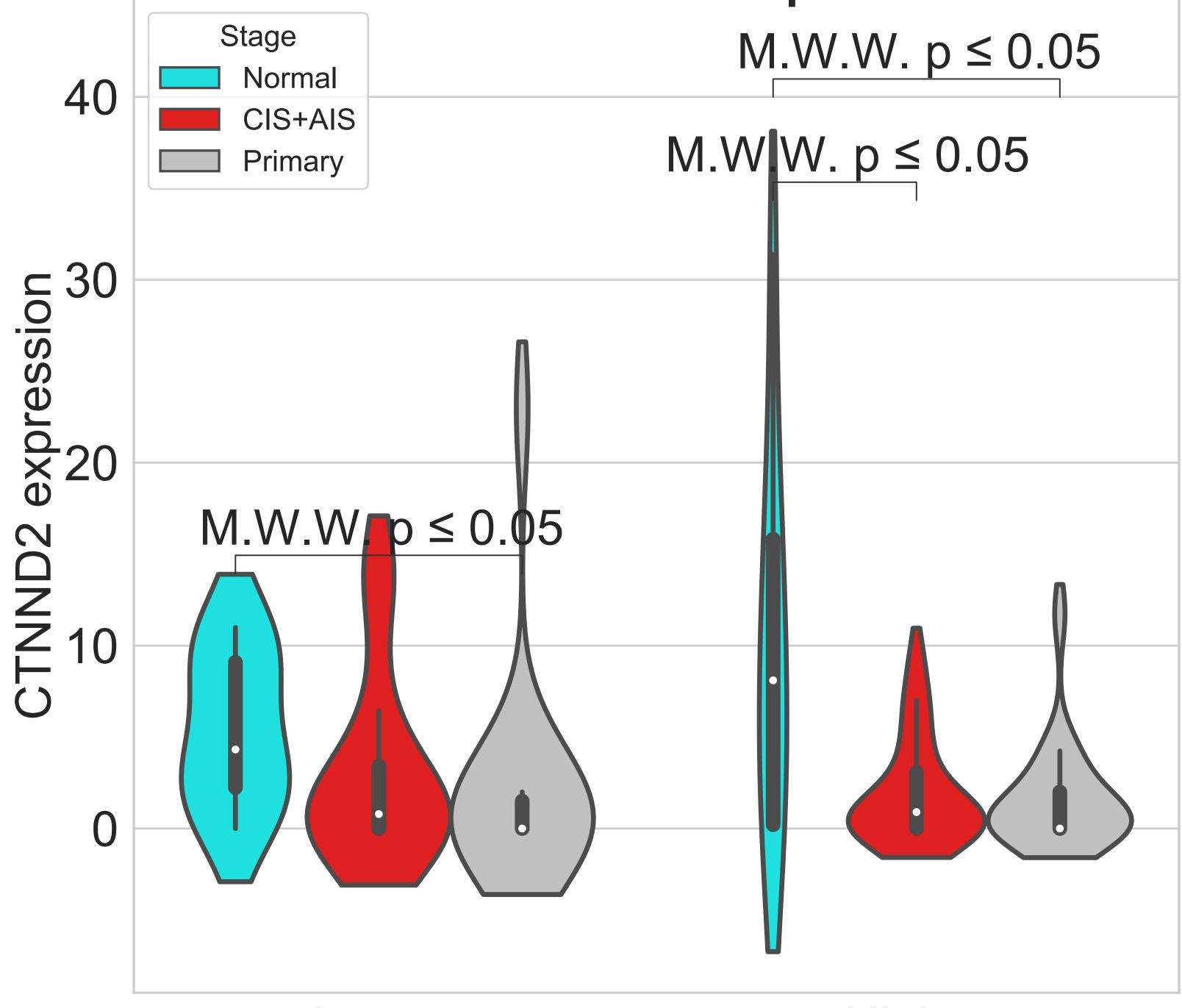
CTNND2: K.W. p=0.012



Lower
Mutation Shared Count (SYN) per TMB