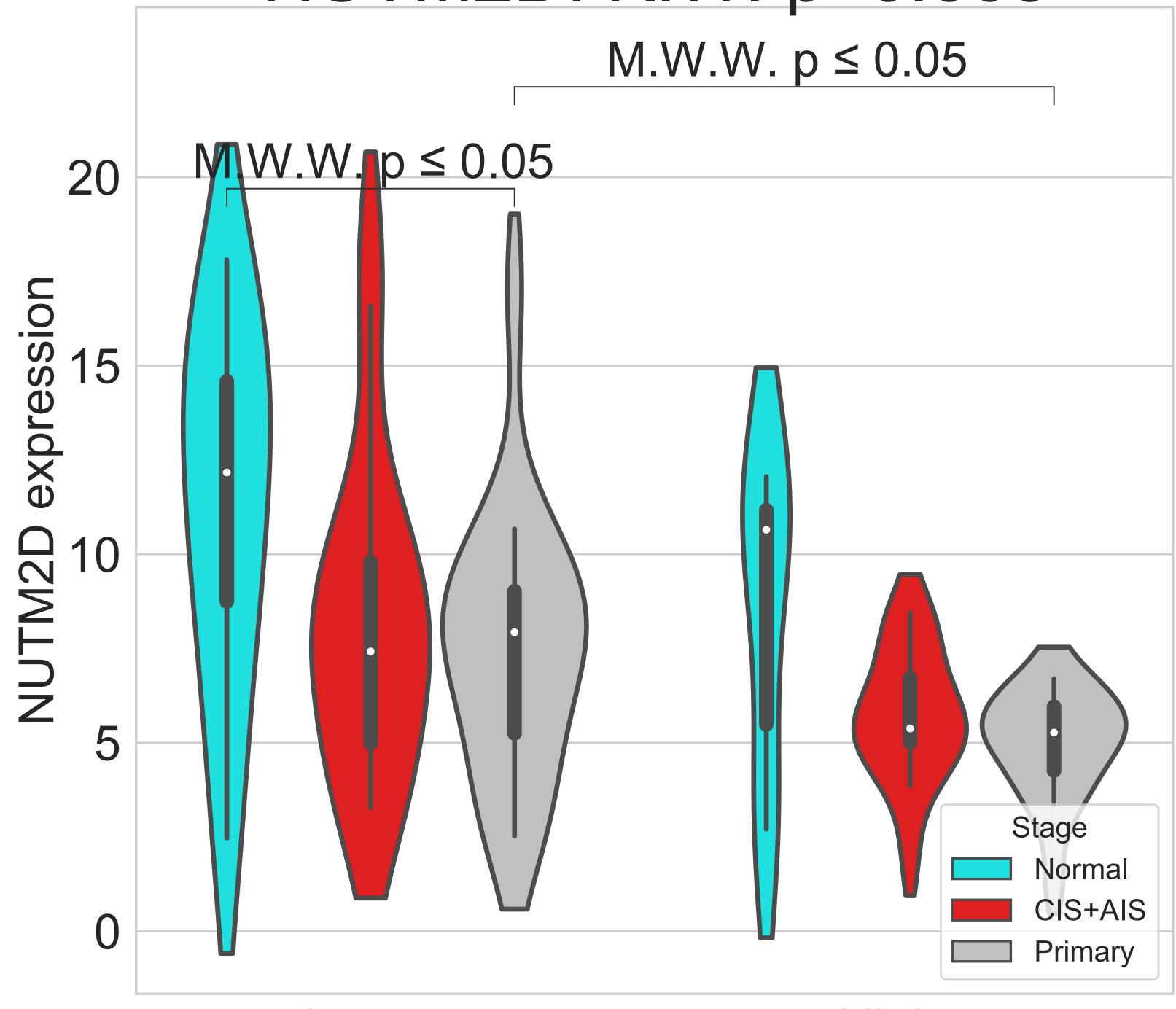
NUTM2D: K.W. p=0.003



Lower
Mutation Shared Count (SYN) per TMB