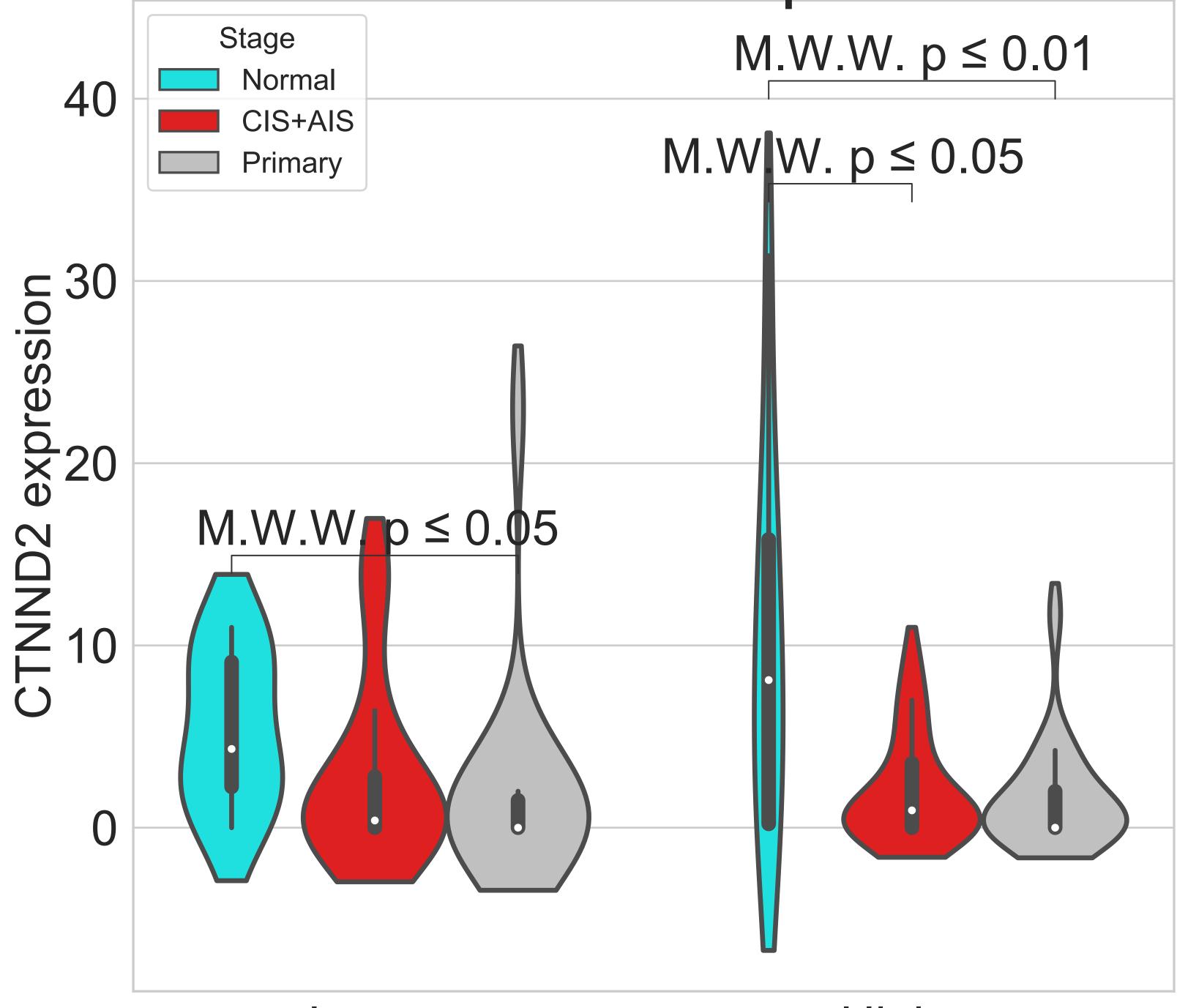
CTNND2: K.W. p=0.013



Lower
Mutation Shared Count per TMB