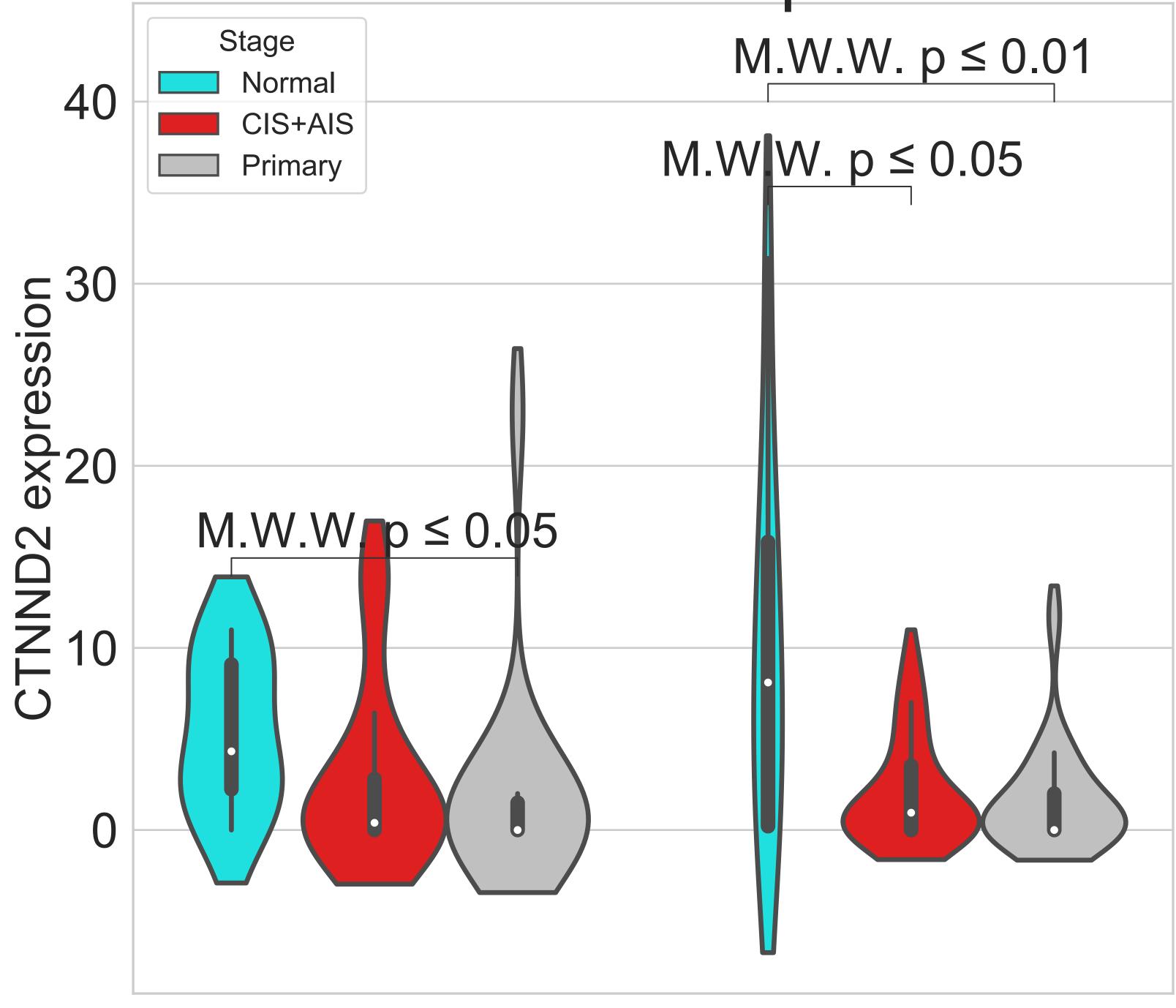
CTNND2: K.W. p=0.013



Lower

Mutation Shared Proportion