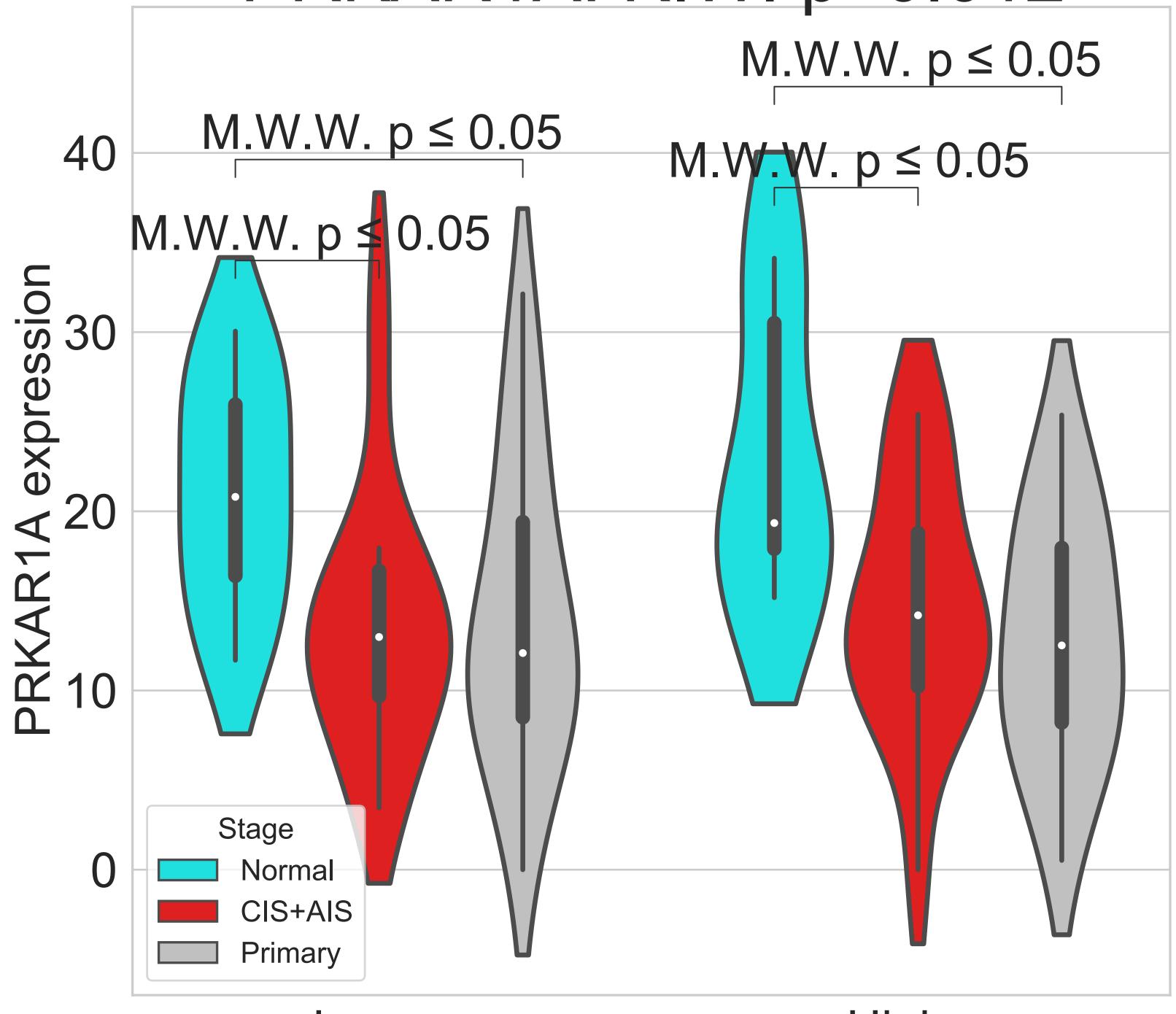
PRKAR1A: K.W. p=0.012



Lower Higher Mutation Shared Count per TMB