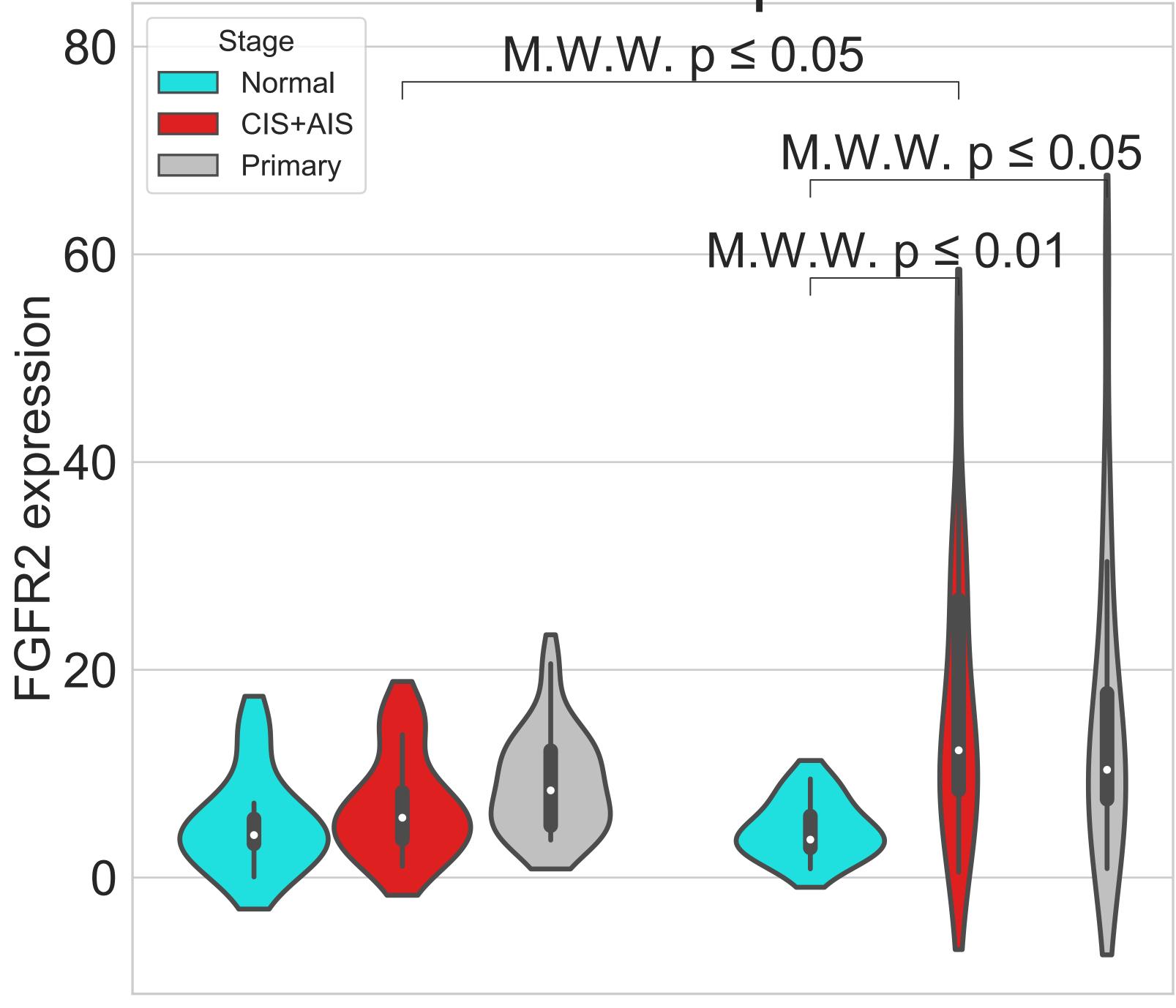
FGFR2: K.W. p=0.004



Lower Higher Mutation Shared Count