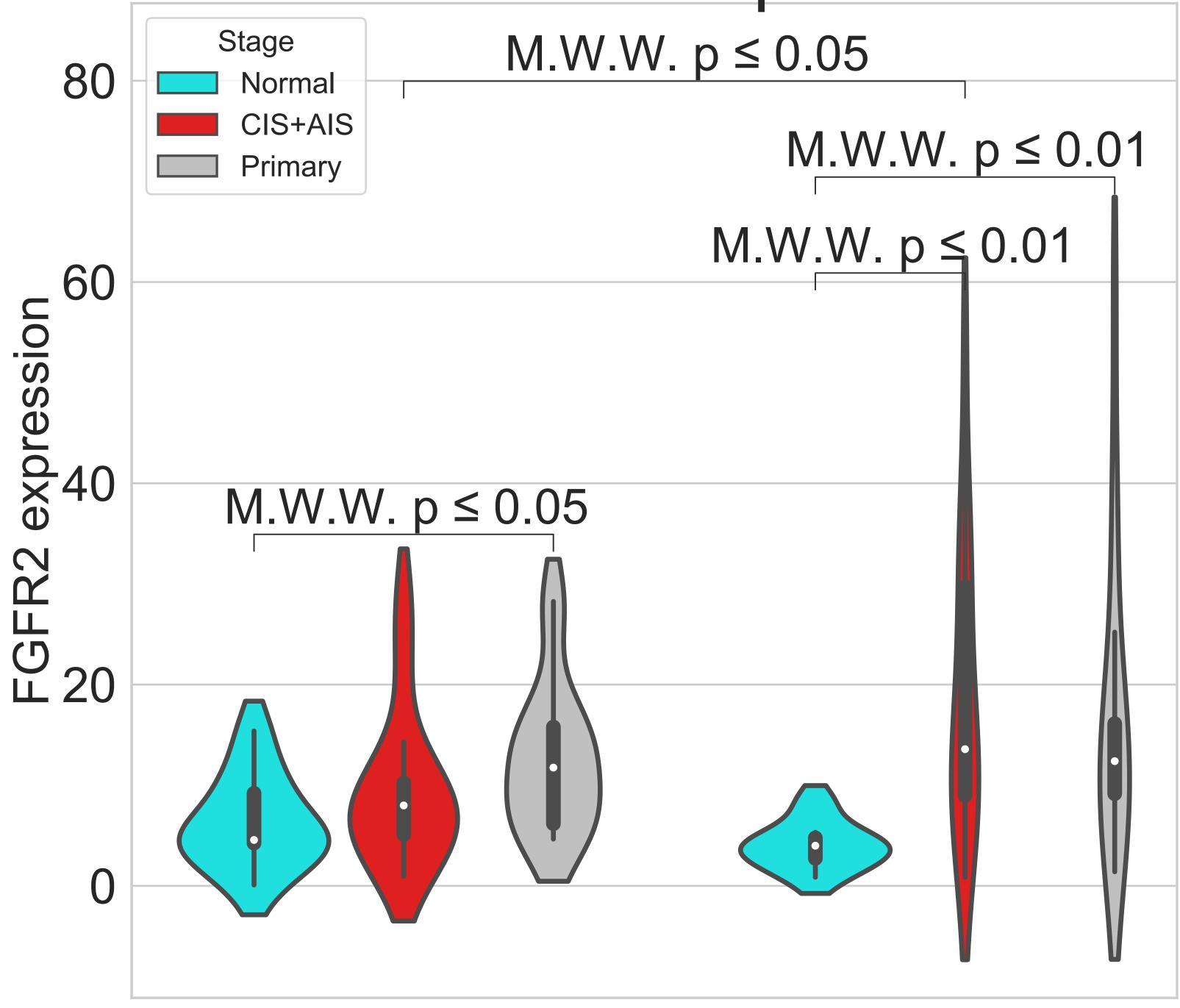
FGFR2: K.W. p=0.001



Lower
Mutation Shared Count (SYN)