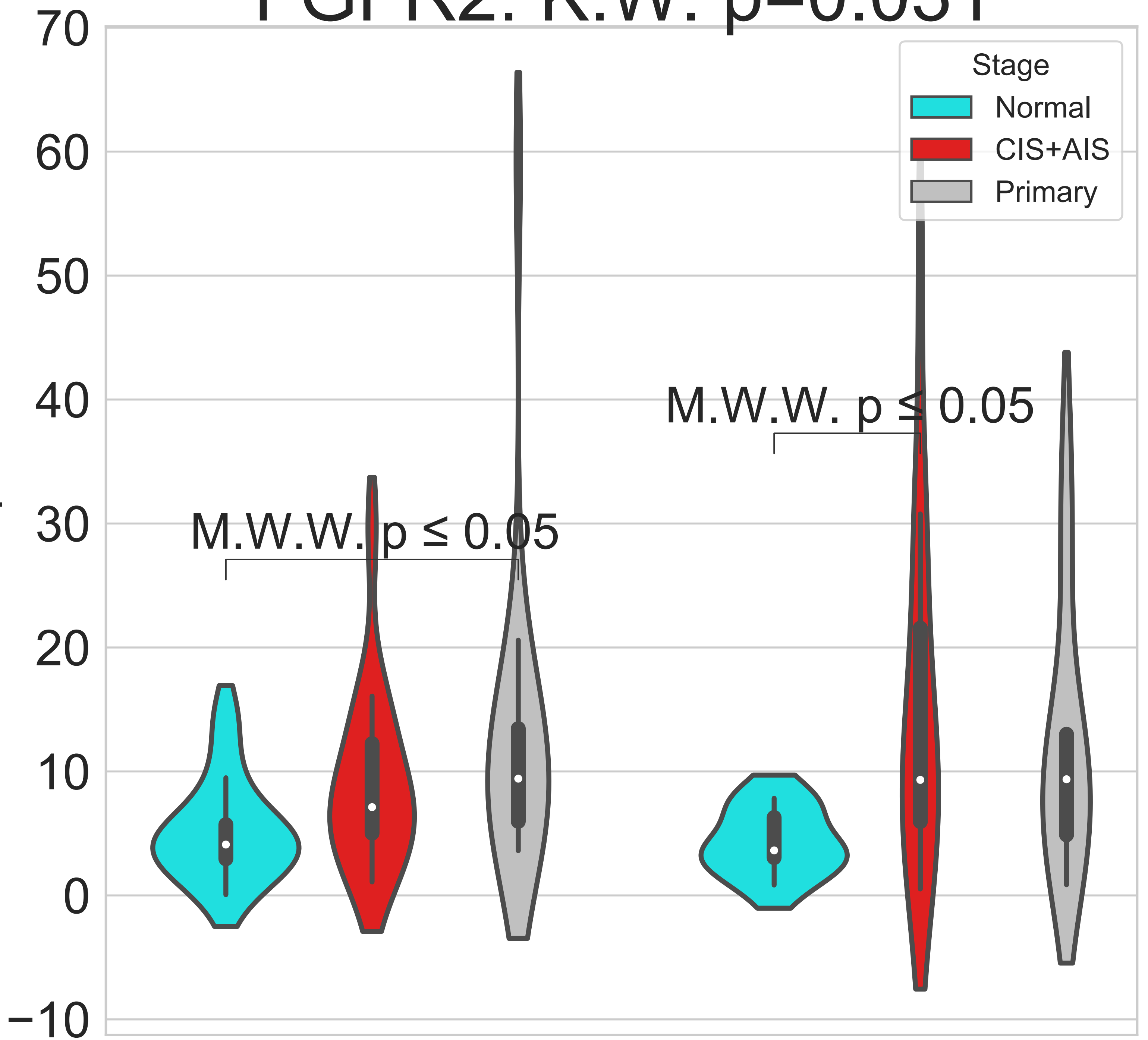


FGFR2: K.W. $p=0.031$

FGFR2 expression



Lower Higher
Mutation Shared Count (SYN) per TMB