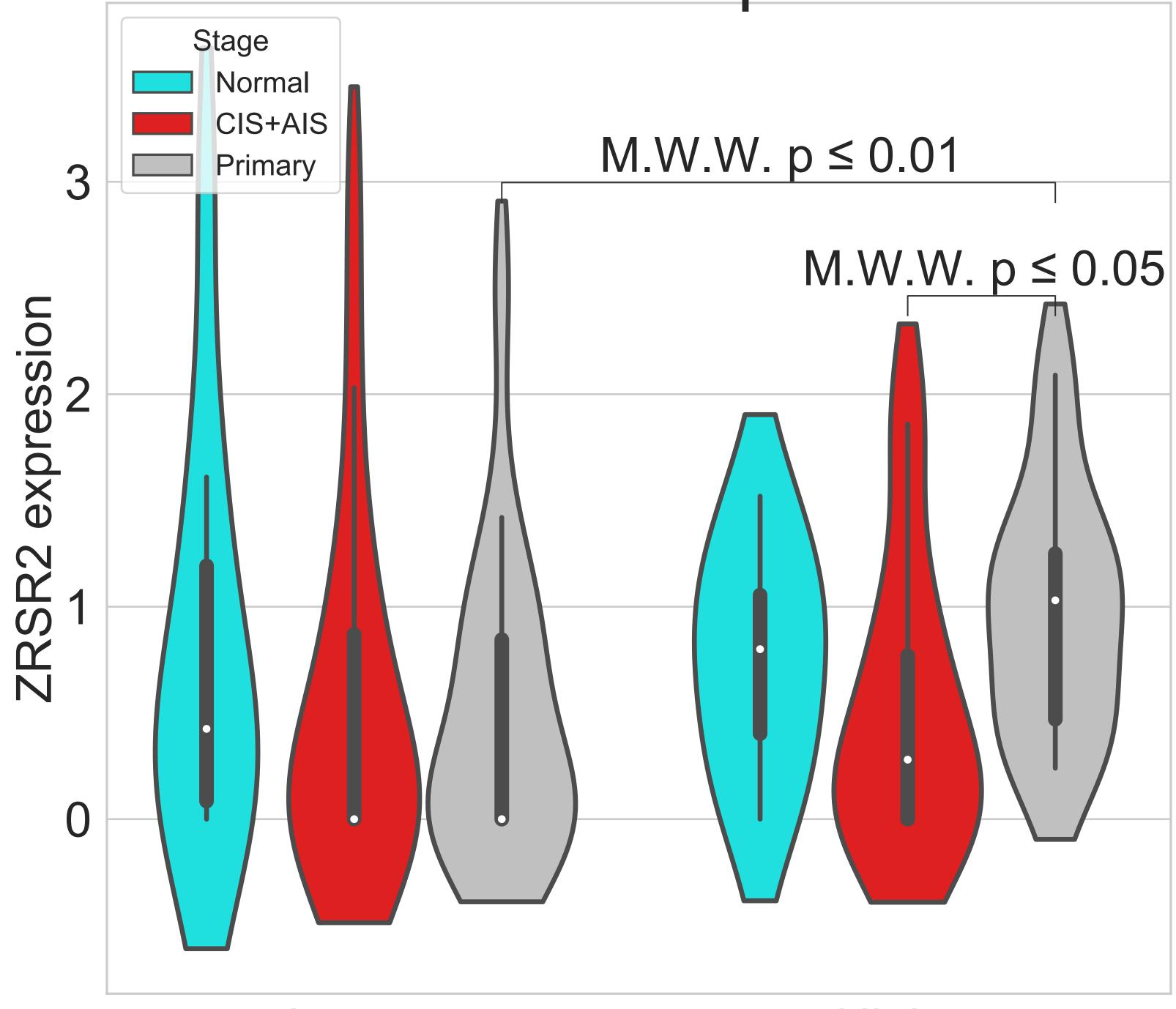
ZRSR2: K.W. p=0.050



Lower Higher Mutation Shared Count (SYN) per TMB