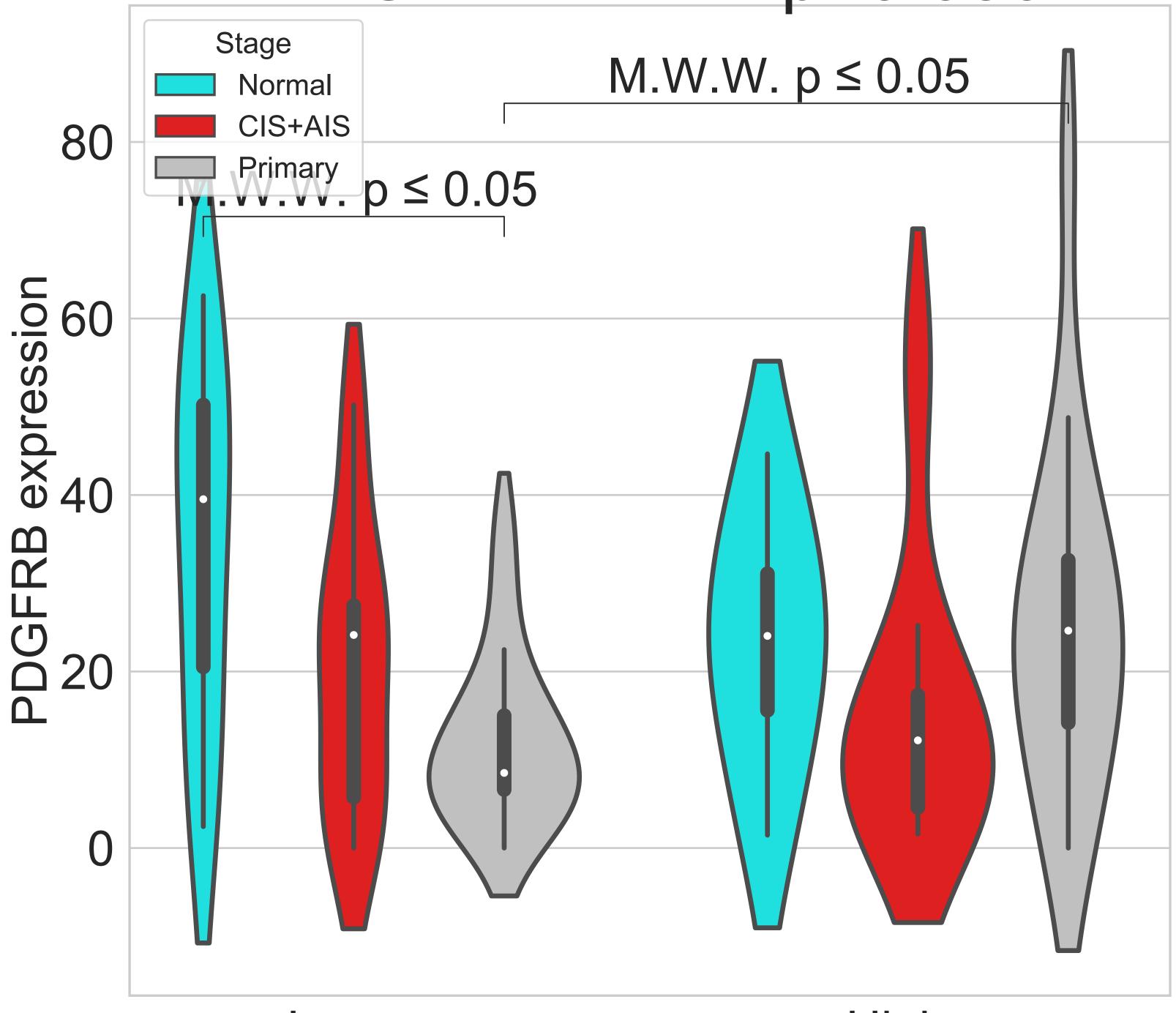
PDGFRB: K.W. p=0.050



Lower
Mutation Shared Count (SYN) per TMB