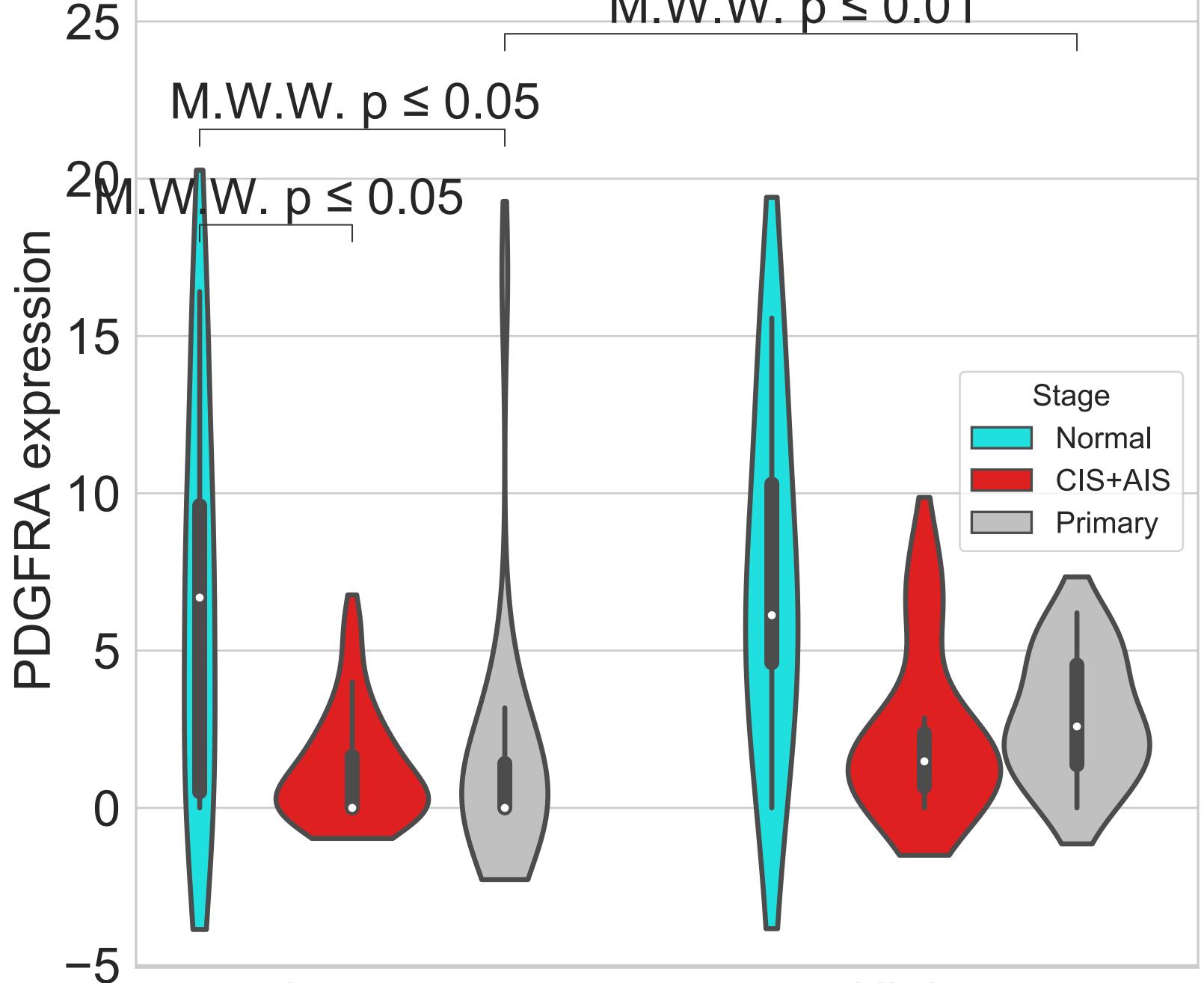
PDGFRA: K.W. p=0.003 M.W.W. p ≤ 0.01



Lower Higher Mutation Shared Count (SYN) per TMB