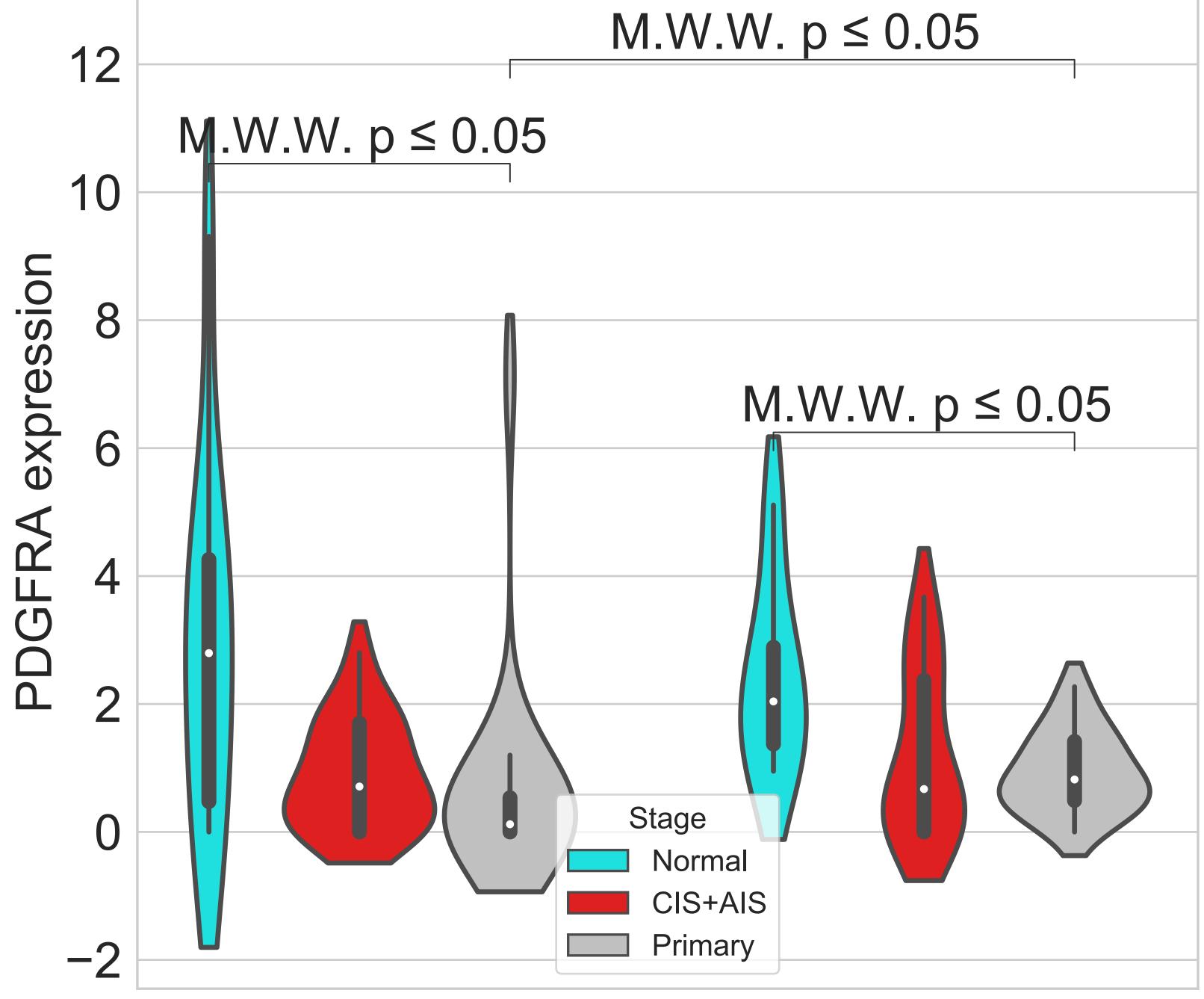
PDGFRA: K.W. p=0.007

M.W.W. p ≤ 0.05



Lower Higher Mutation Shared Count (SYN) per TMB