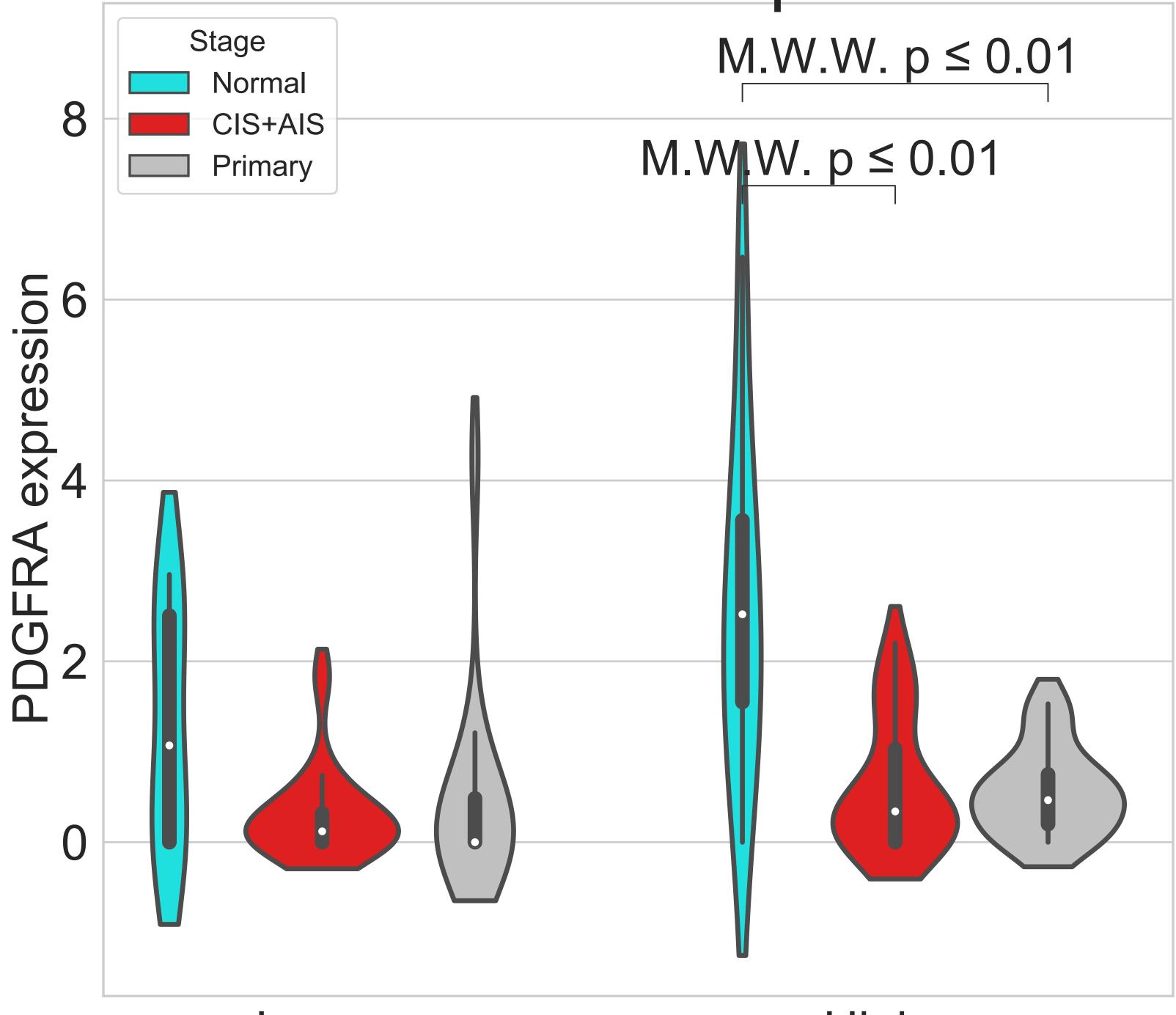
PDGFRA: K.W. p=0.002



Lower Higher Mutation Shared Count per TMB