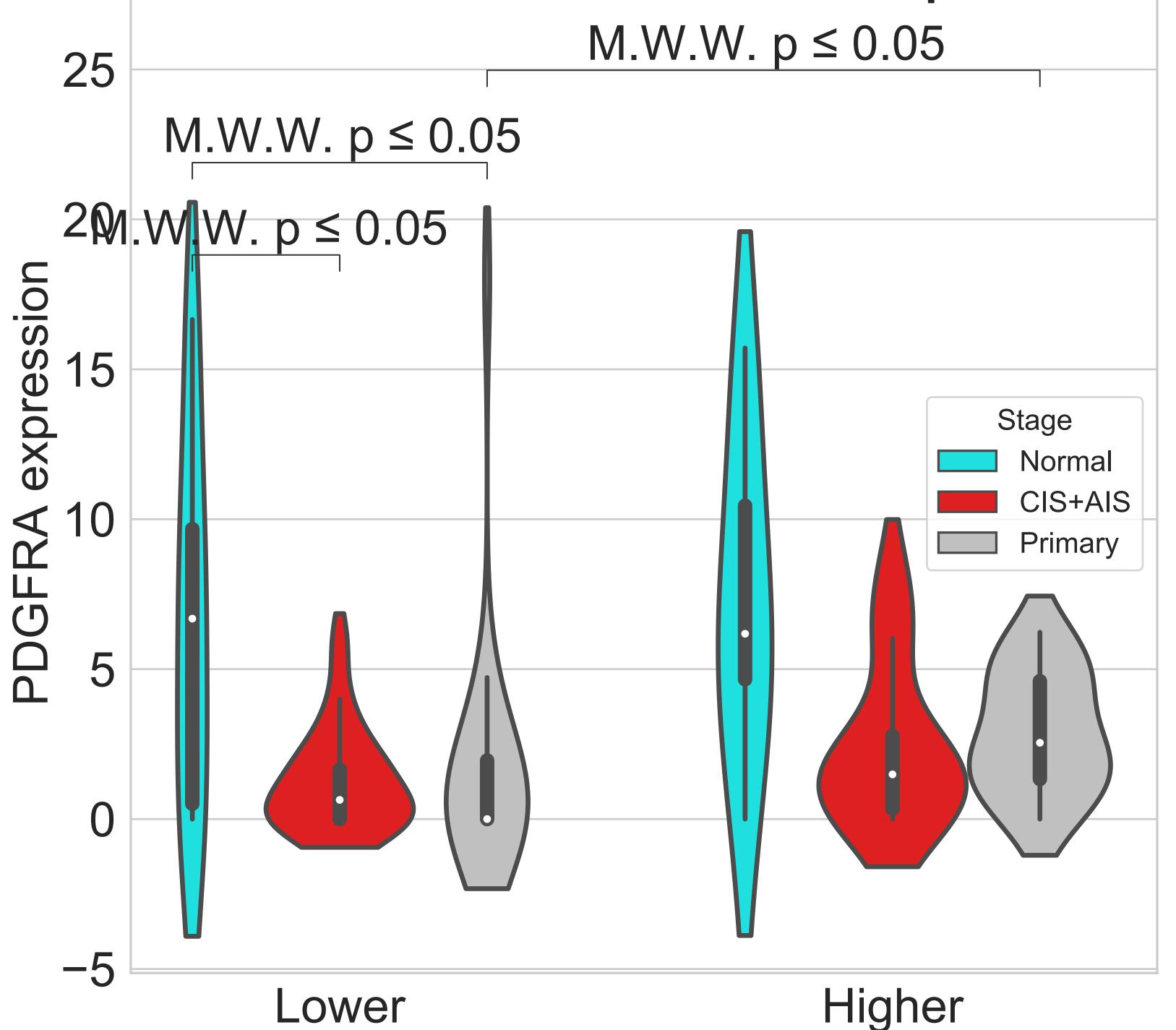
PDGFRA: Kruskal-Wallis p=0.006



Mutation Shared Count per TMB