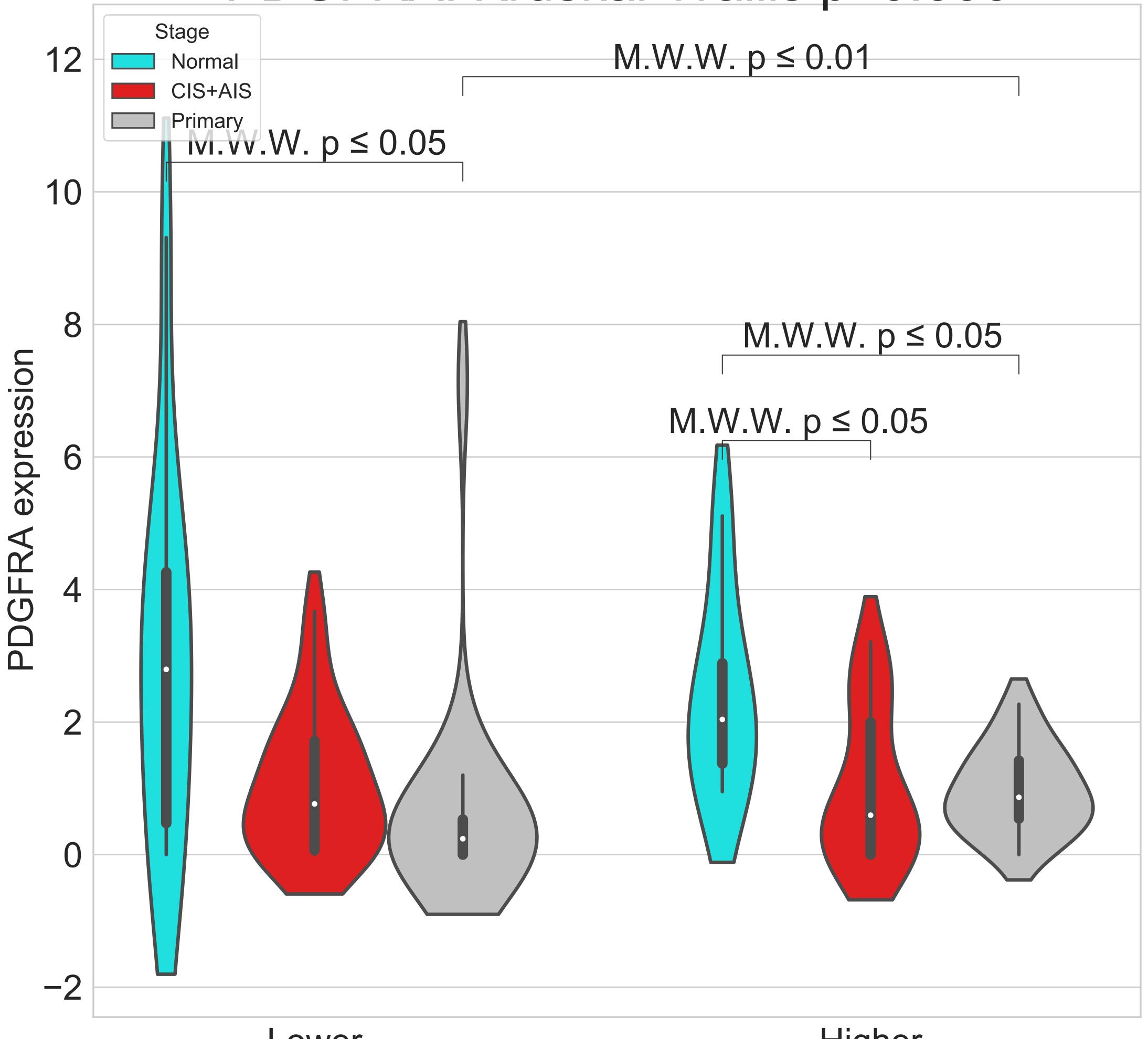
PDGFRA: Kruskal-Wallis p=0.006



Lower
Mutation Shared Count per TMB