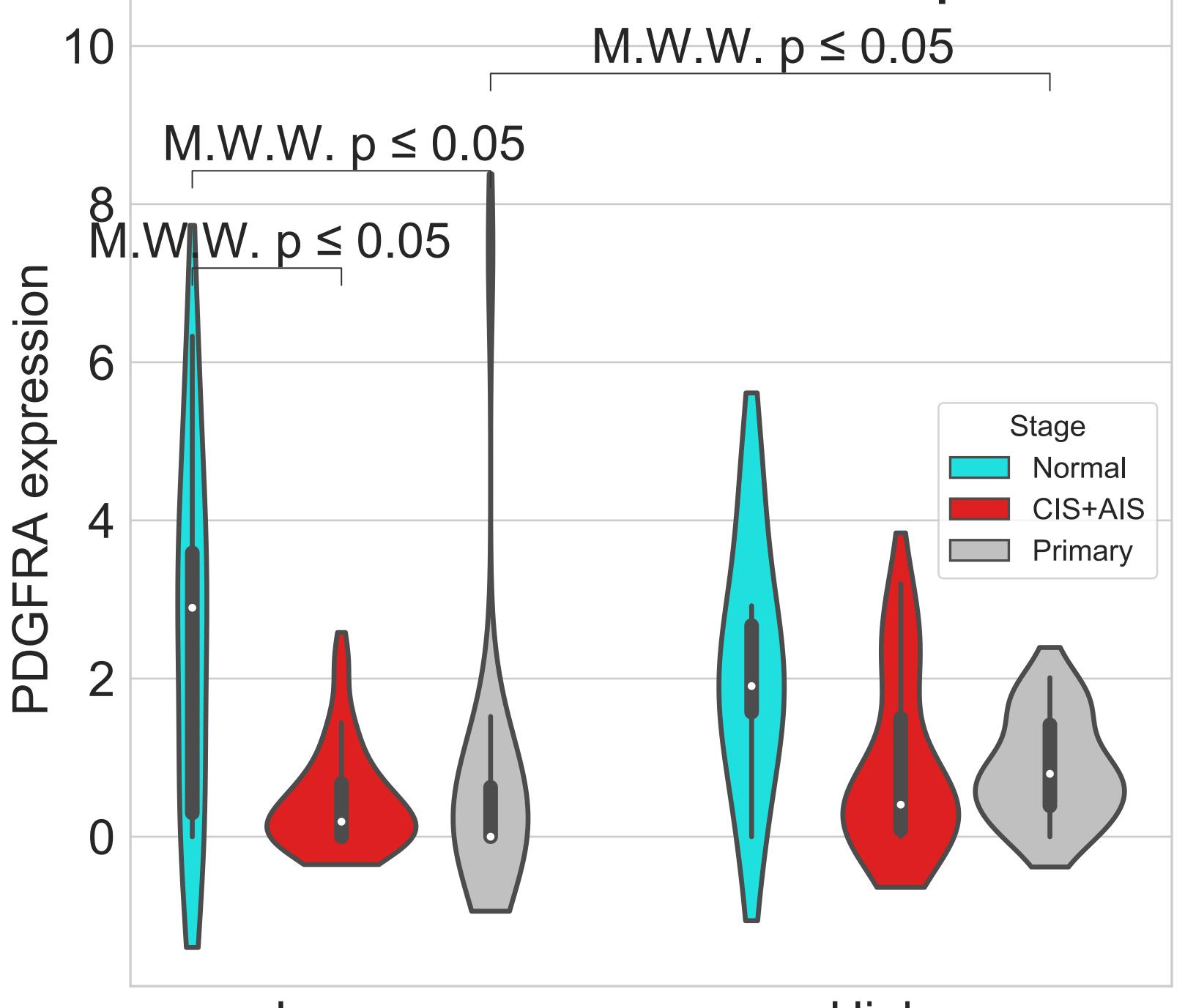
PDGFRA: Kruskal-Wallis p=0.009



Lower

Mutation Shared Count per TMB