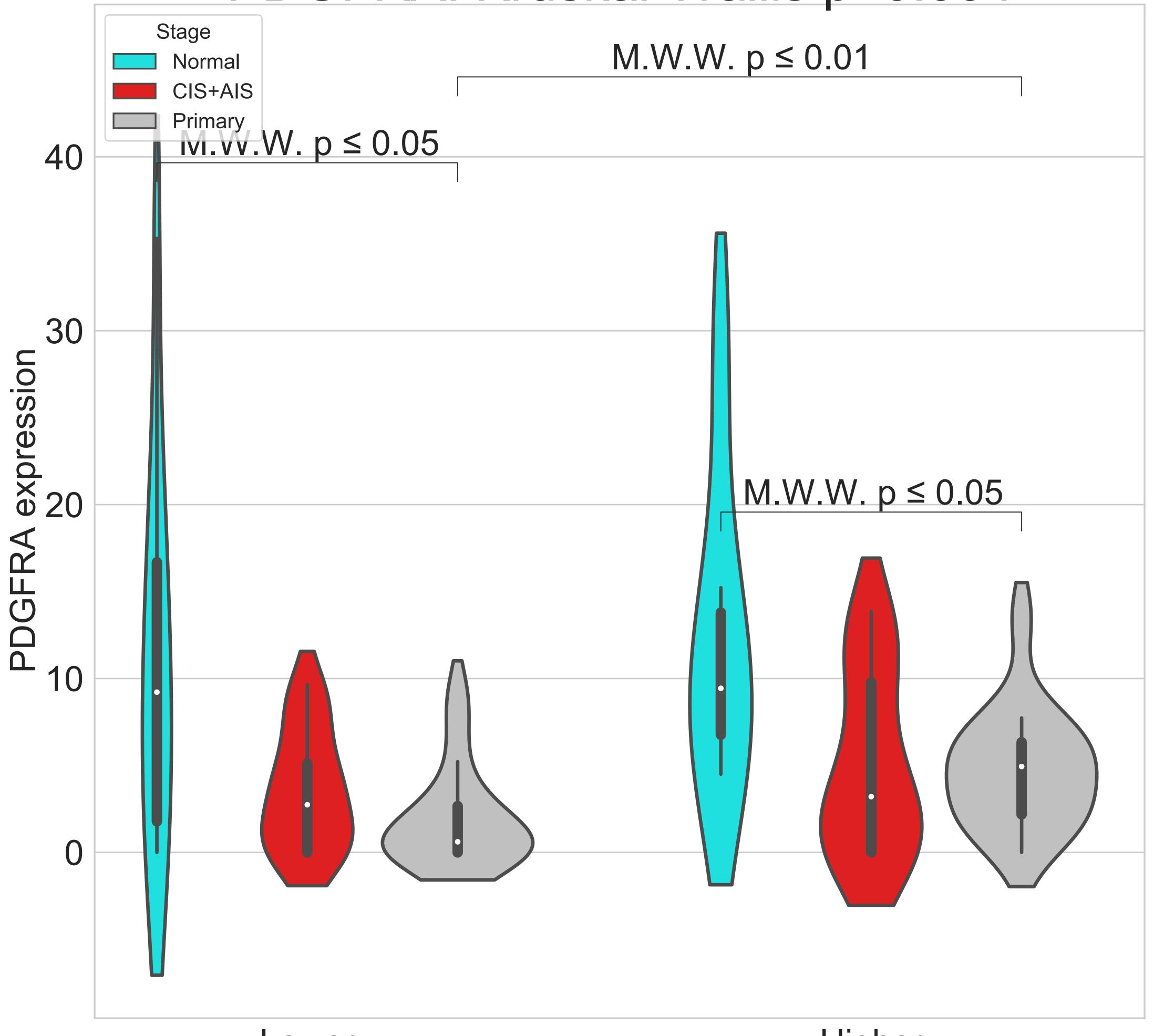
PDGFRA: Kruskal-Wallis p=0.004



Lower
Mutation Shared Count (SYN) per TMB