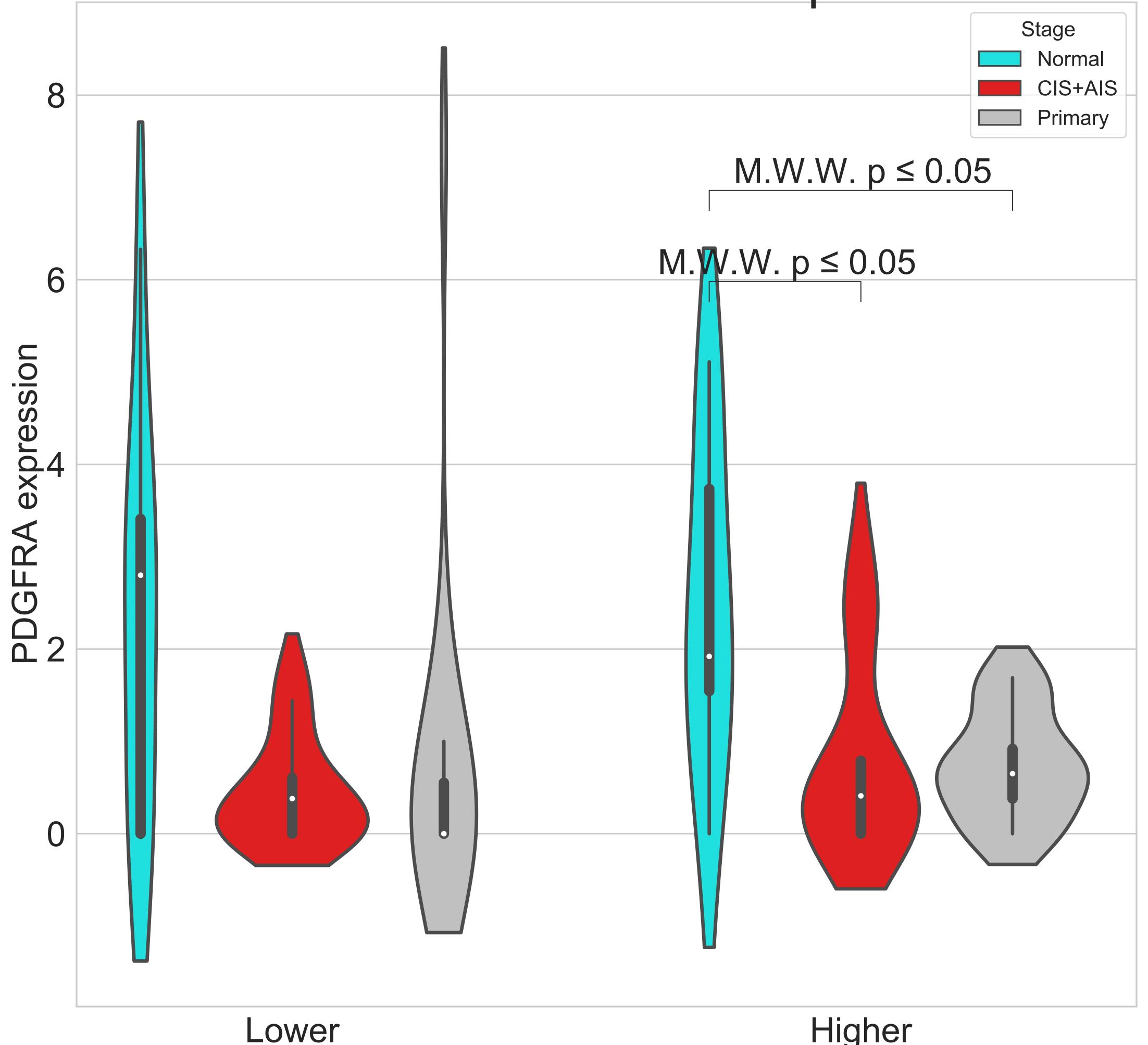
PDGFRA: Kruskal-Wallis p=0.016



Higher Mutation Shared Count (SYN)