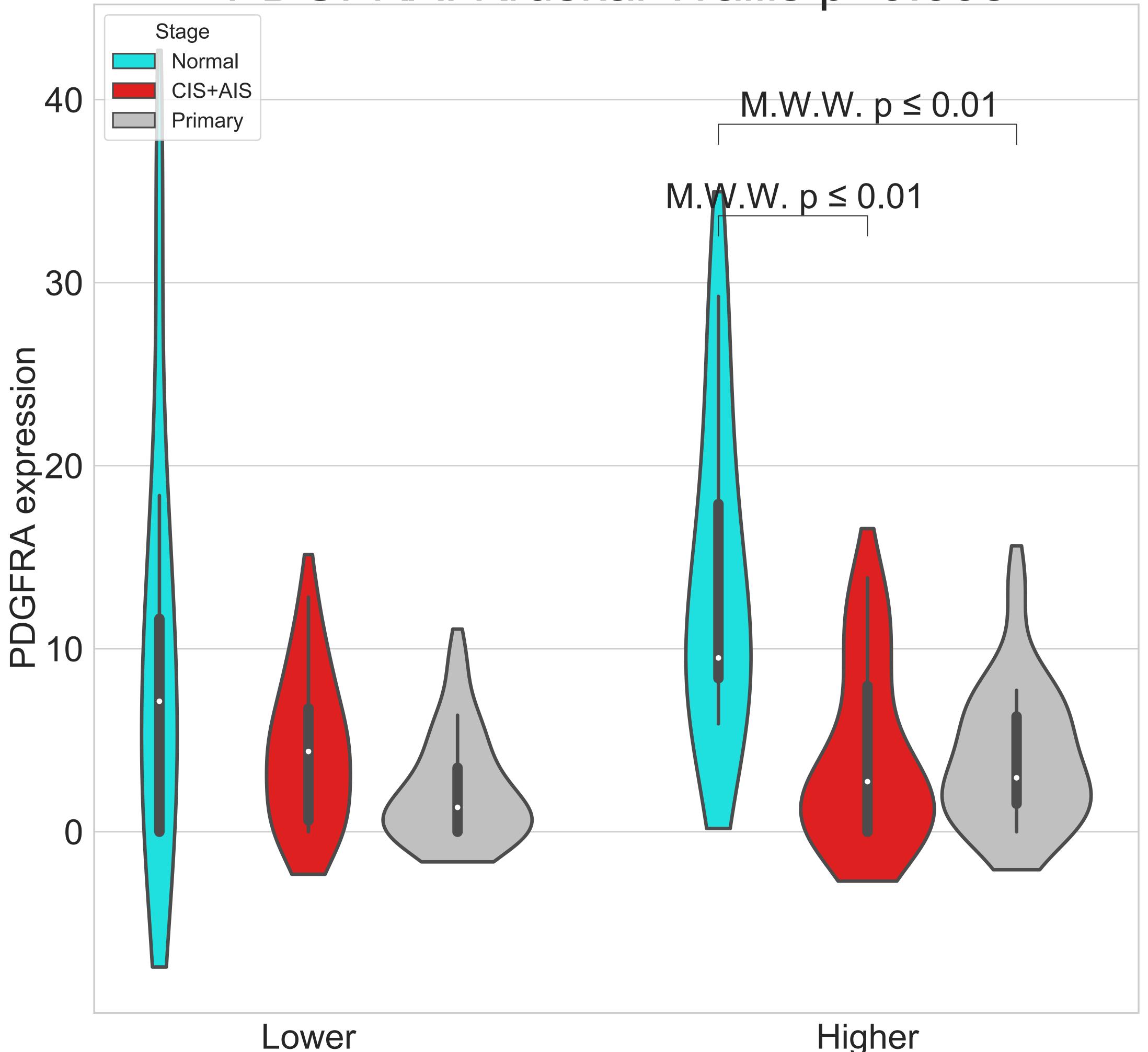
PDGFRA: Kruskal-Wallis p=0.008



Higher Mutation Shared Count (SYN)