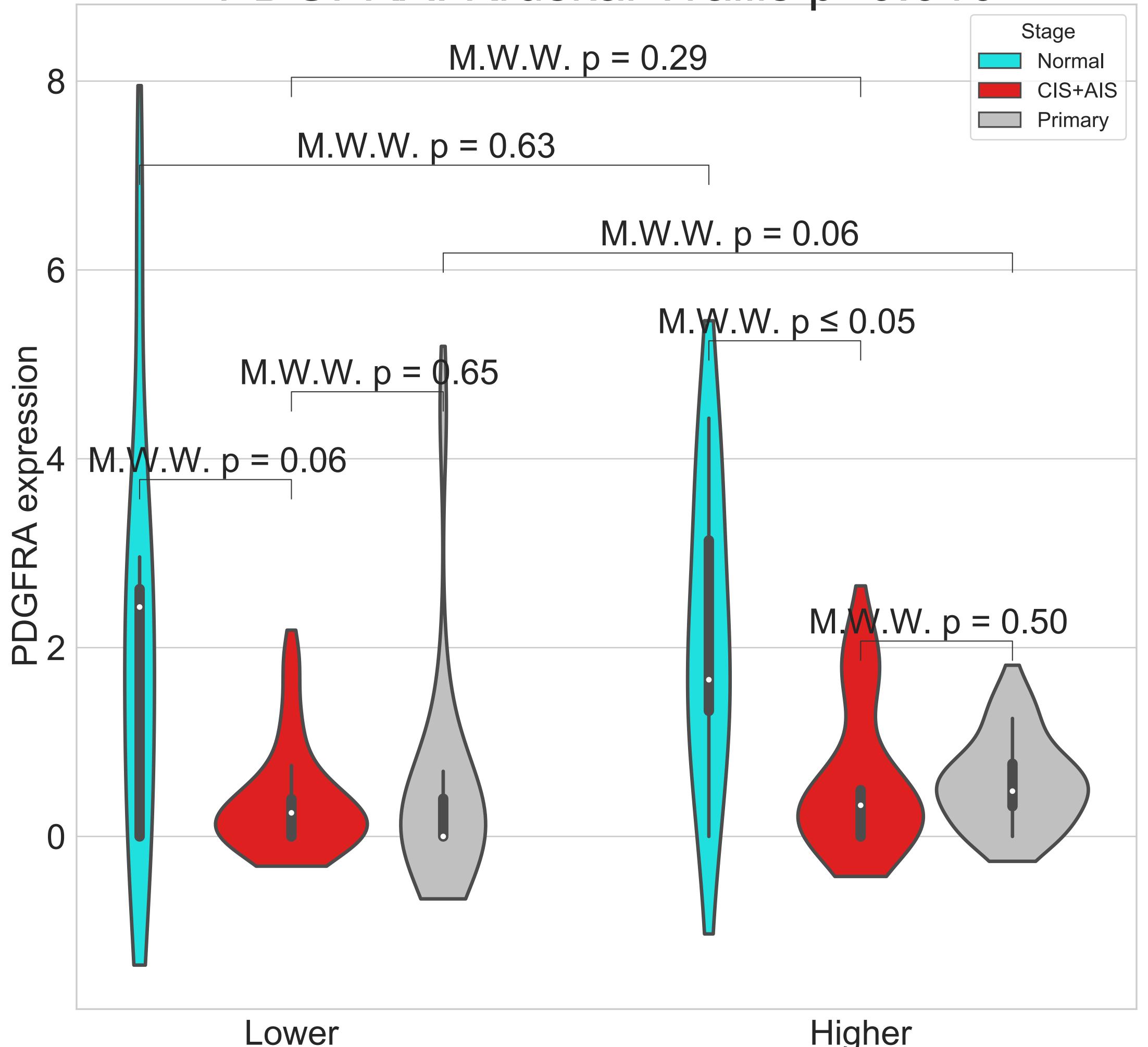
PDGFRA: Kruskal-Wallis p=0.010



Higher Mutation Shared Count (SYN)