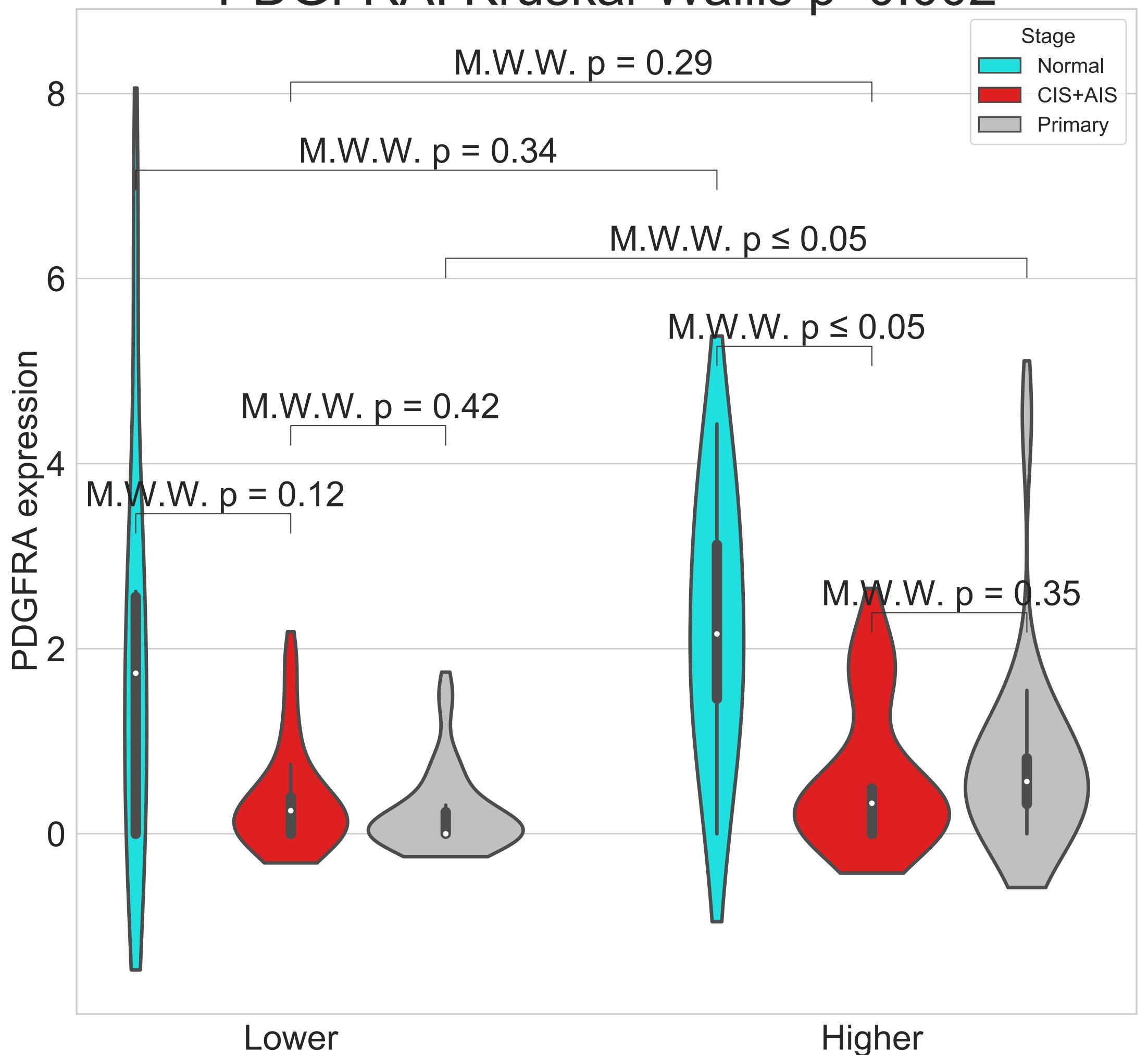
PDGFRA: Kruskal-Wallis p=0.002



Higher Mutation Shared Count (SYN)