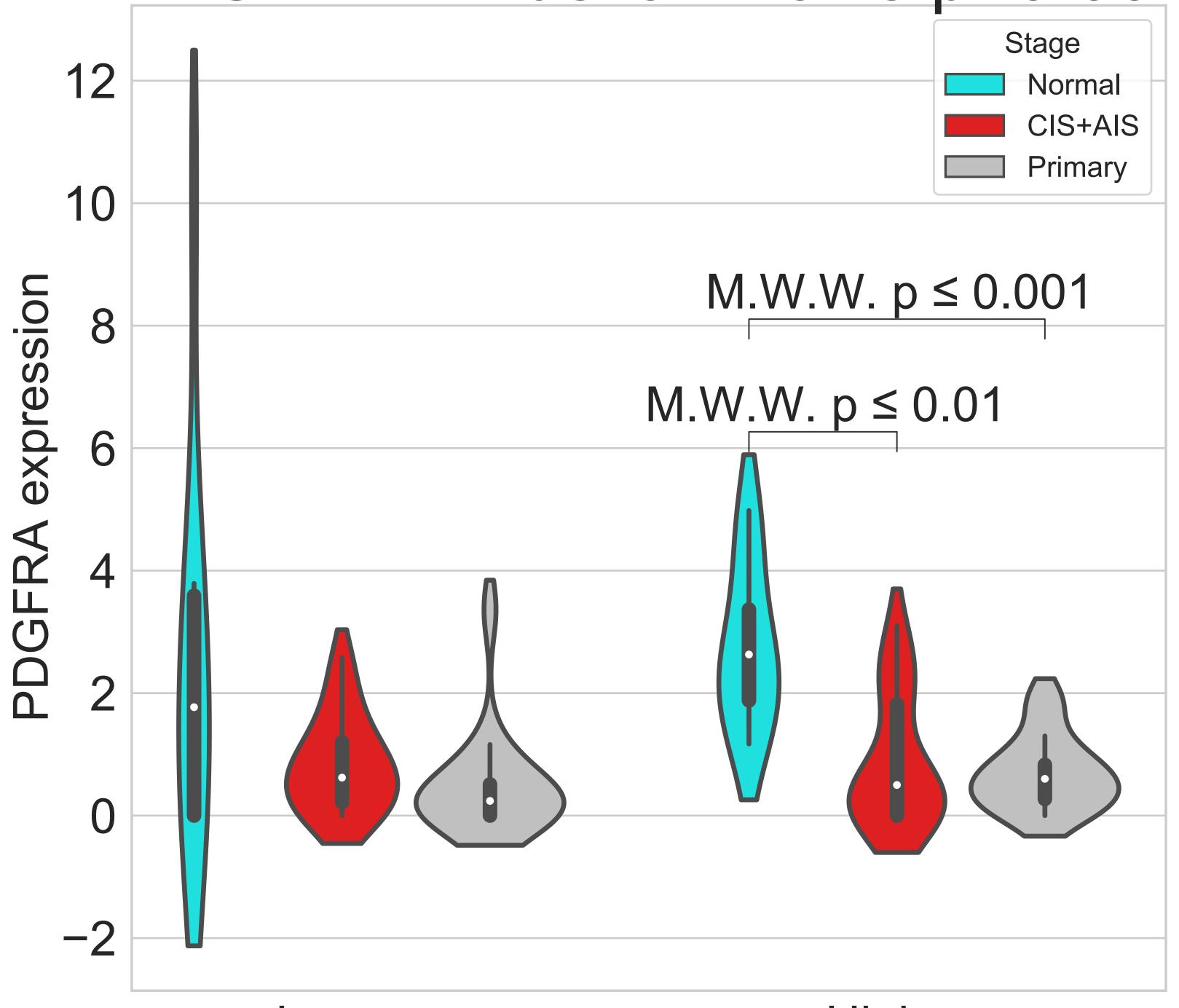
PDGFRA: Kruskal-Wallis p=0.004



Lower

Higher

Mutation Shared Count (SYN)