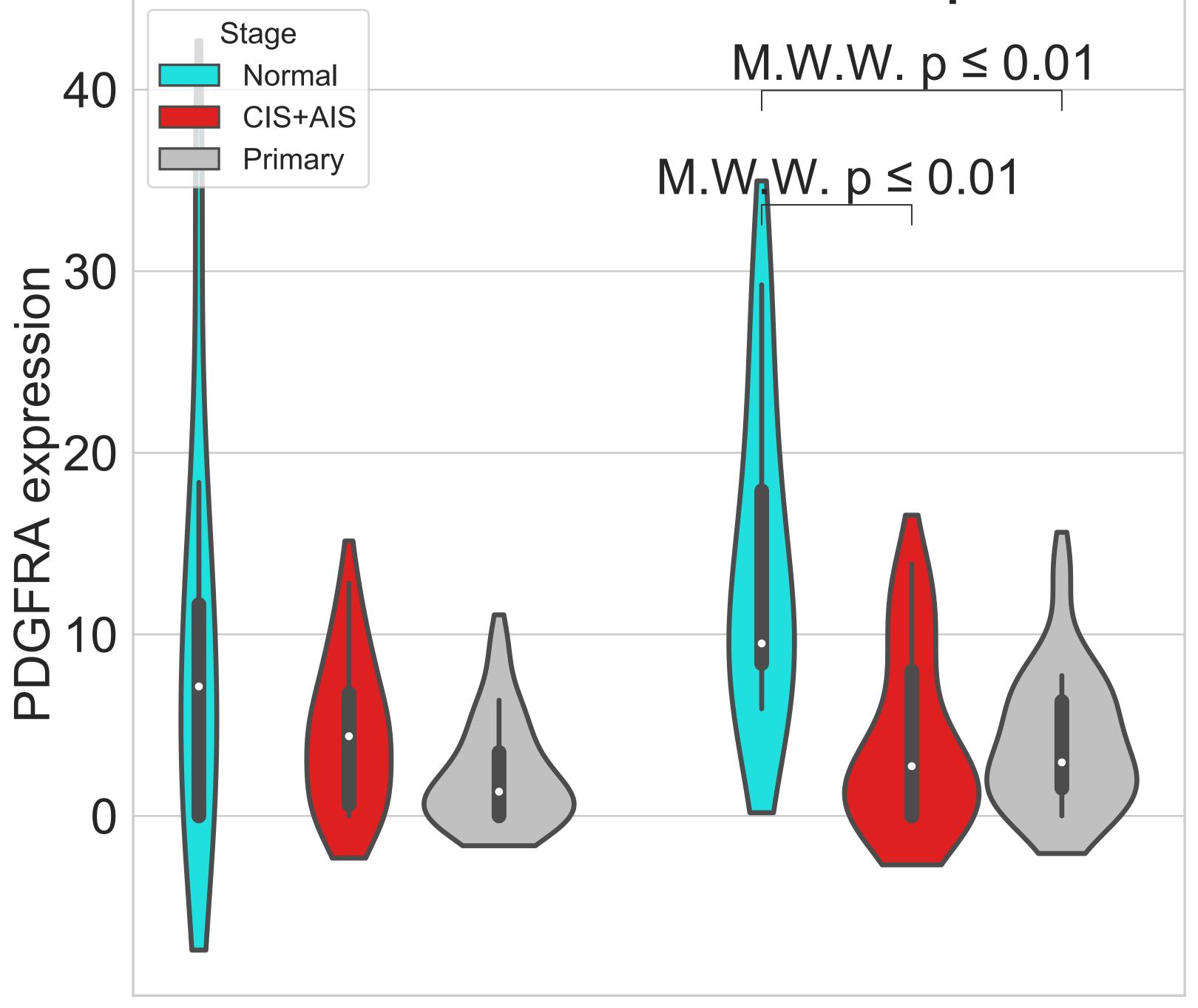
PDGFRA: Kruskal-Wallis p=0.008



Lower
Higher
Mutation Shared Count (SYN)