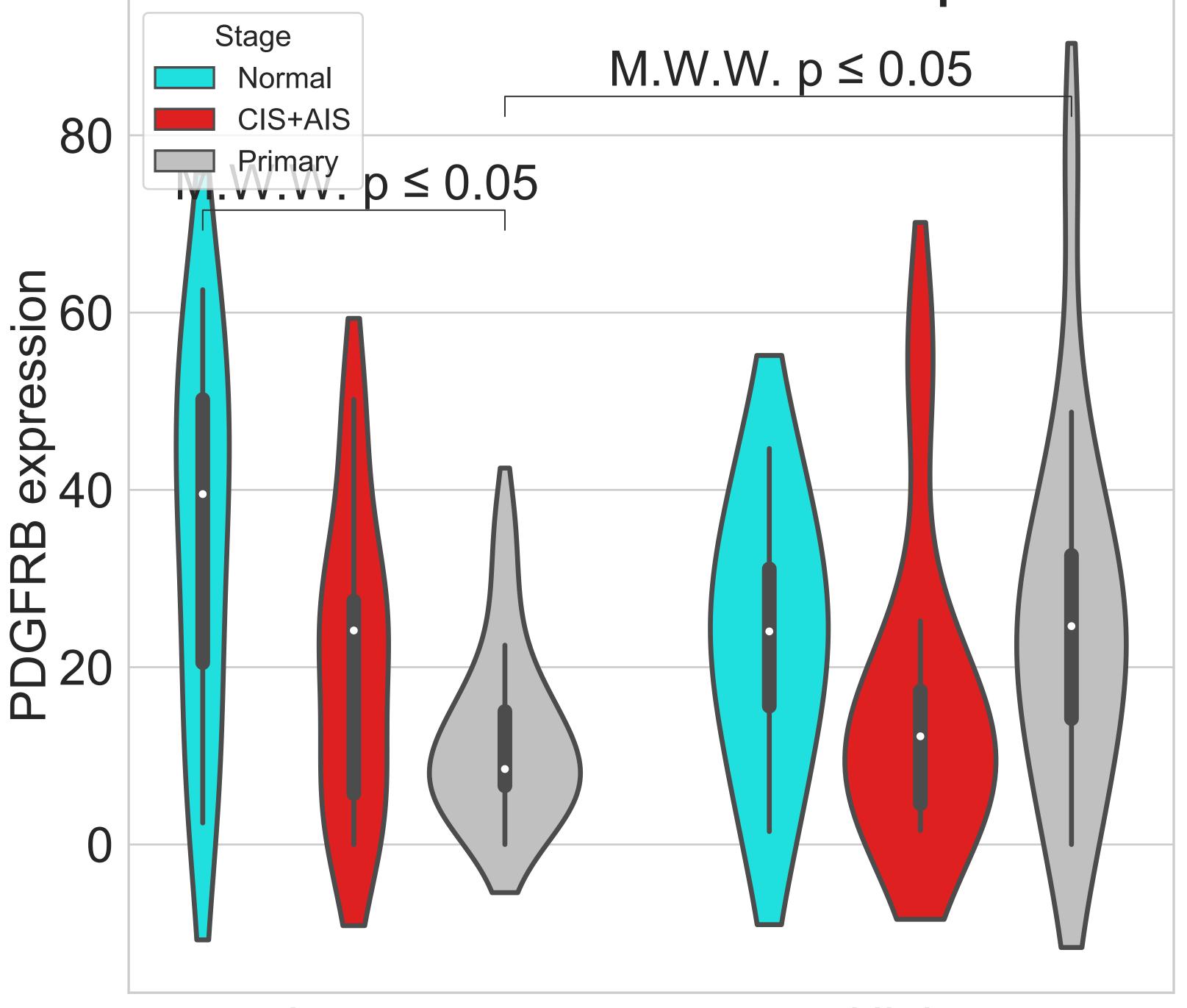
PDGFRB: Kruskal-Wallis p=0.050



Lower
Mutation Shared Proportion (Union & SYN)