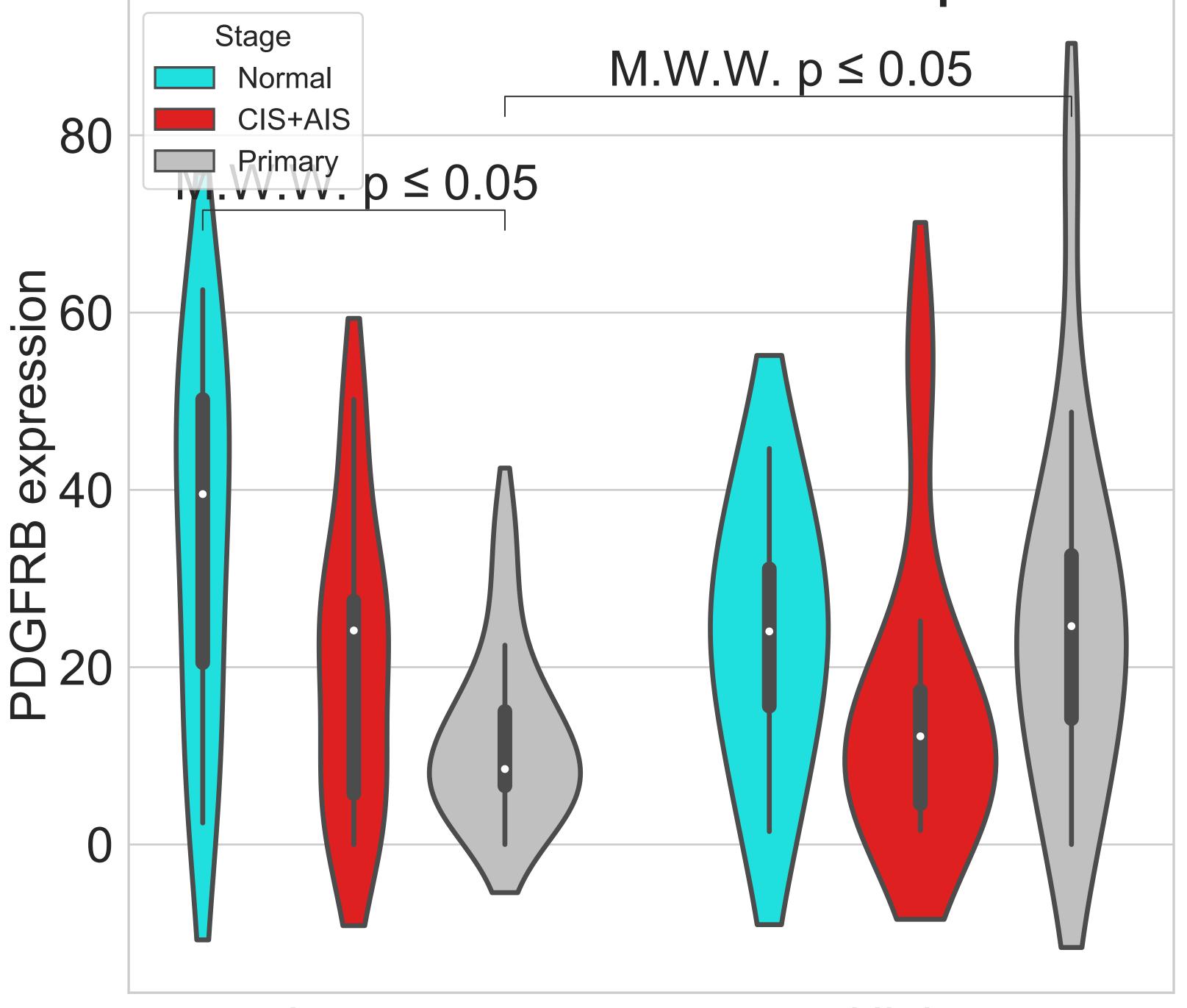
PDGFRB: Kruskal-Wallis p=0.050



Lower
Mutation Shared Count (SYN) per TMB