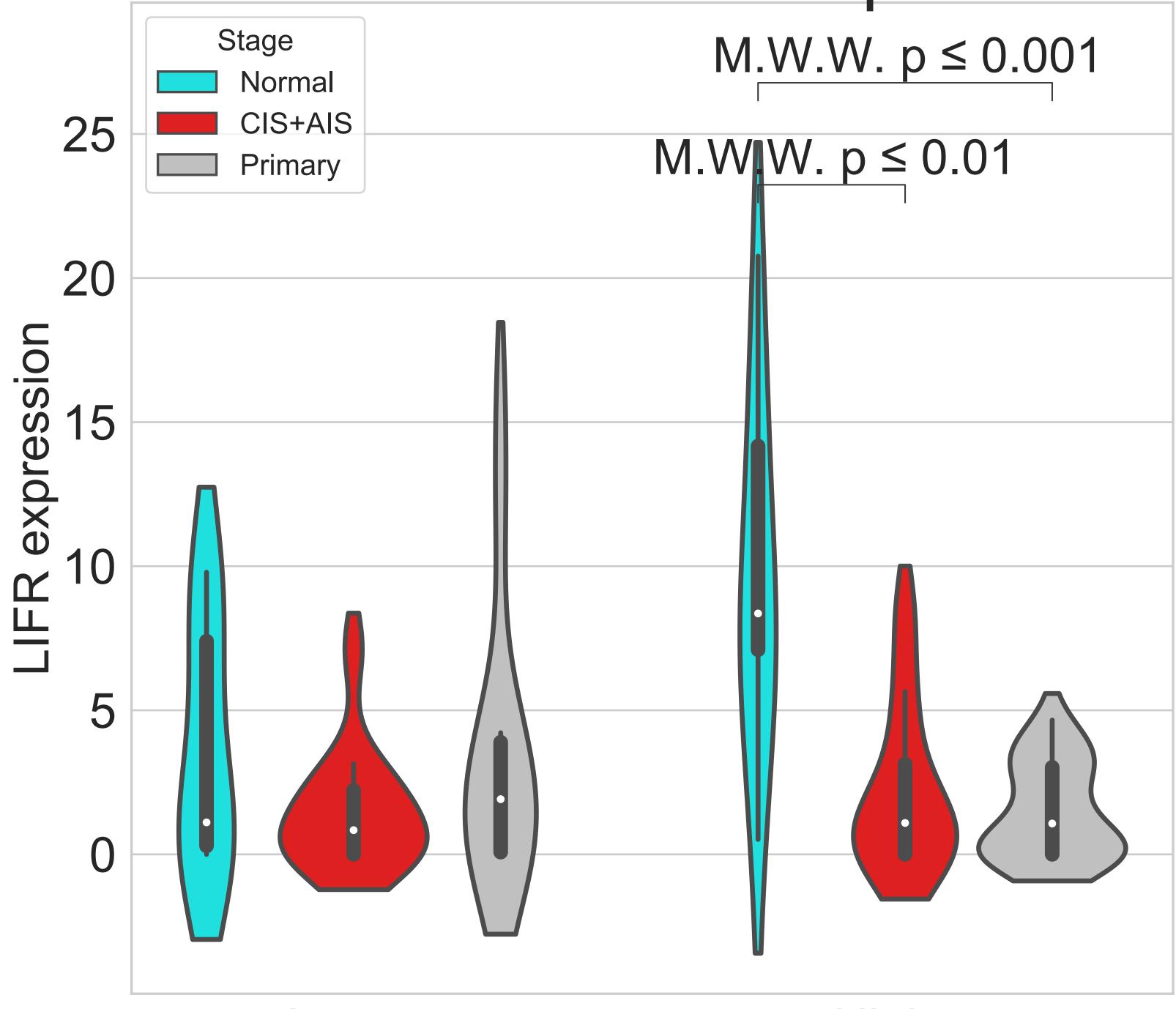
LIFR: Kruskal-Wallis p=0.006



Lower
Mutation Shared Count (SYN) per TMB