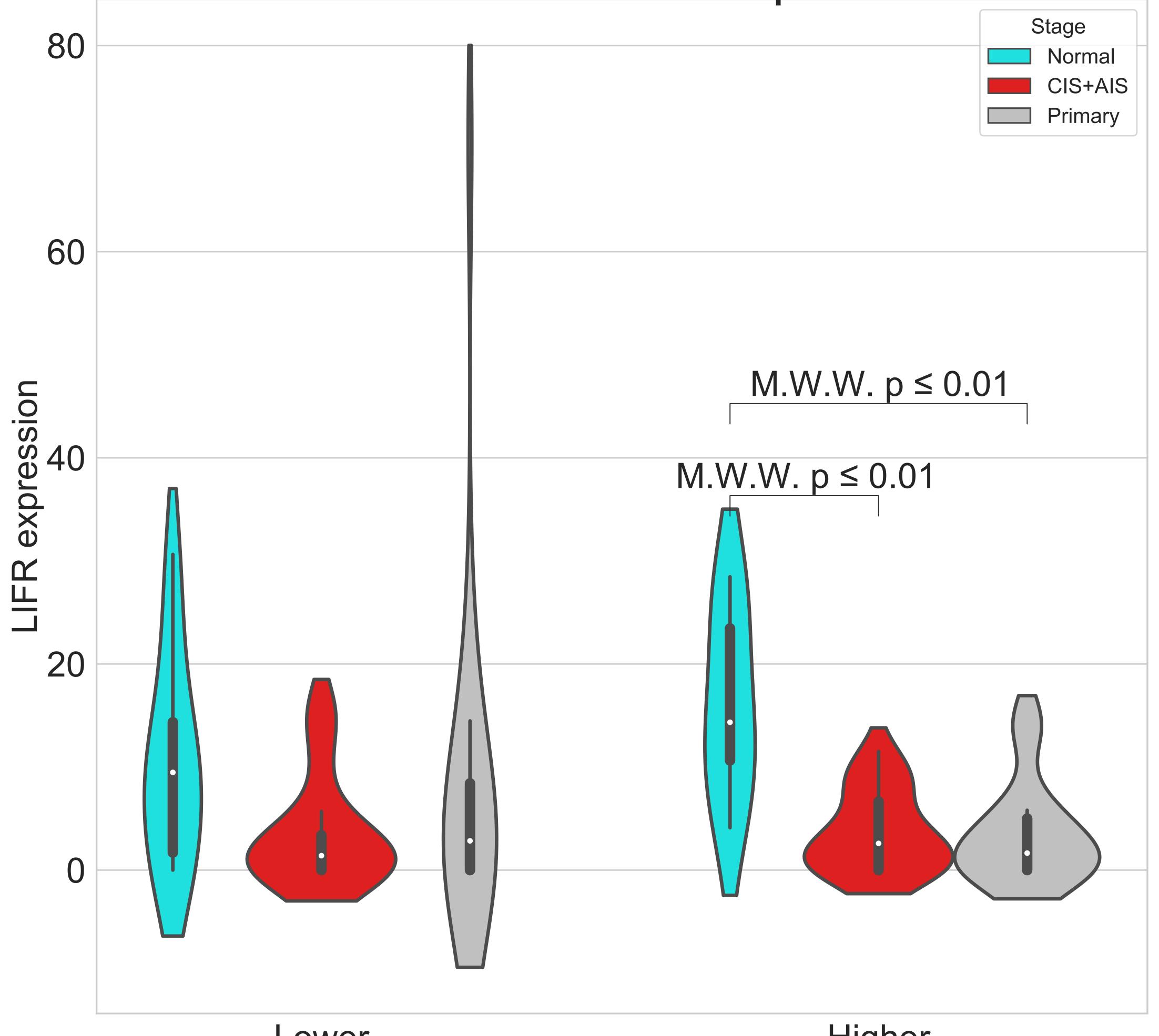
LIFR: Kruskal-Wallis p=0.021



Lower
Mutation Shared Count (SYN) per TMB