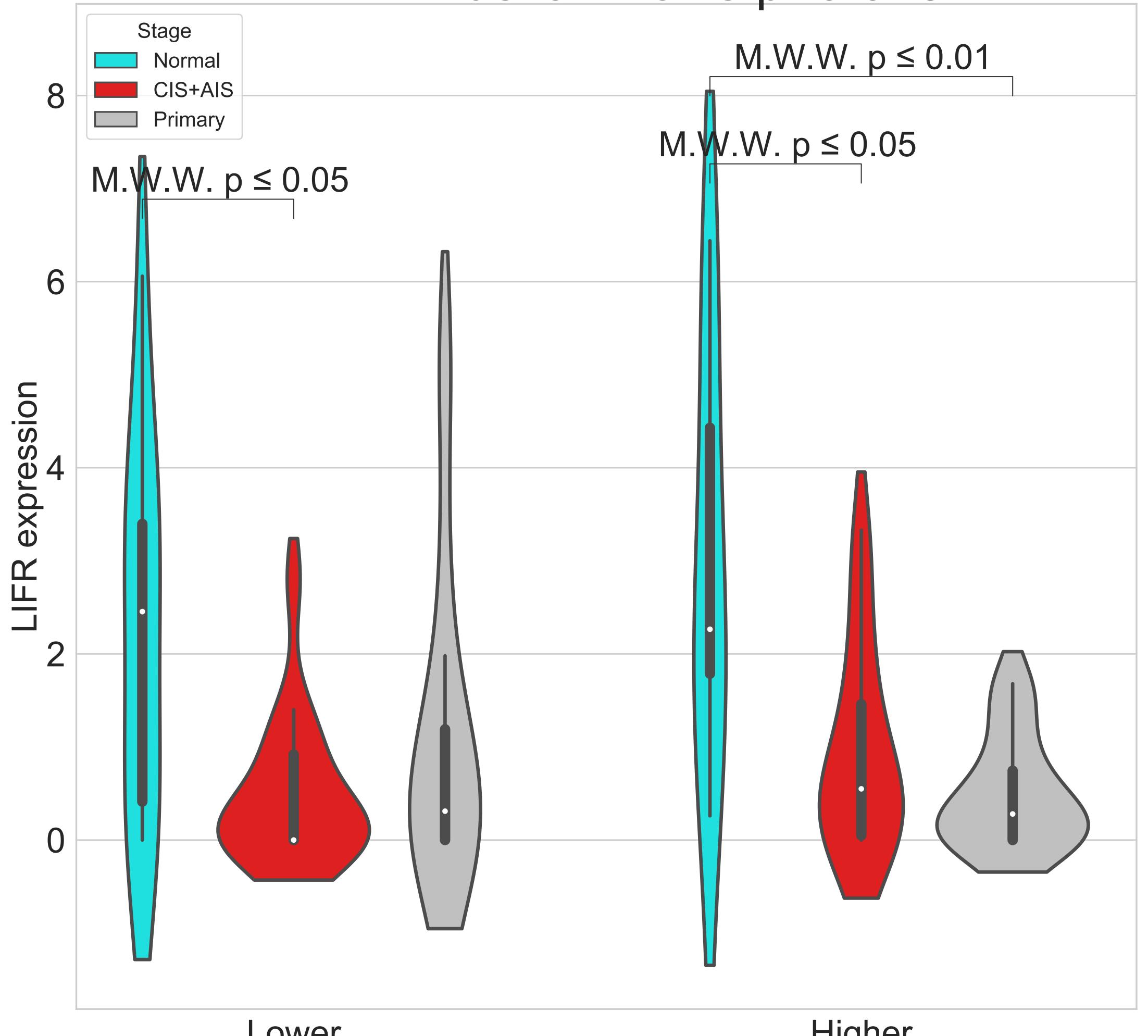
LIFR: Kruskal-Wallis p=0.013



Lower
Mutation Shared Count (SYN) per TMB