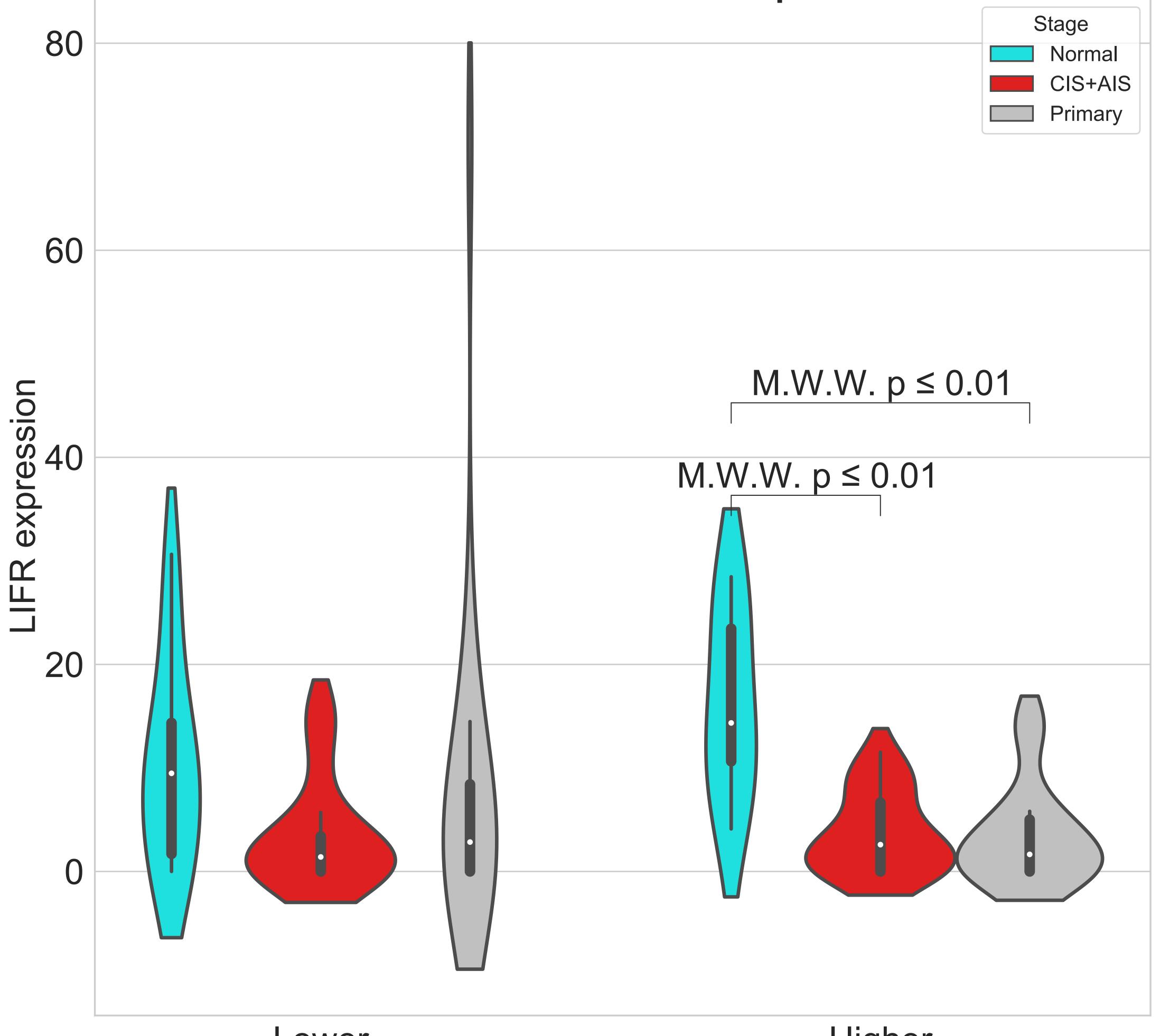
LIFR: Kruskal-Wallis p=0.021



Lower
Mutation Shared Proportion (Union & SYN)