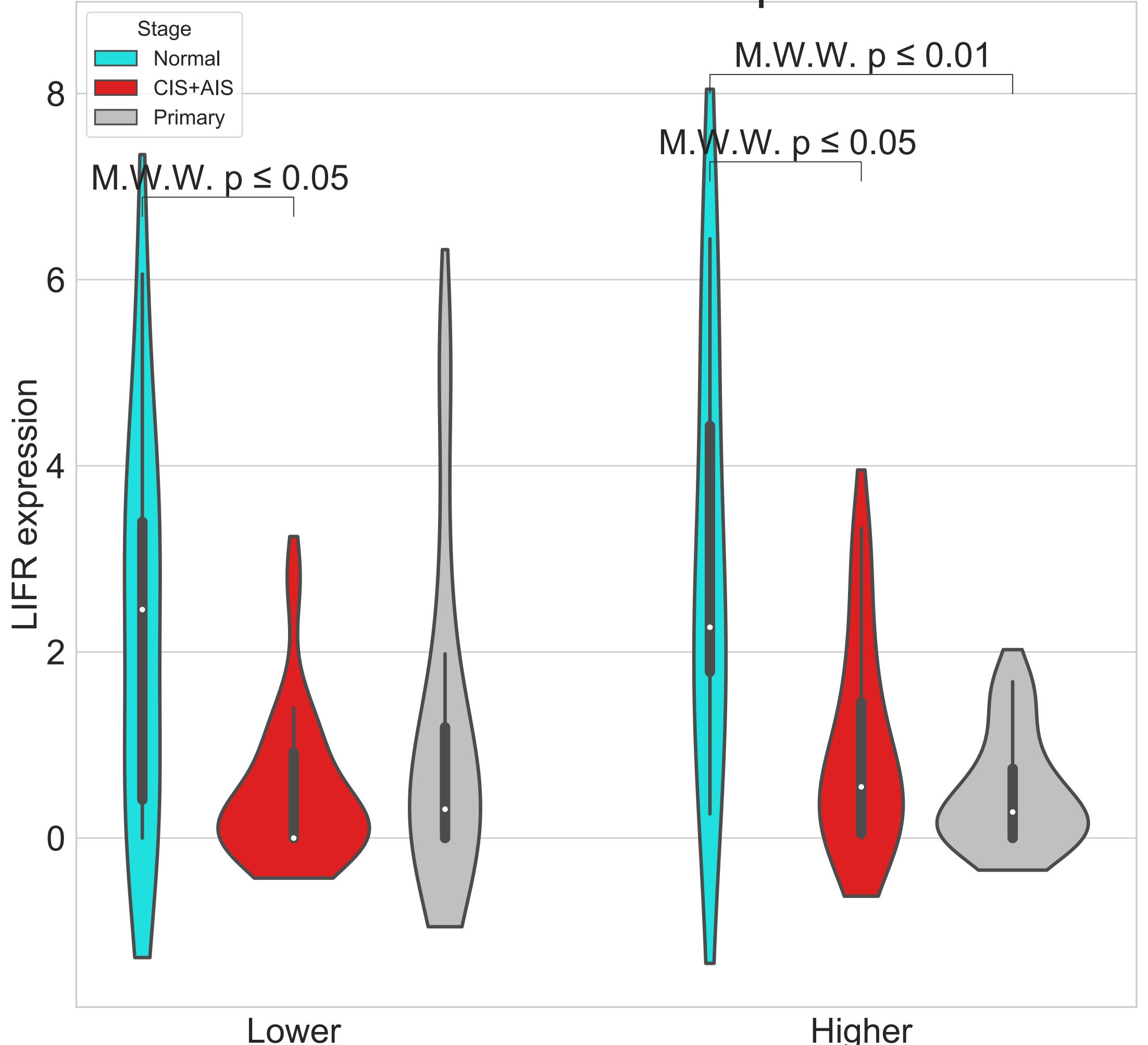
LIFR: Kruskal-Wallis p=0.013



Higher Mutation Shared Proportion (SYN)