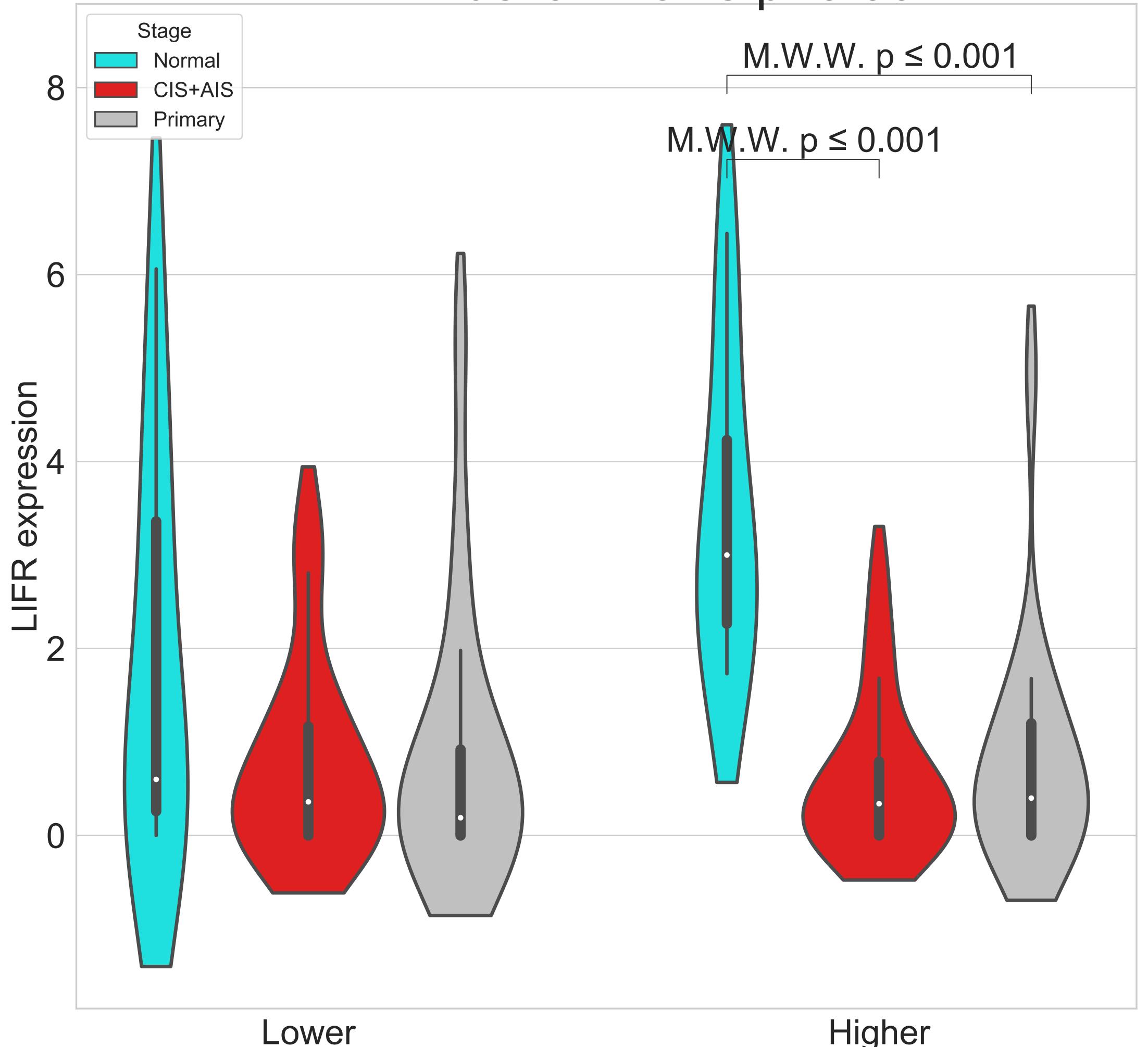
LIFR: Kruskal-Wallis p=0.007



Higher Mutation Shared Count (SYN)