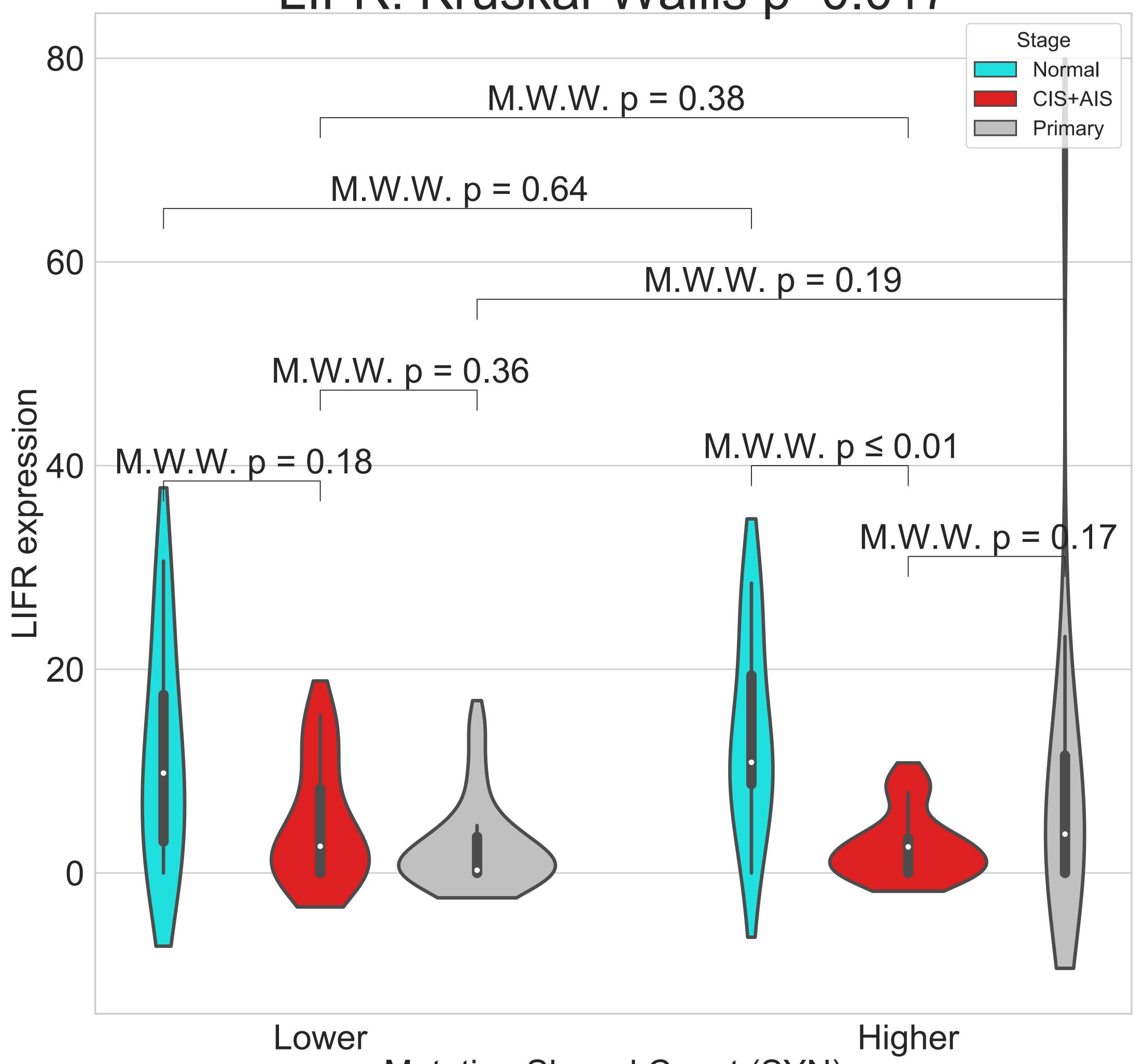
LIFR: Kruskal-Wallis p=0.017



Mutation Shared Count (SYN)