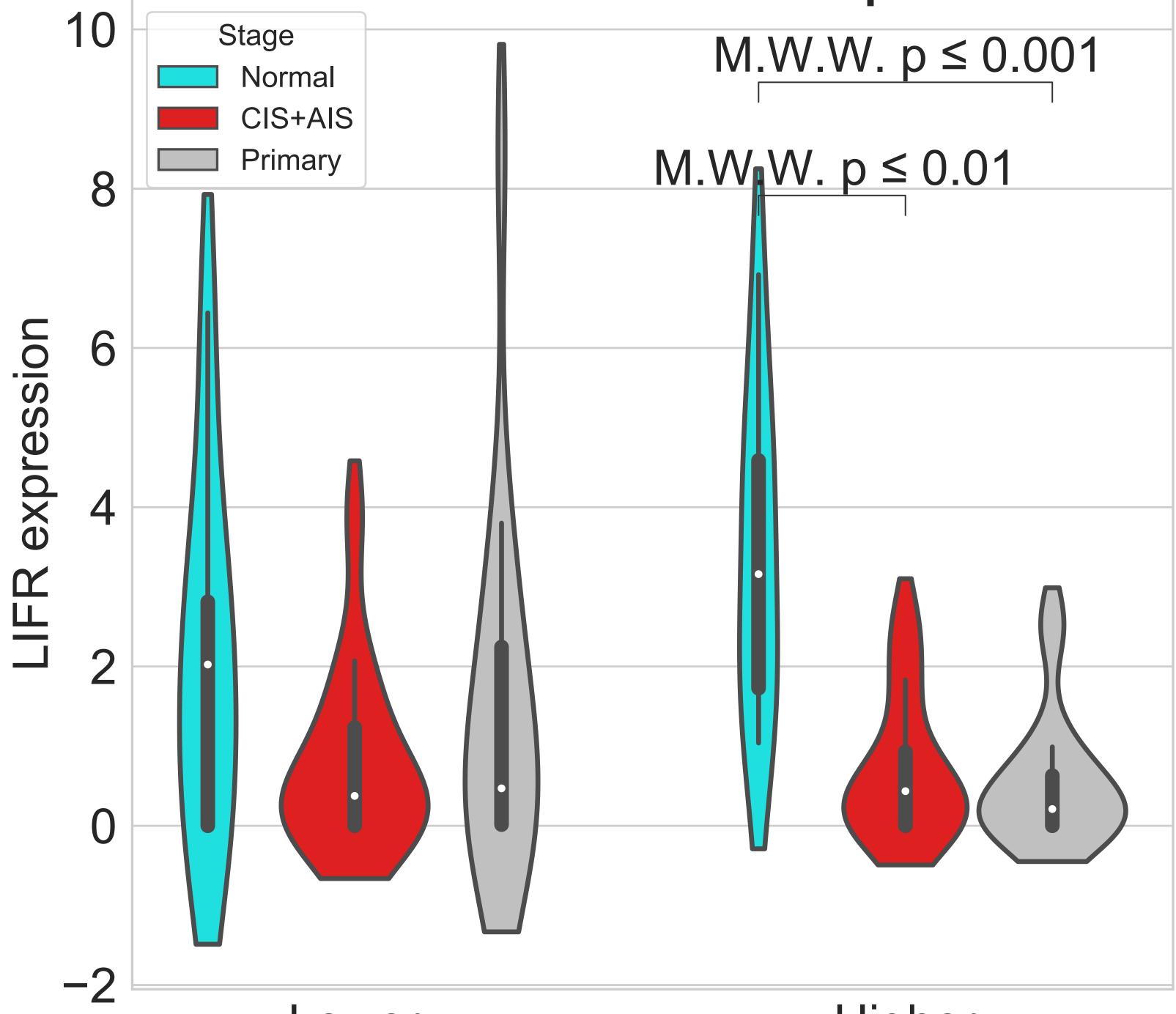
LIFR: Kruskal-Wallis p=0.007



Lower
Mutation Shared Proportion (Union & SYN)