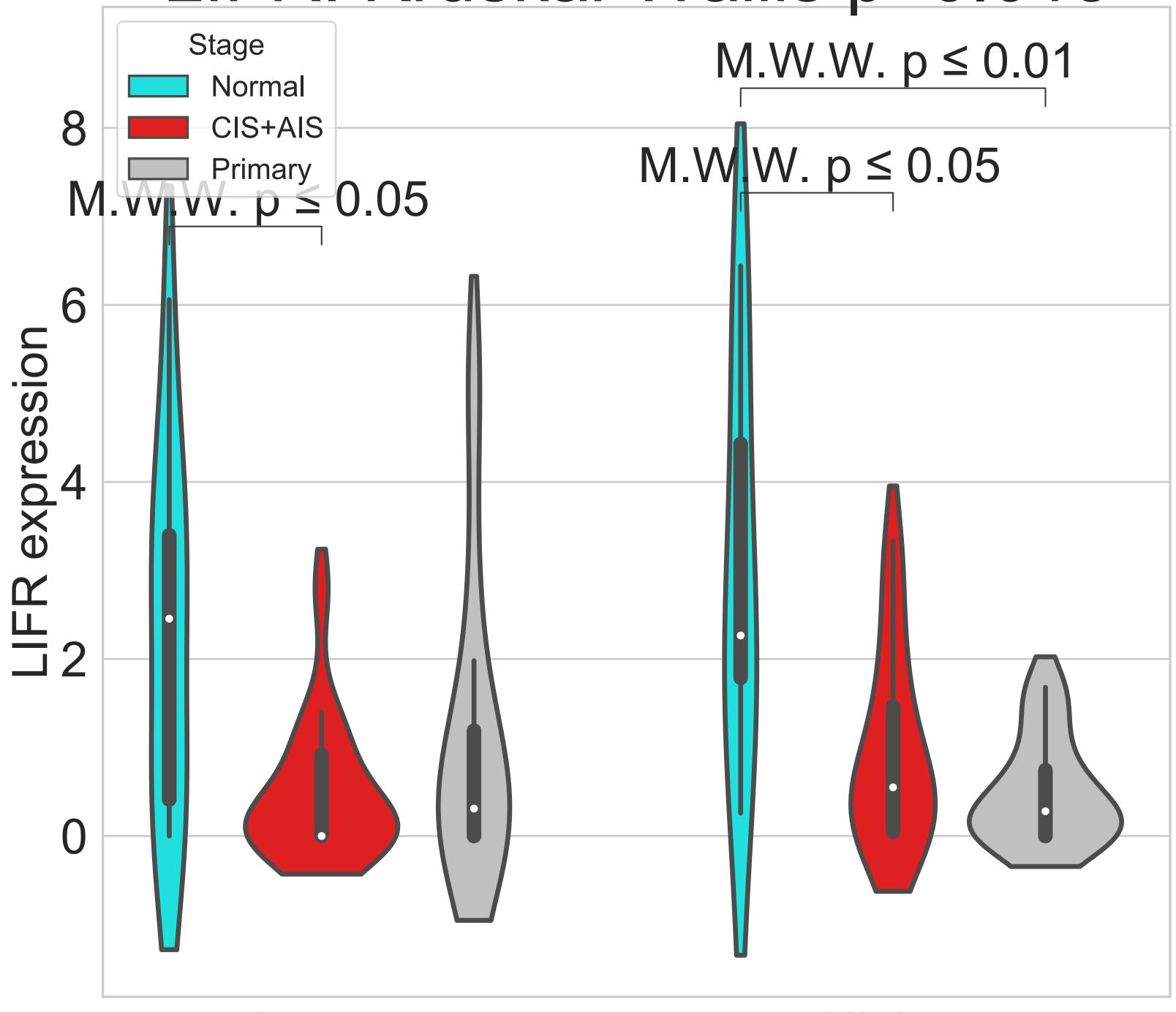
LIFR: Kruskal-Wallis p=0.013



Lower
Mutation Shared Proportion (Union & SYN)