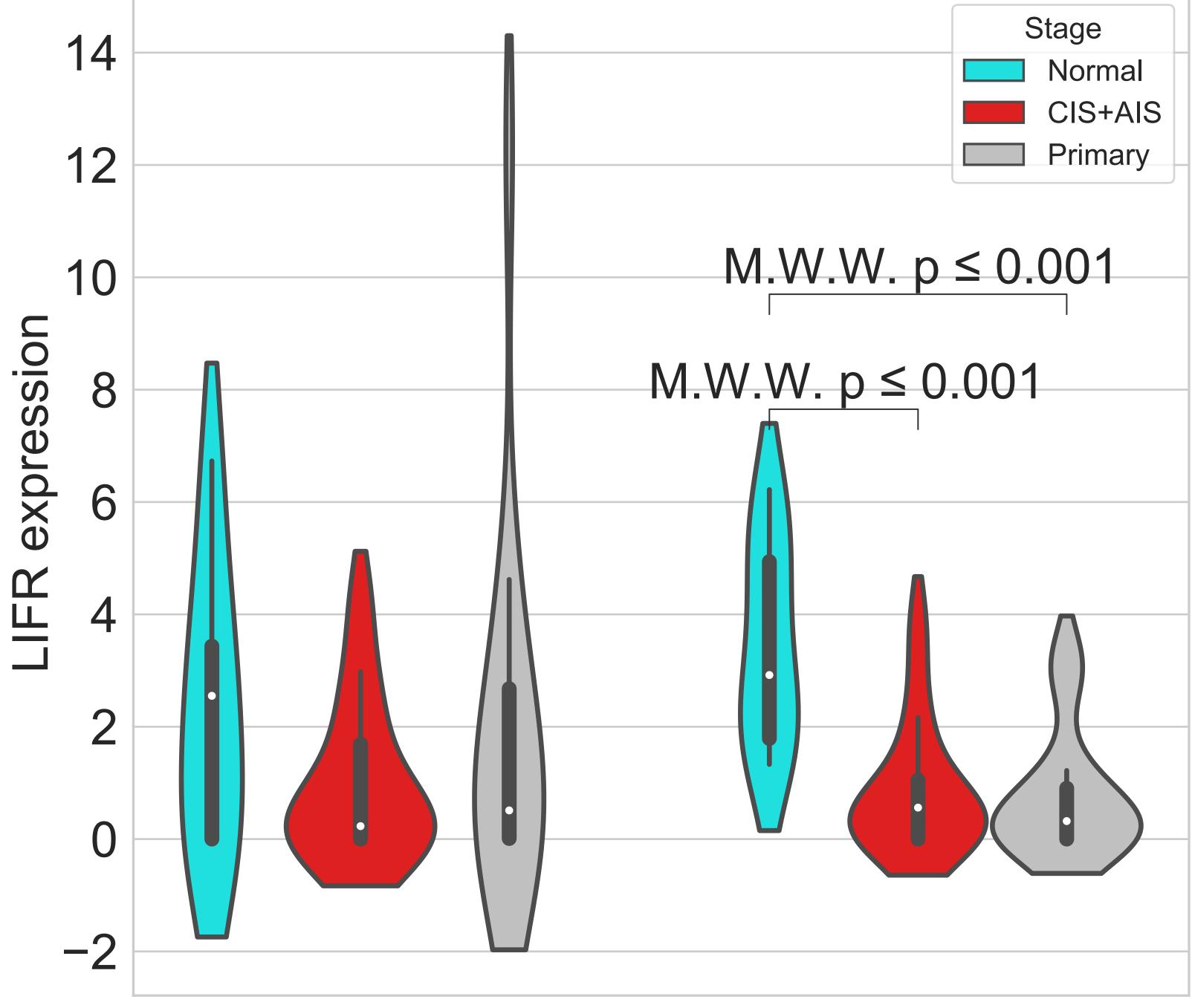
LIFR: Kruskal-Wallis p=0.013

Stage
Normal



Lower Higher Mutation Shared Proportion (SYN)