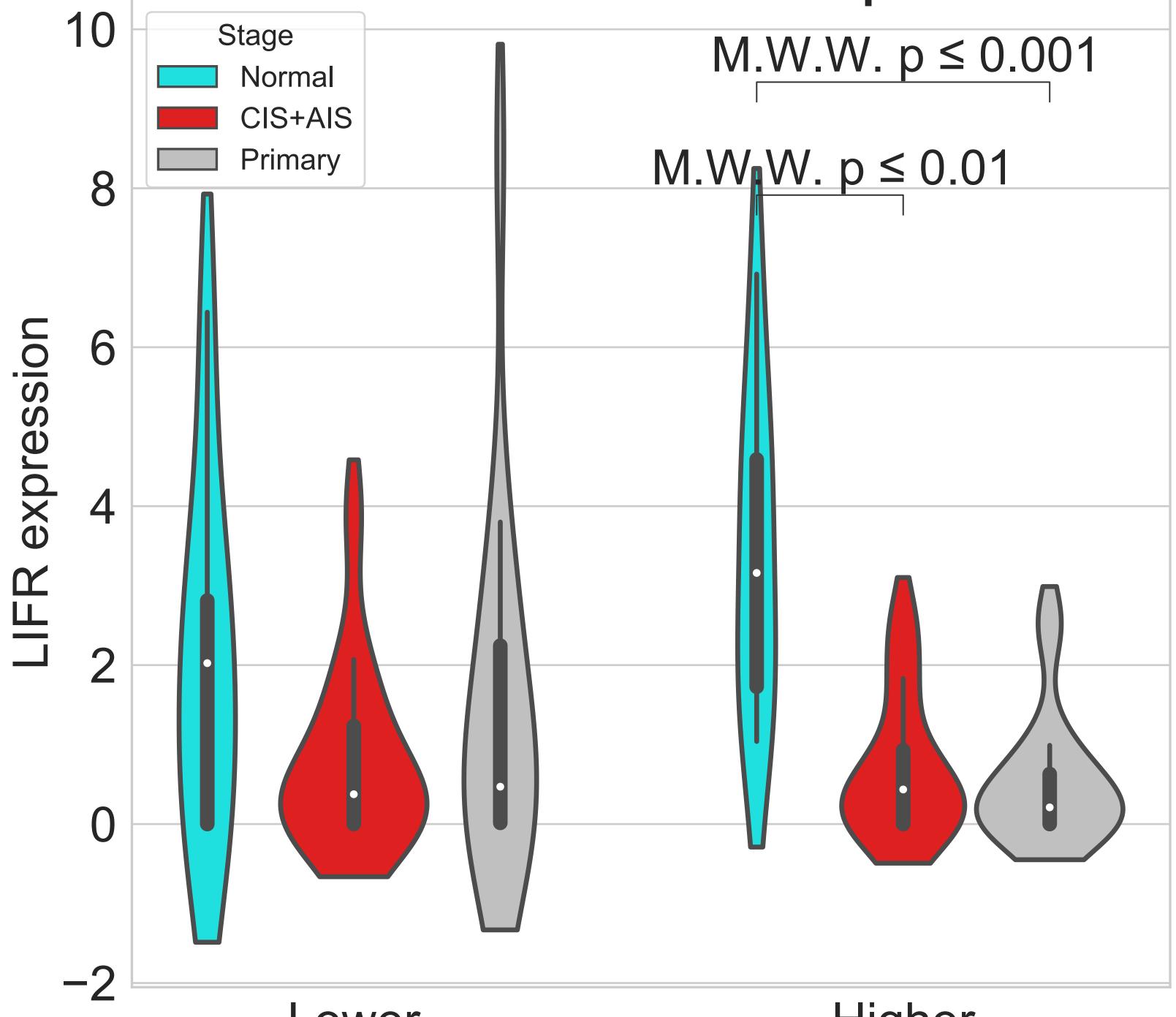
LIFR: Kruskal-Wallis p=0.007



Lower Higher Mutation Shared Proportion (Union)