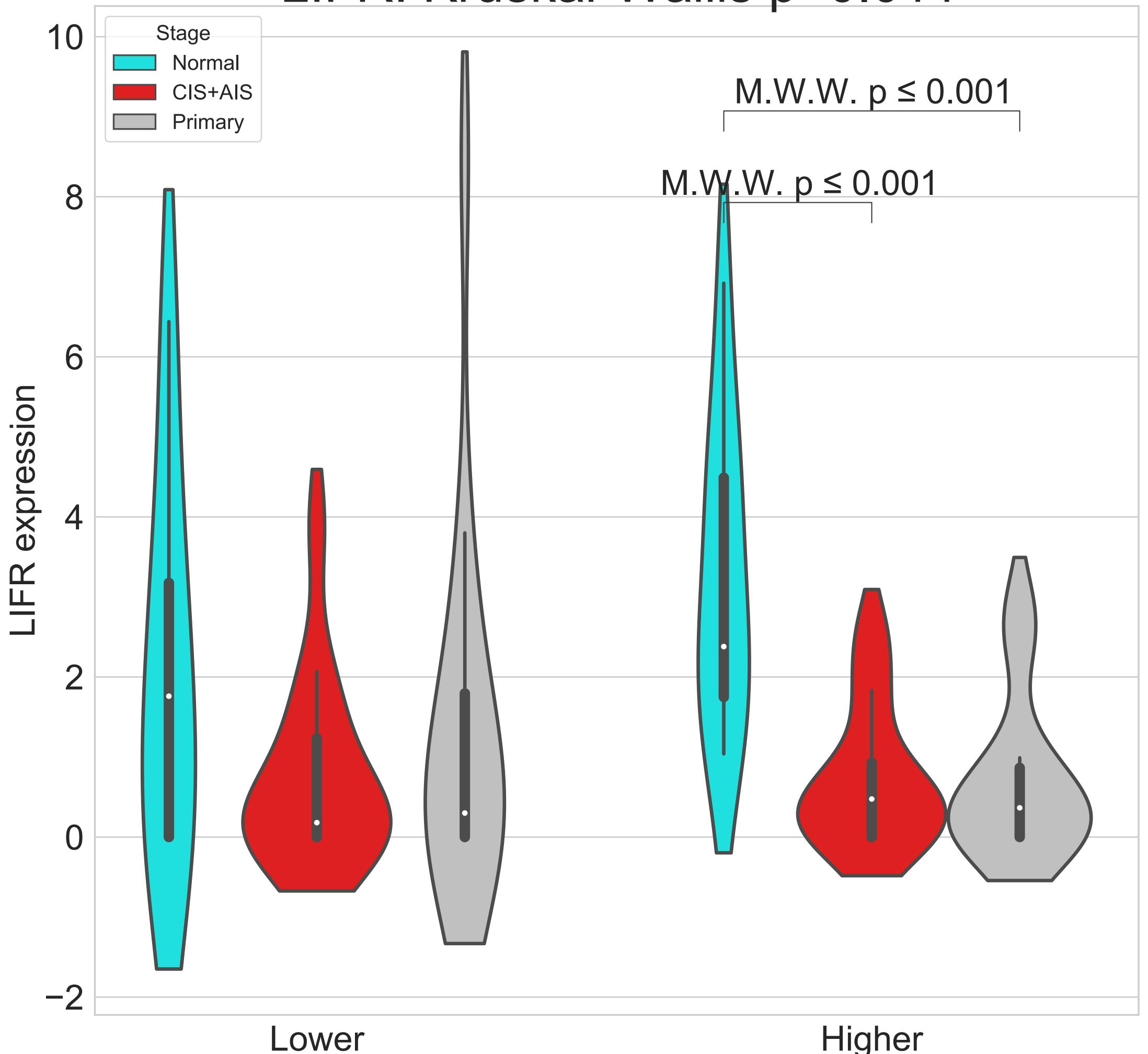
LIFR: Kruskal-Wallis p=0.011



Higher Mutation Shared Proportion