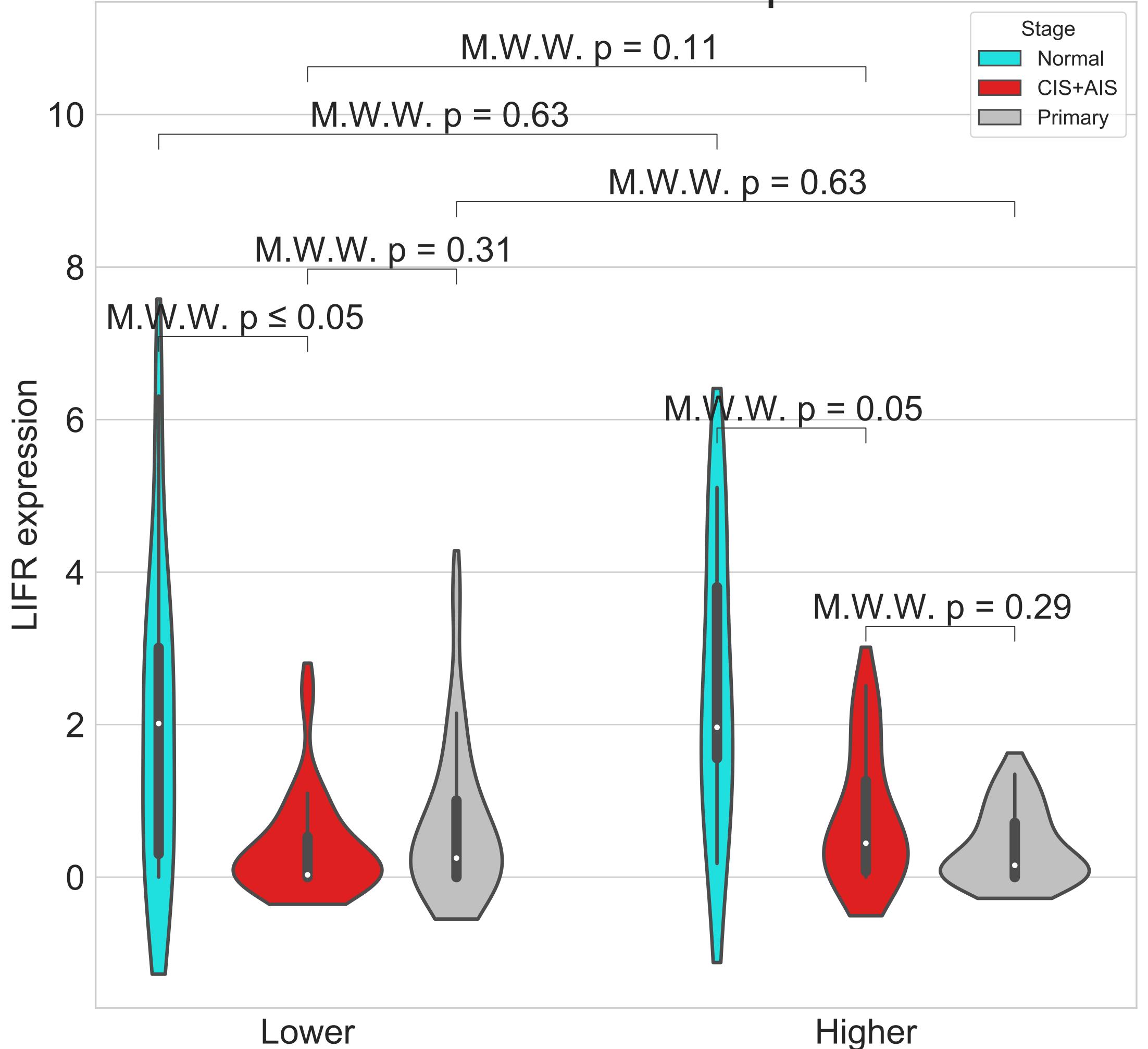
LIFR: Kruskal-Wallis p=0.008



Higher Mutation Shared Proportion