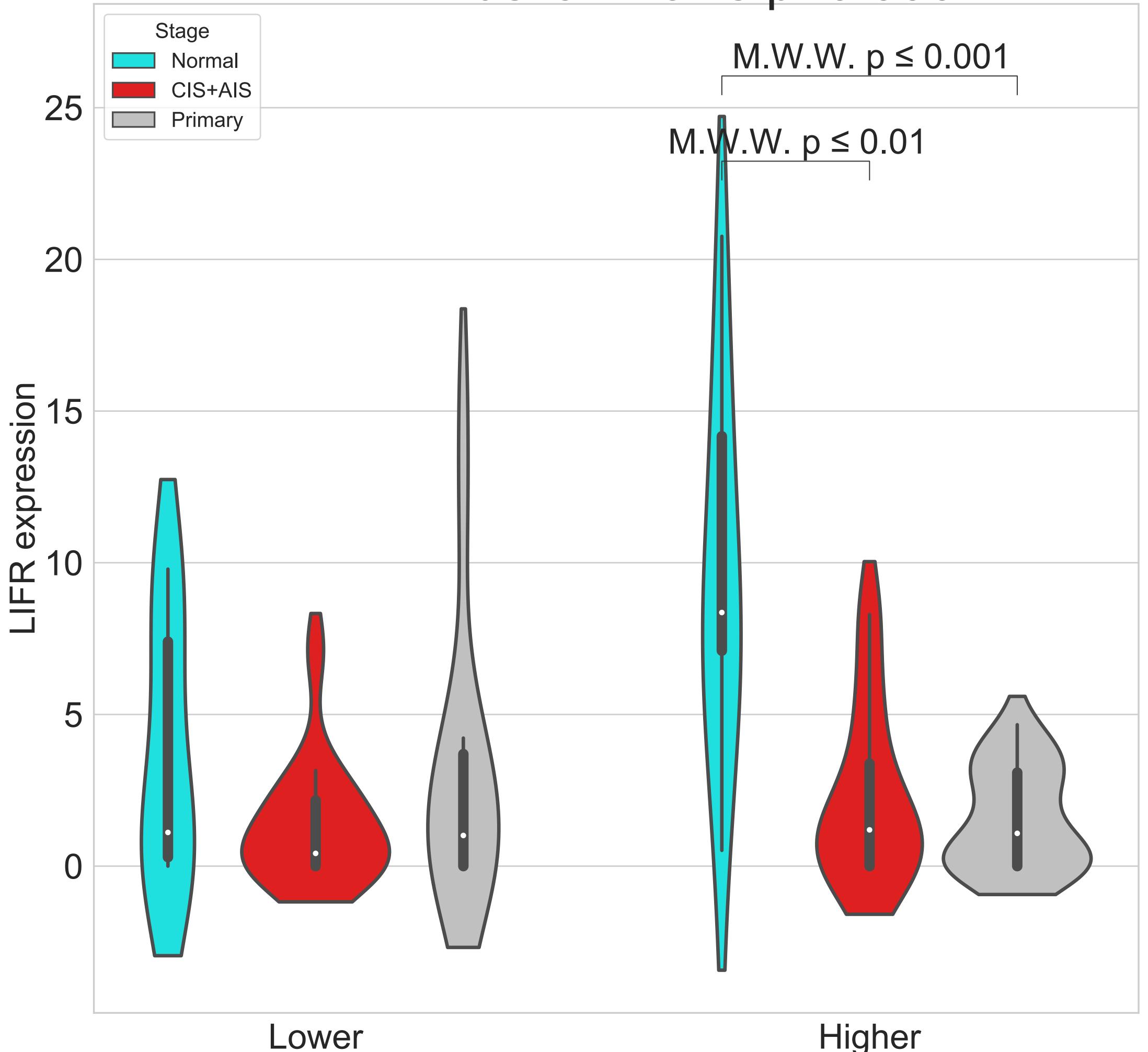
LIFR: Kruskal-Wallis p=0.006



Higher Mutation Shared Proportion