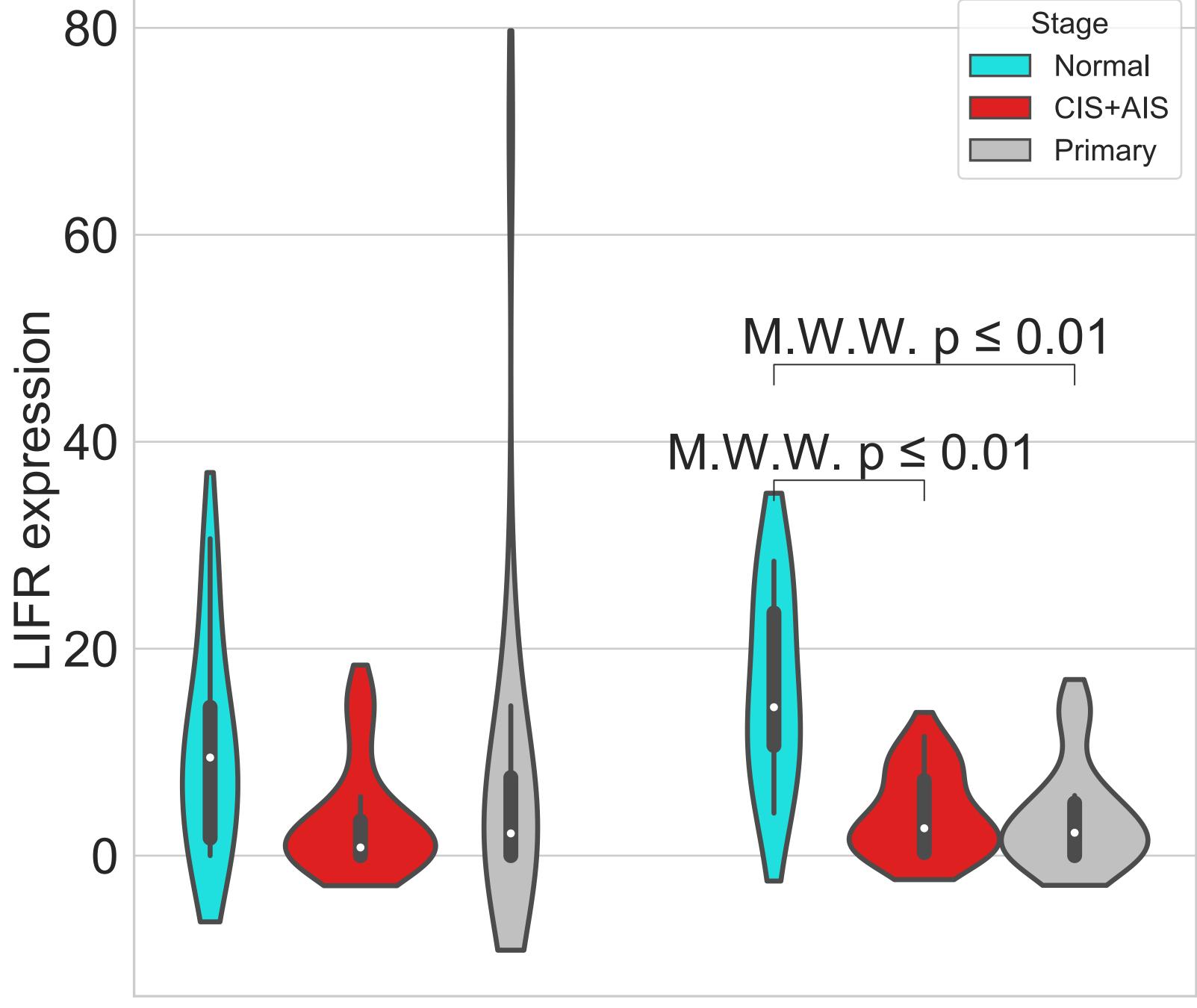
LIFR: Kruskal-Wallis p=0.022 Stage Normal



Higher Lower Mutation Shared Count per TMB