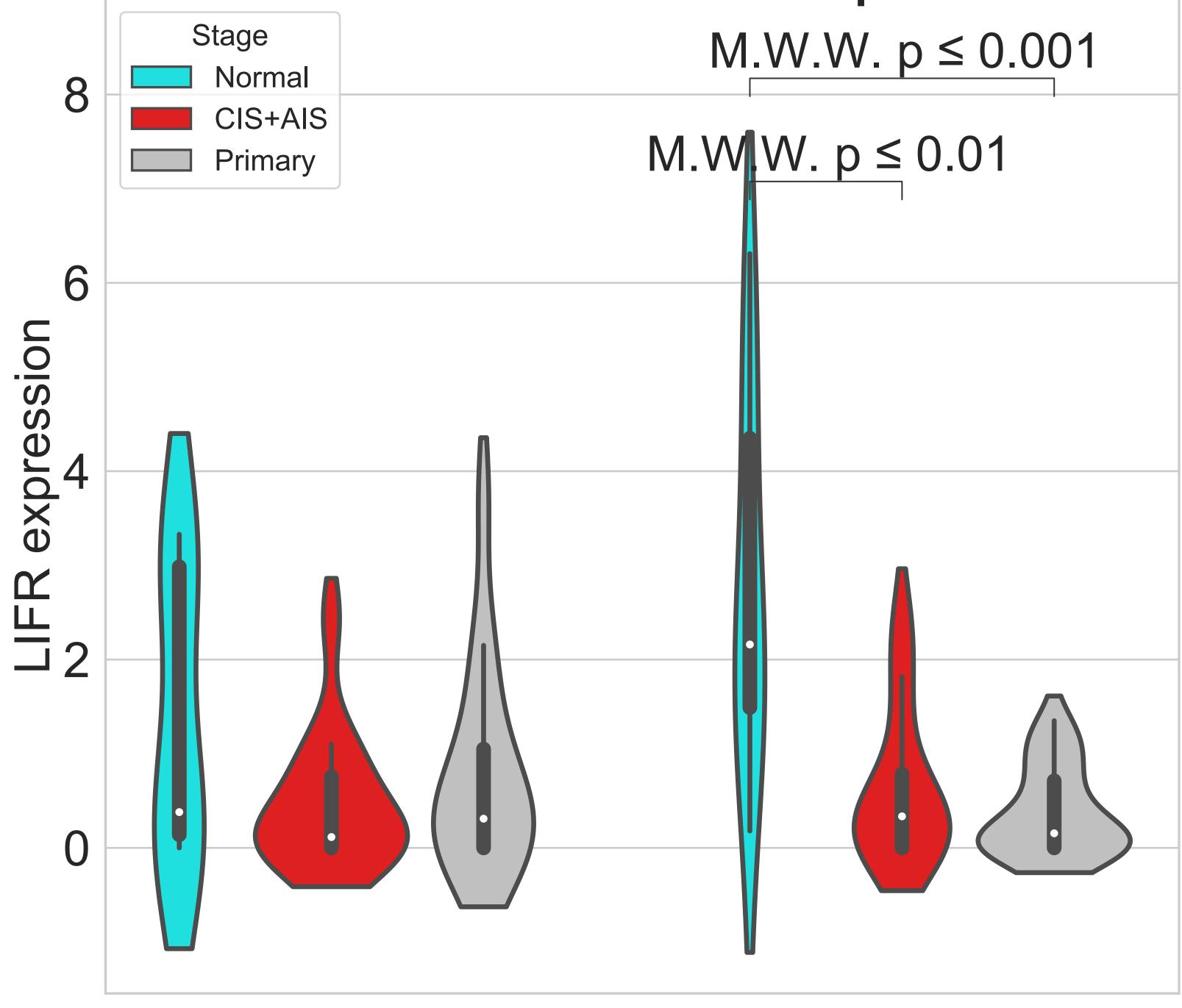
LIFR: Kruskal-Wallis p=0.005



Lower Higher Mutation Shared Count per TMB