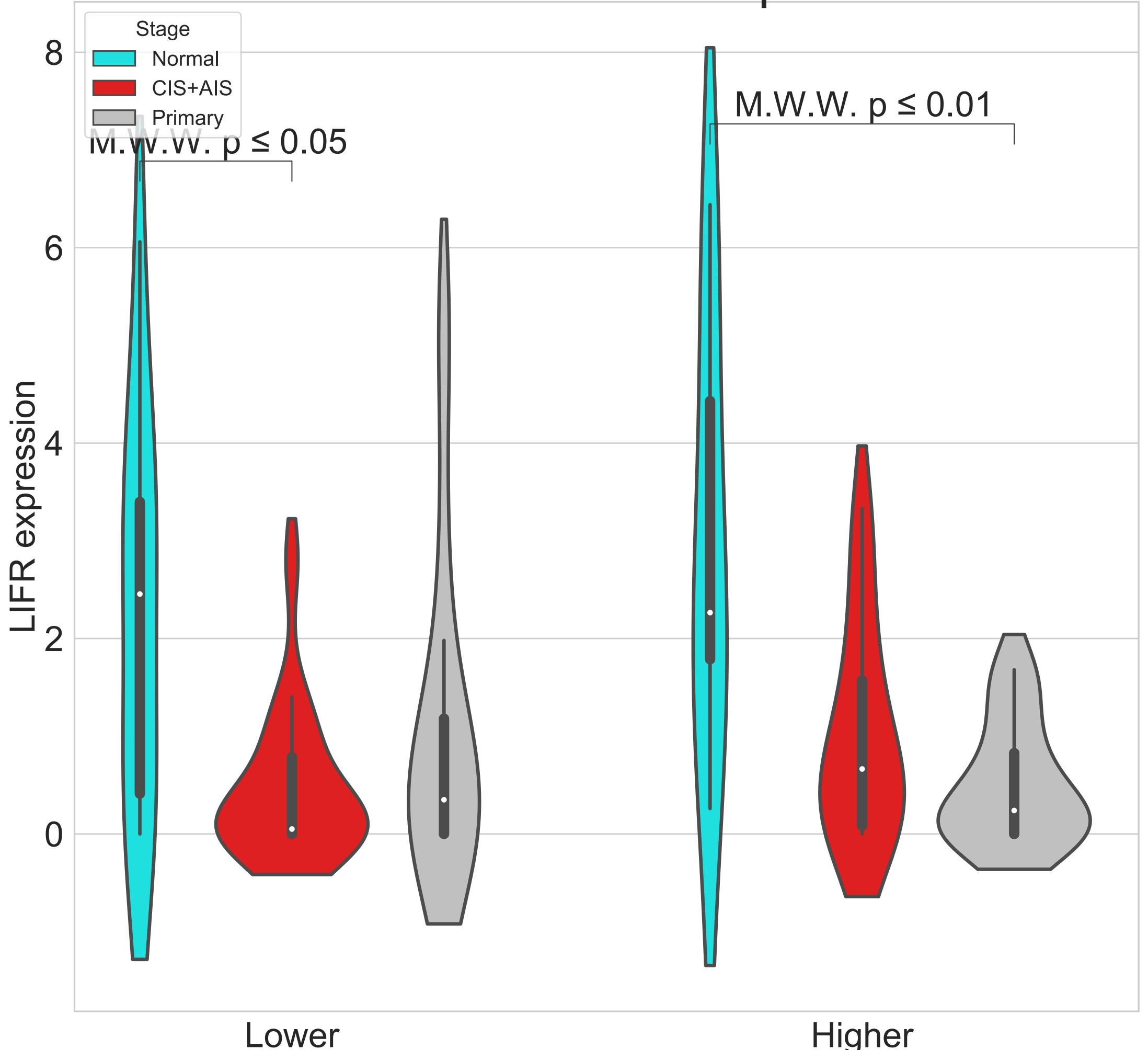
LIFR: Kruskal-Wallis p=0.011



wer
Mutation Shared Count per TMB