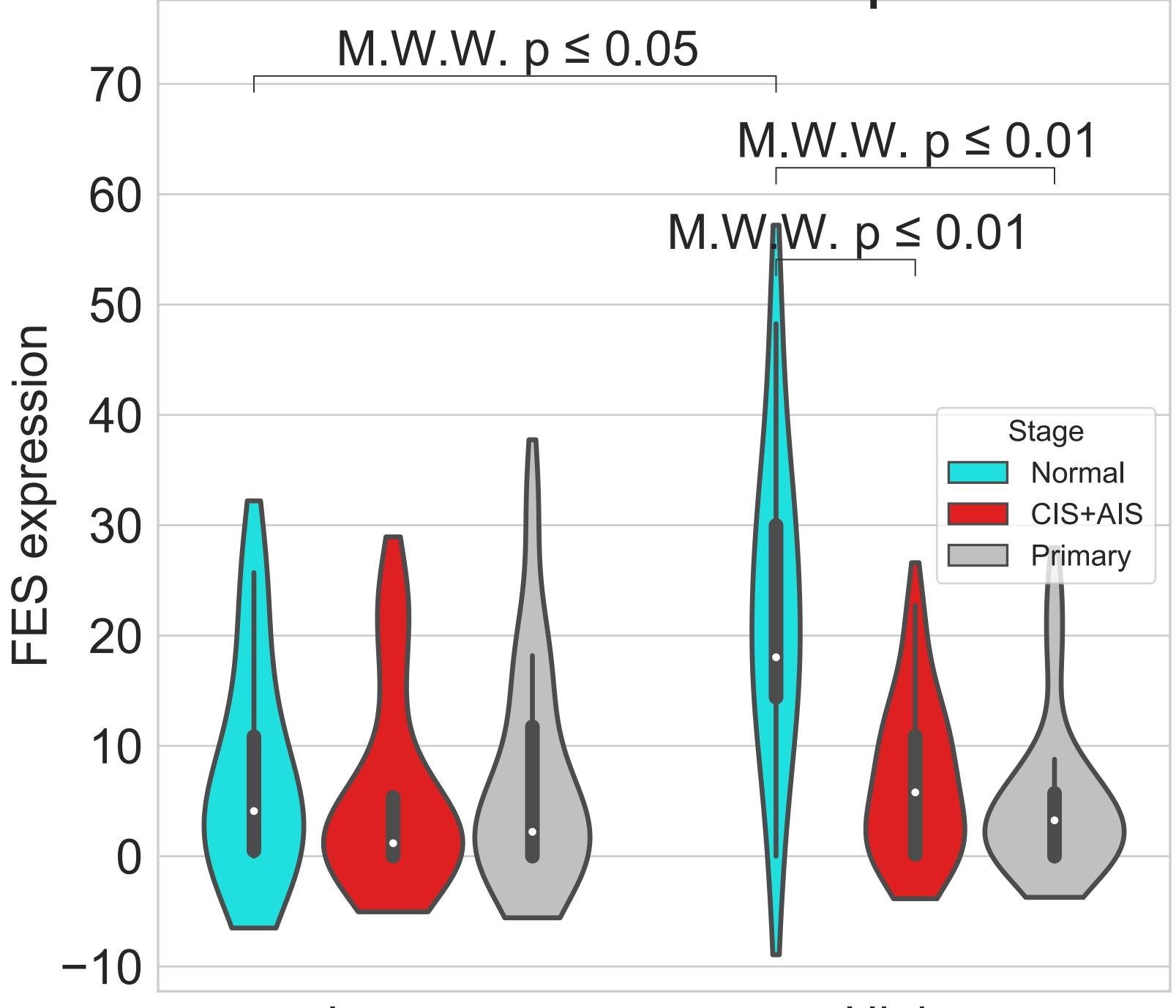
FES: Kruskal-Wallis p=0.034



Lower Higher Mutation Shared Count per TMB