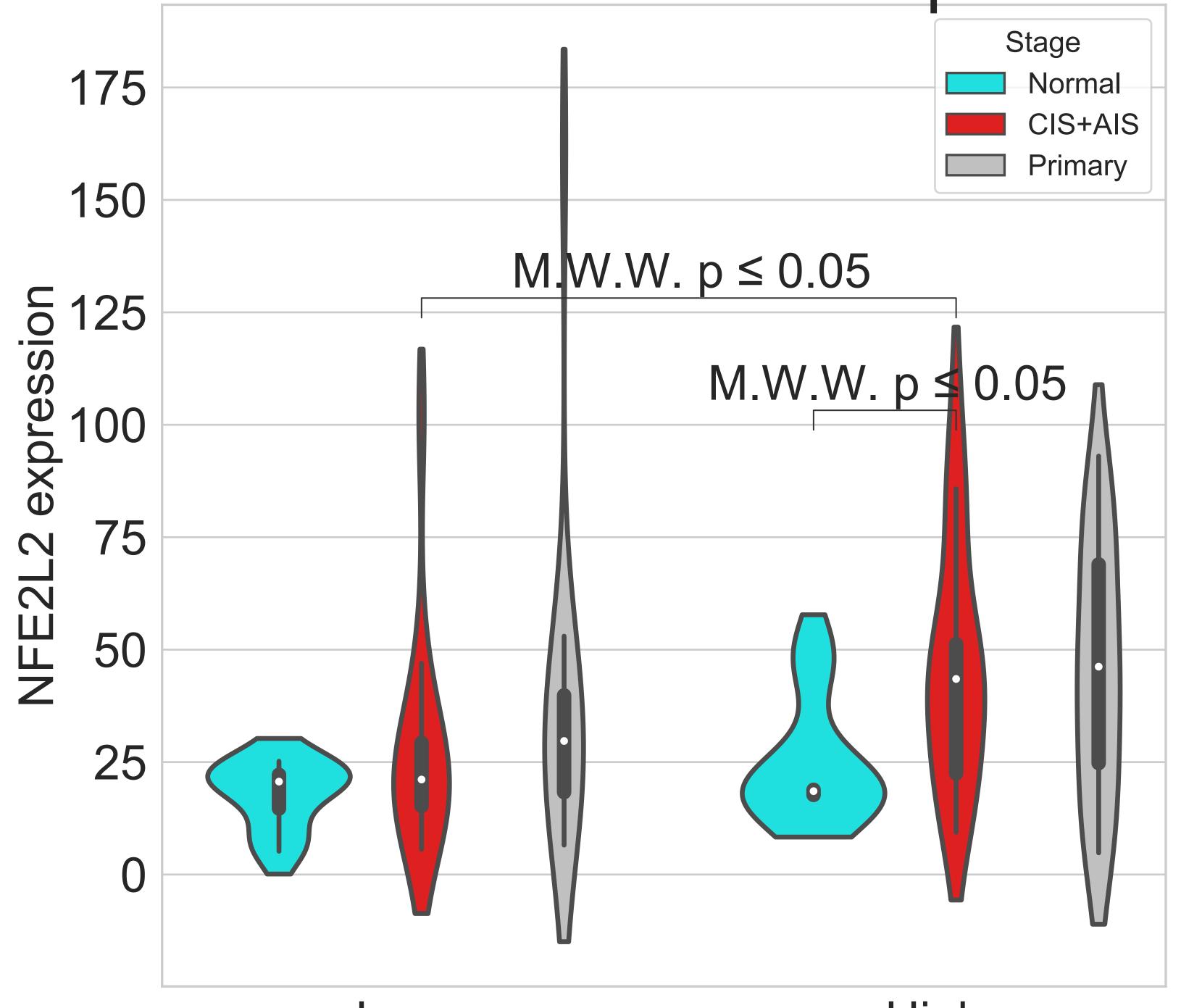
NFE2L2: Kruskal-Wallis p=0.035



Lower Higher Mutation Shared Count per TMB