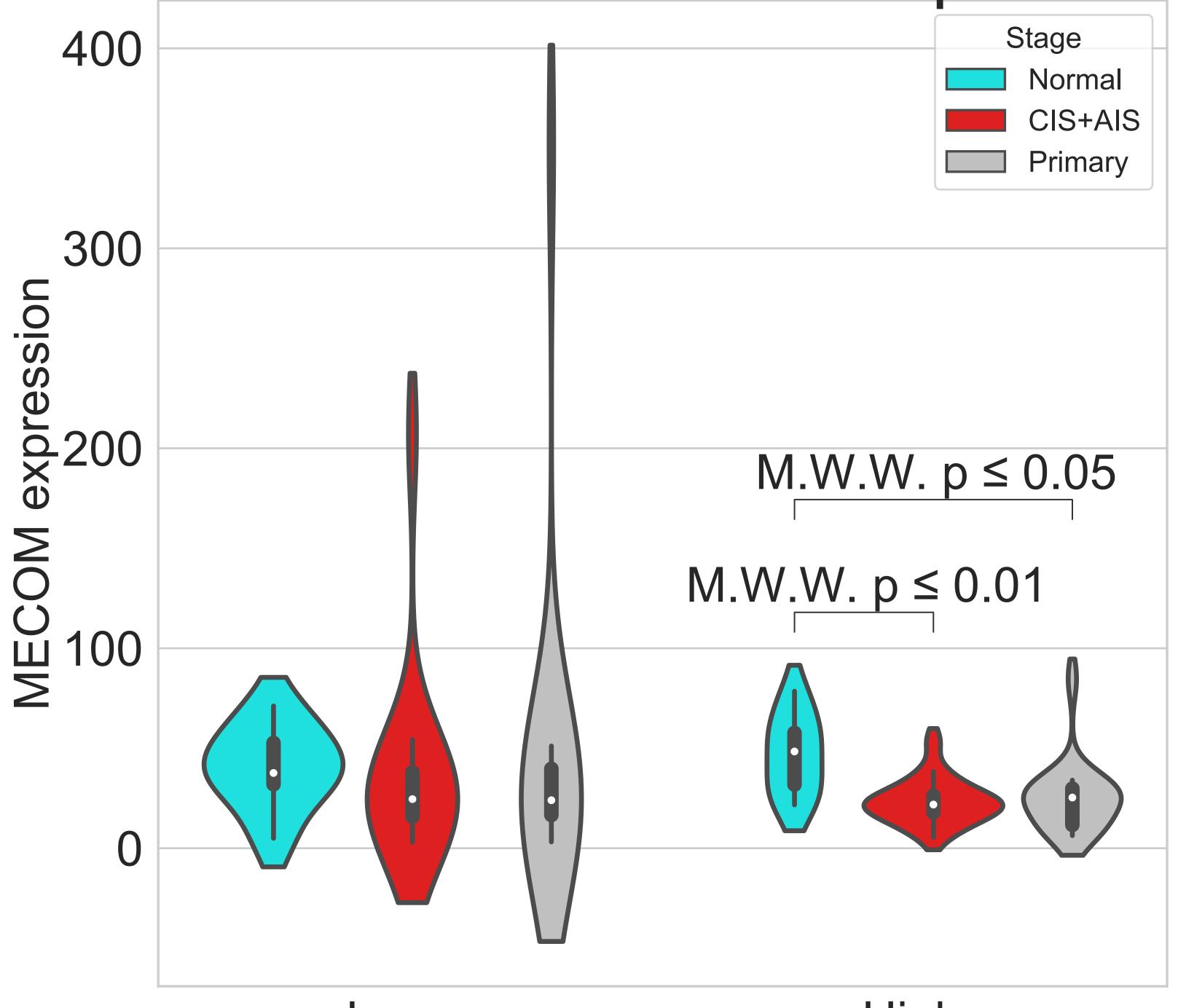
MECOM: Kruskal-Wallis p=0.038



Lower Higher Mutation Shared Count per TMB