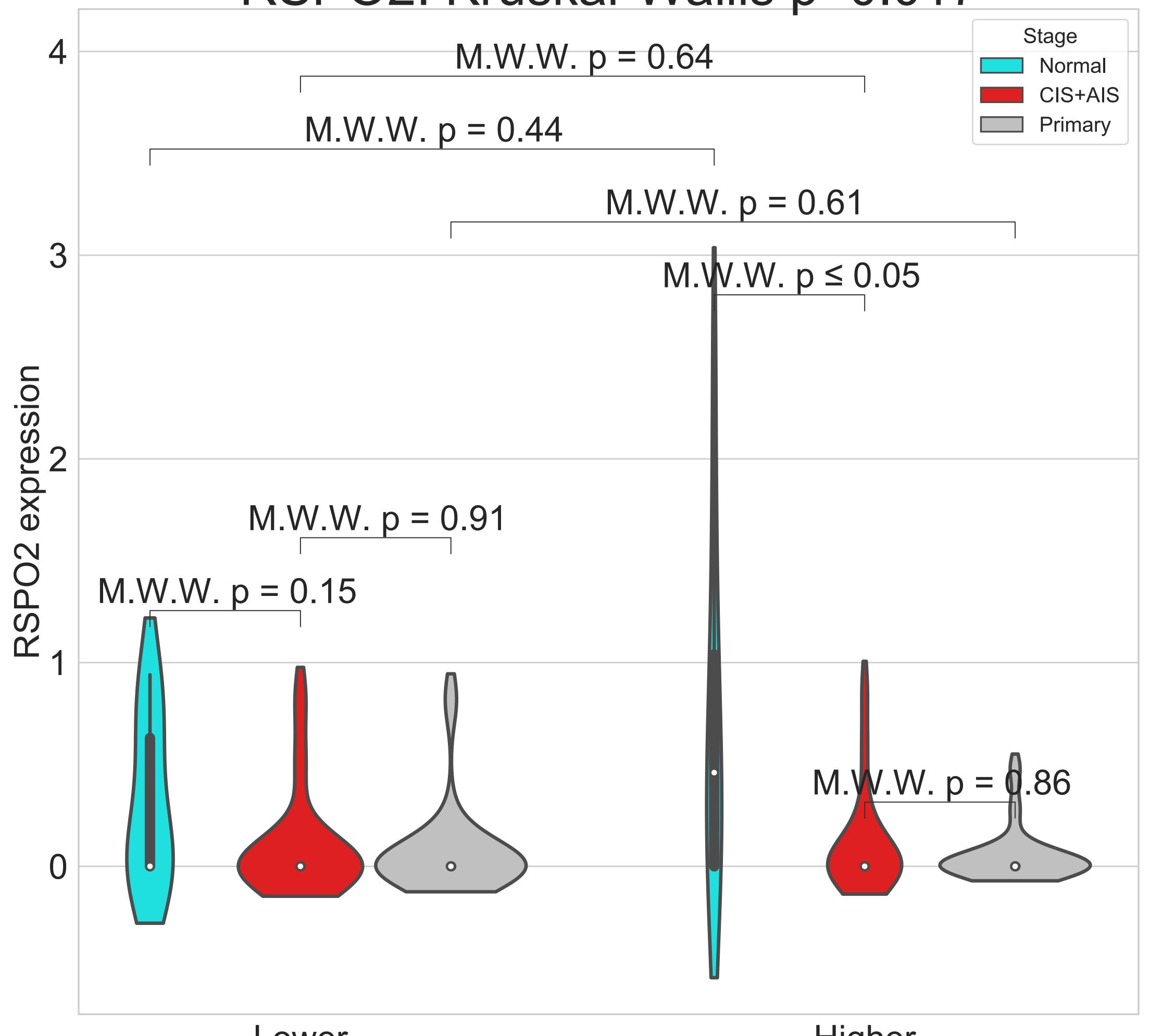
RSP02: Kruskal-Wallis p=0.017



Lower Higher Mutation Shared Count per TMB