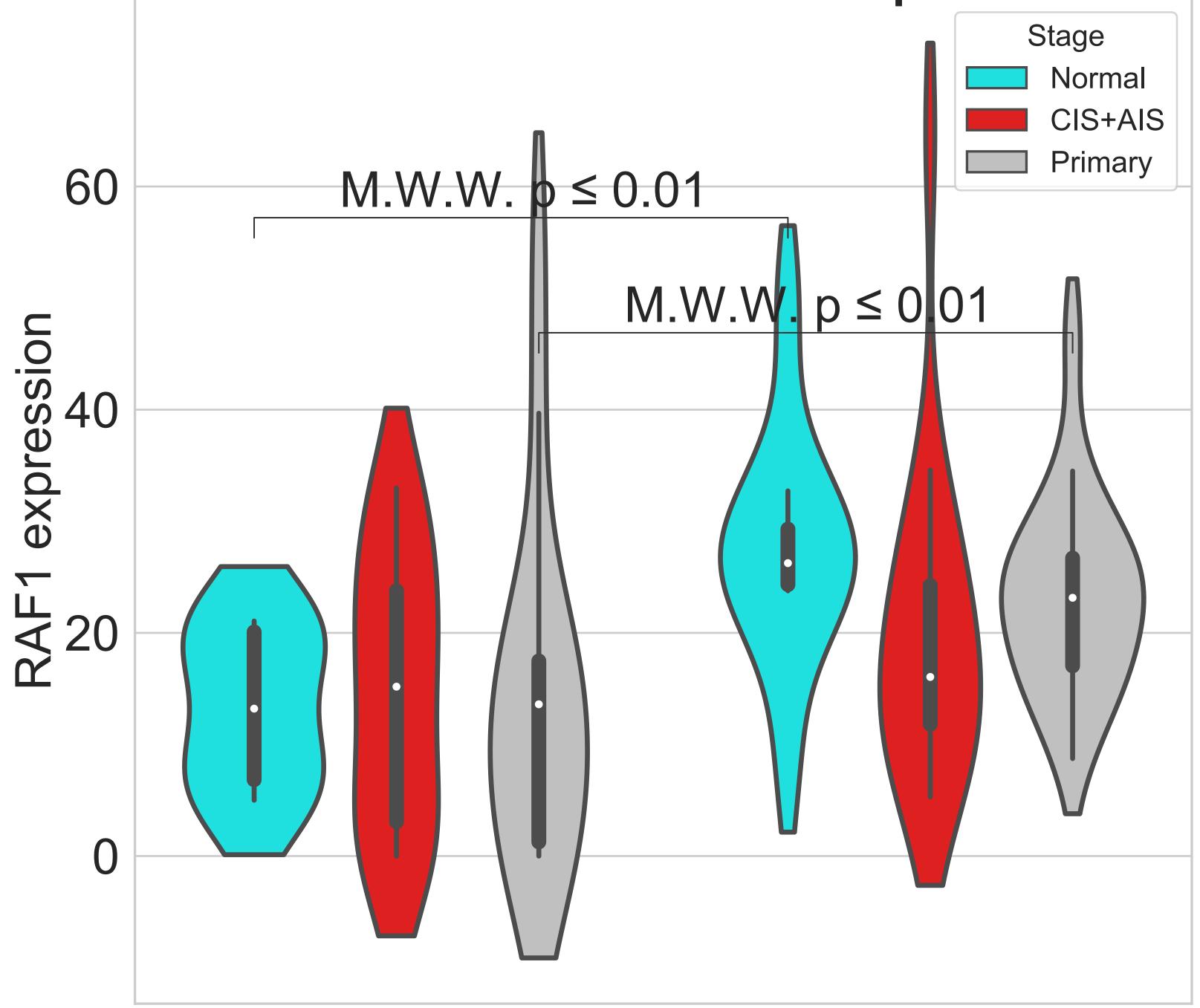
RAF1: Kruskal-Wallis p=0.007

Stage
Normal



Lower Higher Mutation Shared Count per TMB