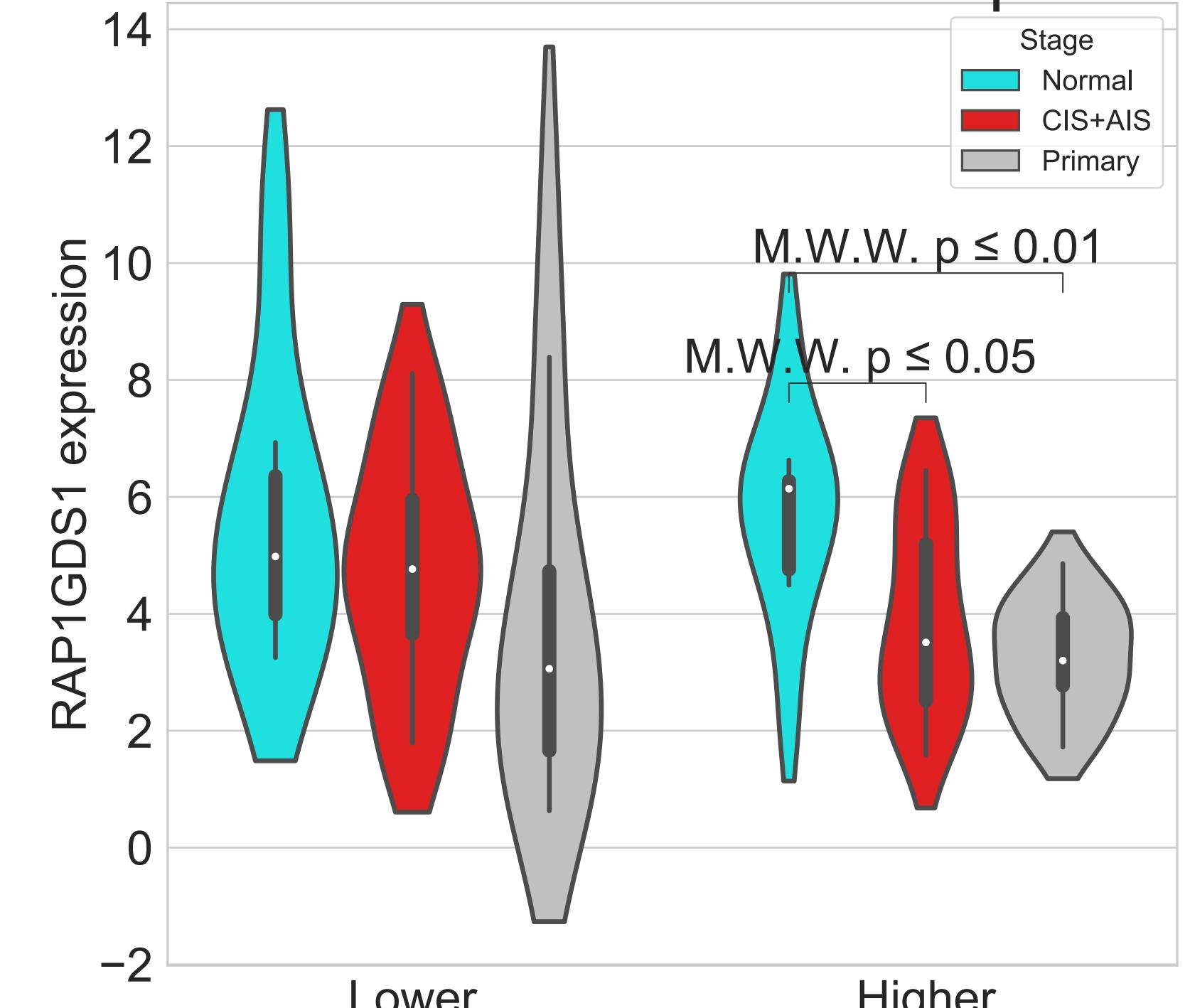
RAP1GDS1: Kruskal-Wallis p=0.010



Lower Higher Mutation Shared Count per TMB