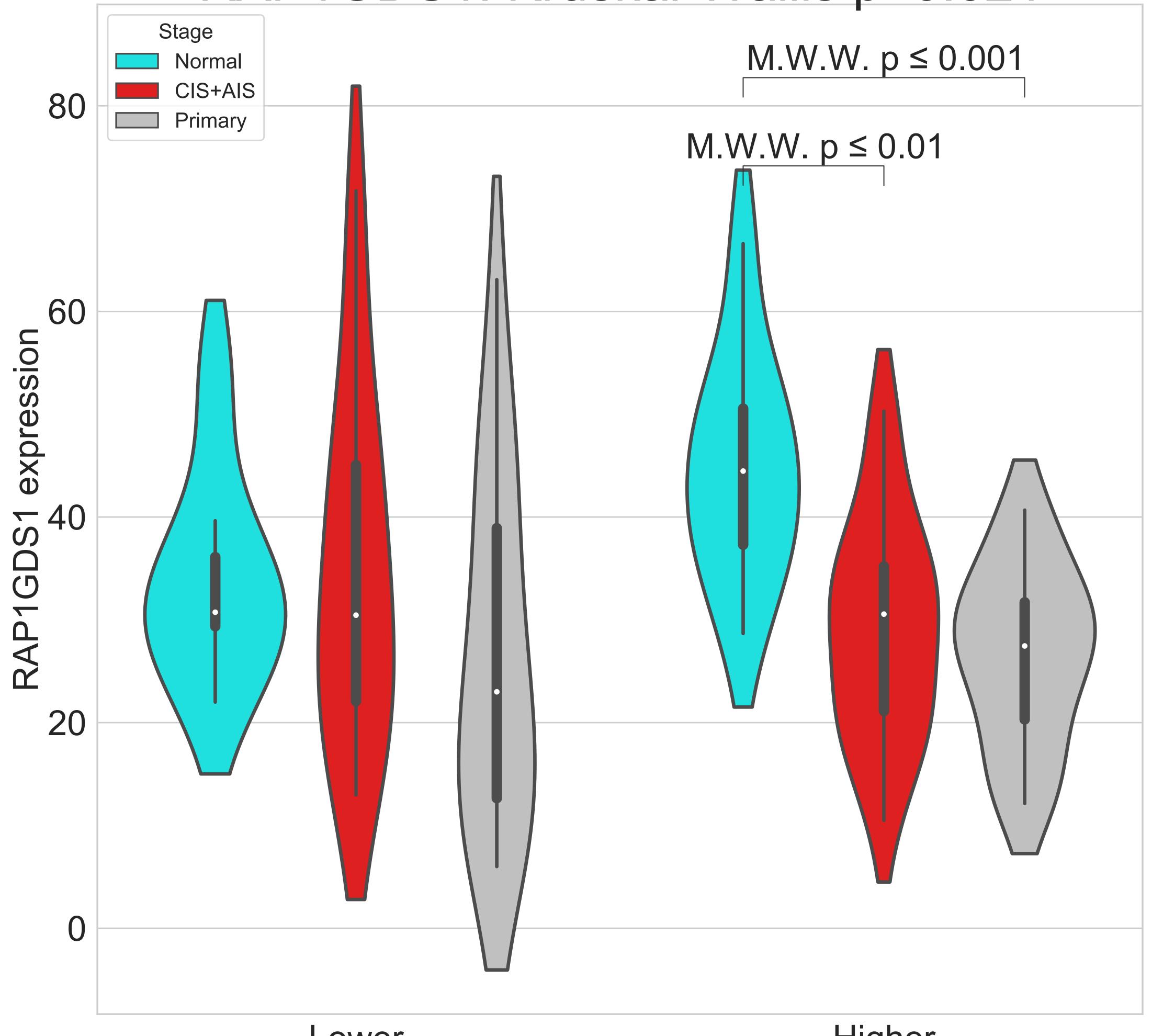
RAP1GDS1: Kruskal-Wallis p=0.021



Lower
Mutation Shared Count per TMB