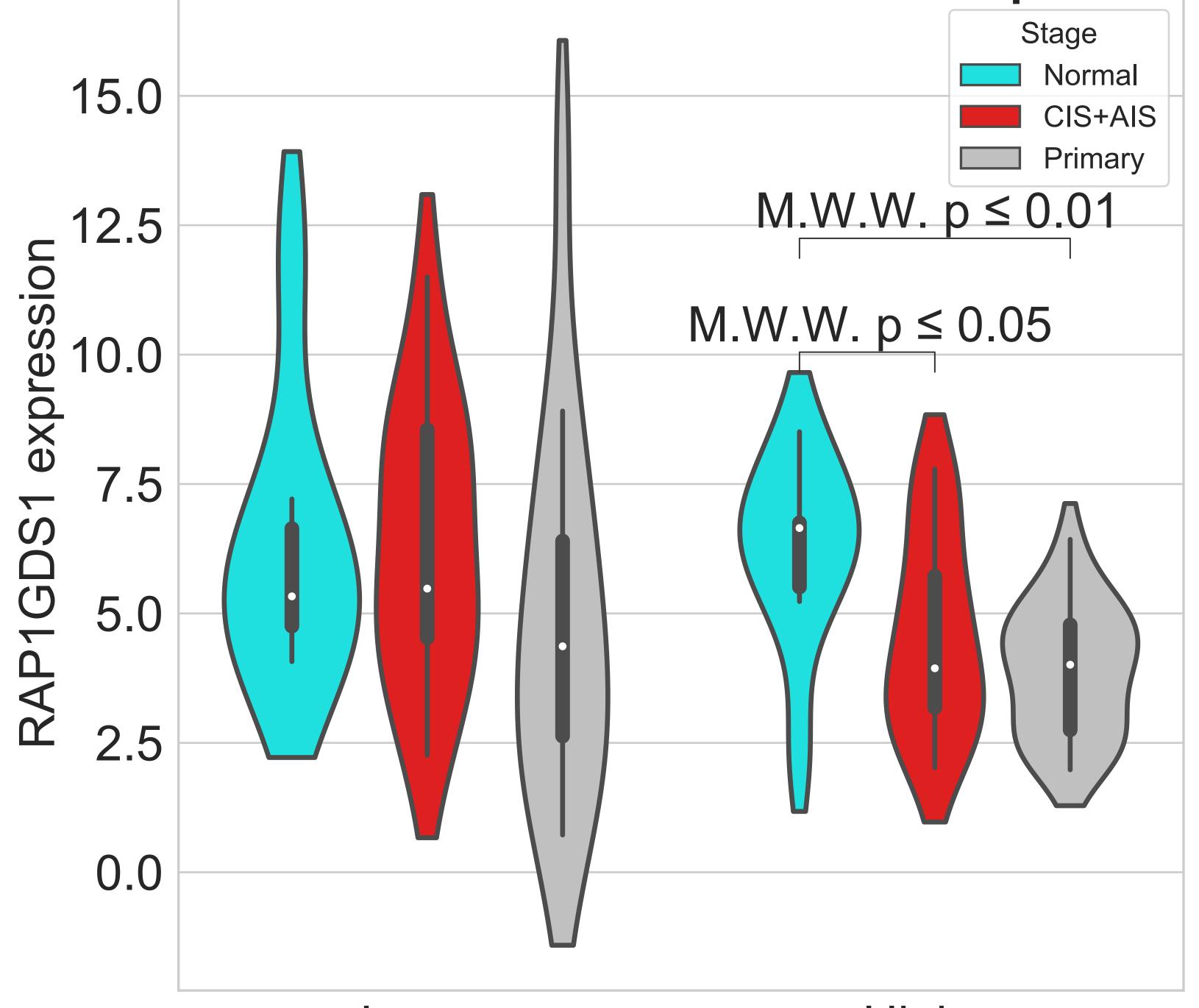
RAP1GDS1: Kruskal-Wallis p=0.01



Lower Higher Mutation Shared Proportion (SYN)