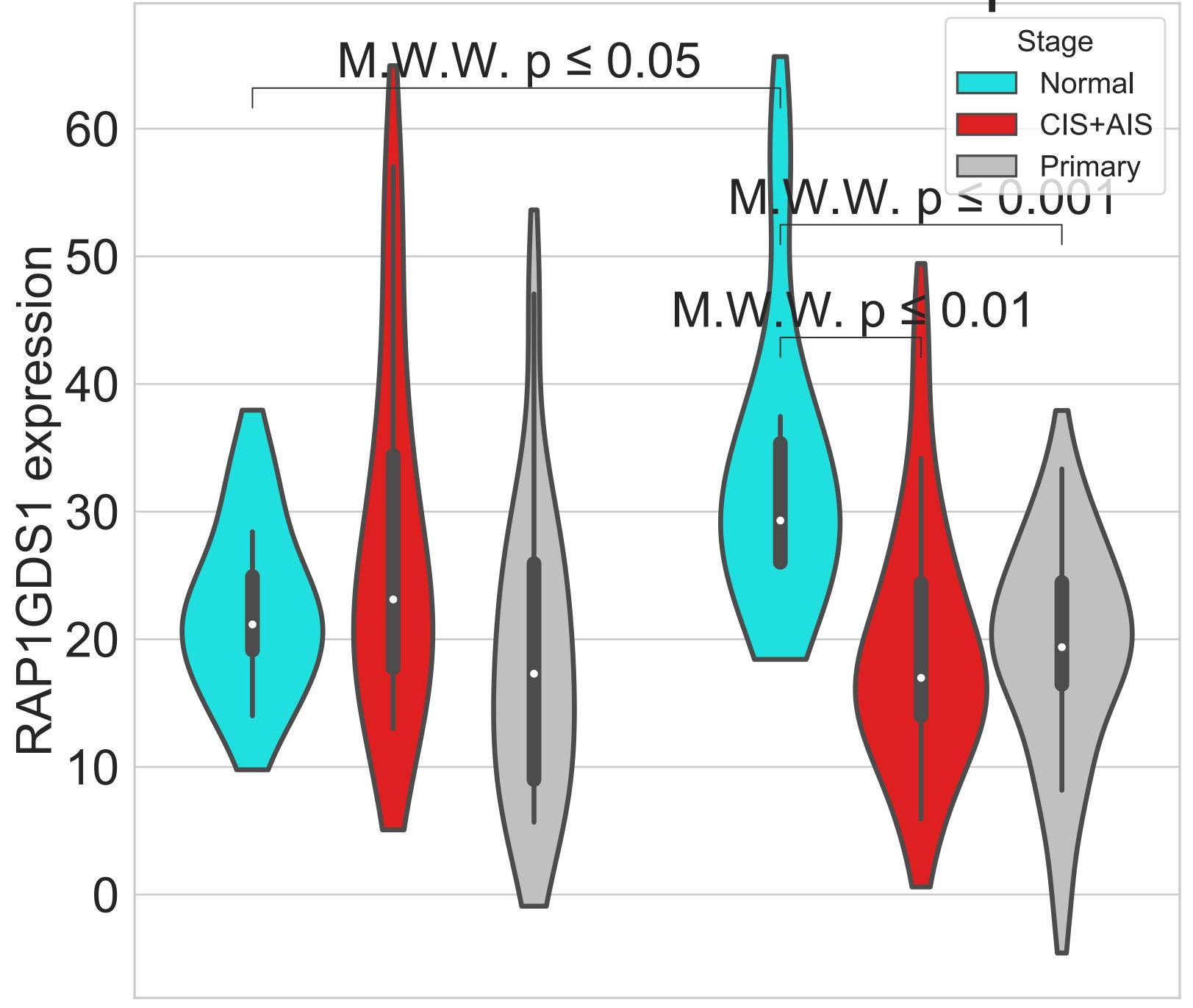
RAP1GDS1: Kruskal-Wallis p=0.007



Lower Higher Mutation Shared Proportion (Union & SYN)