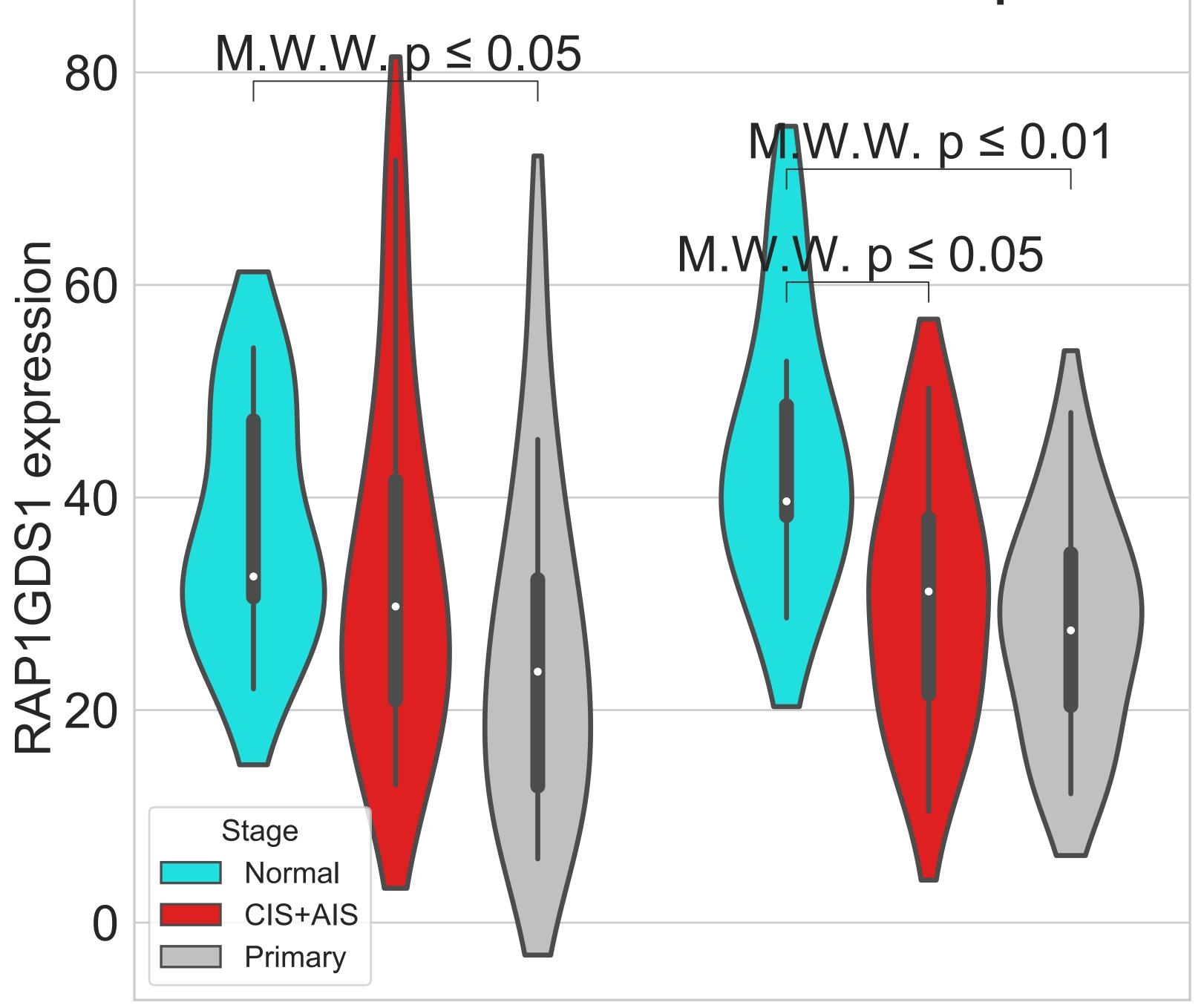
RAP1GDS1: Kruskal-Wallis p=0.035



Lower
Mutation Shared Count (SYN)