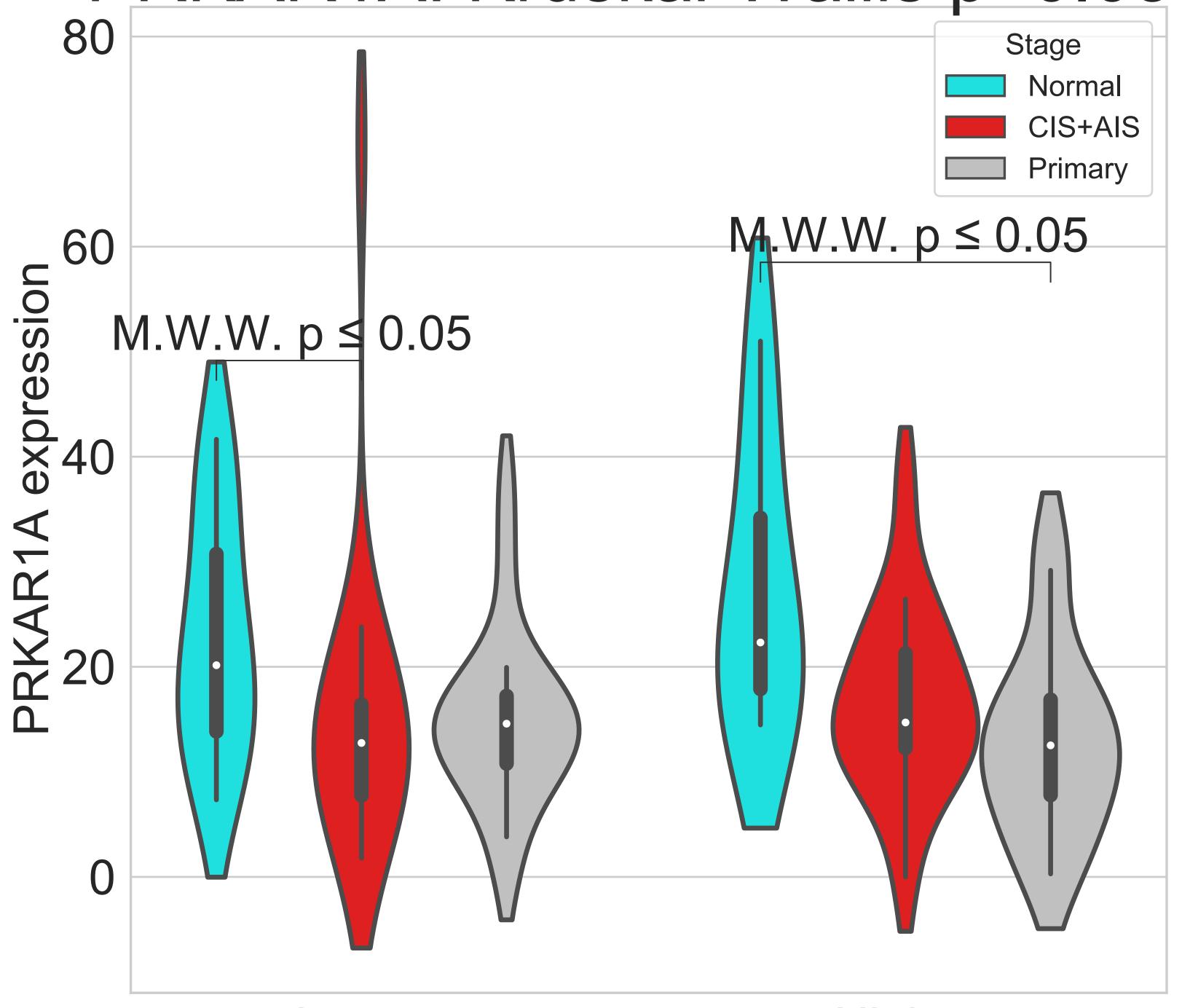
PRKAR1A: Kruskal-Wallis p=0.035



Lower
Mutation Shared Count (SYN) per TMB