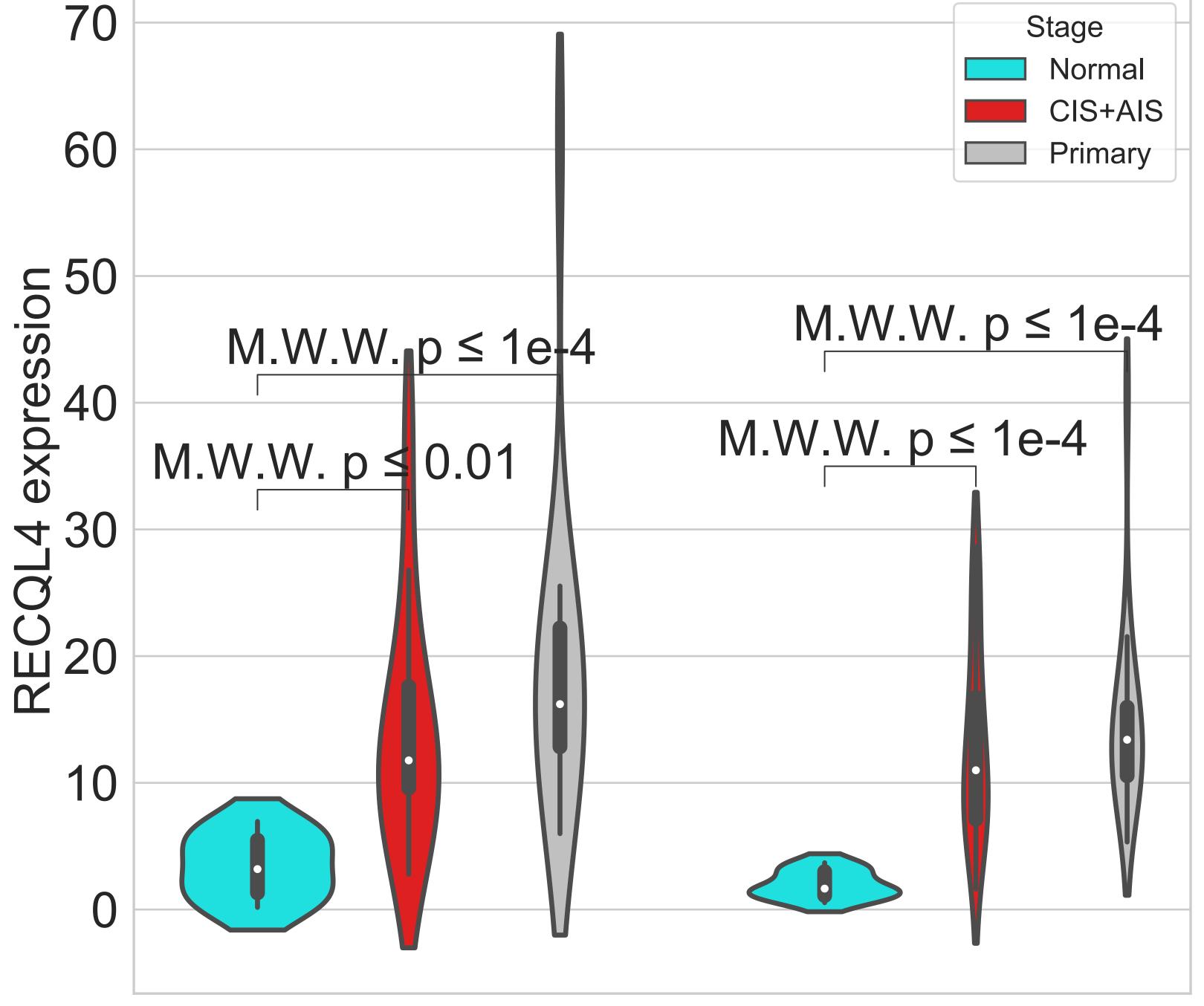
RECQL4: Kruskal-Wallis p=0.000 Stage Normal CIS+AIS Primary



Higher Lower Mutation Shared Count (SYN) per TMB