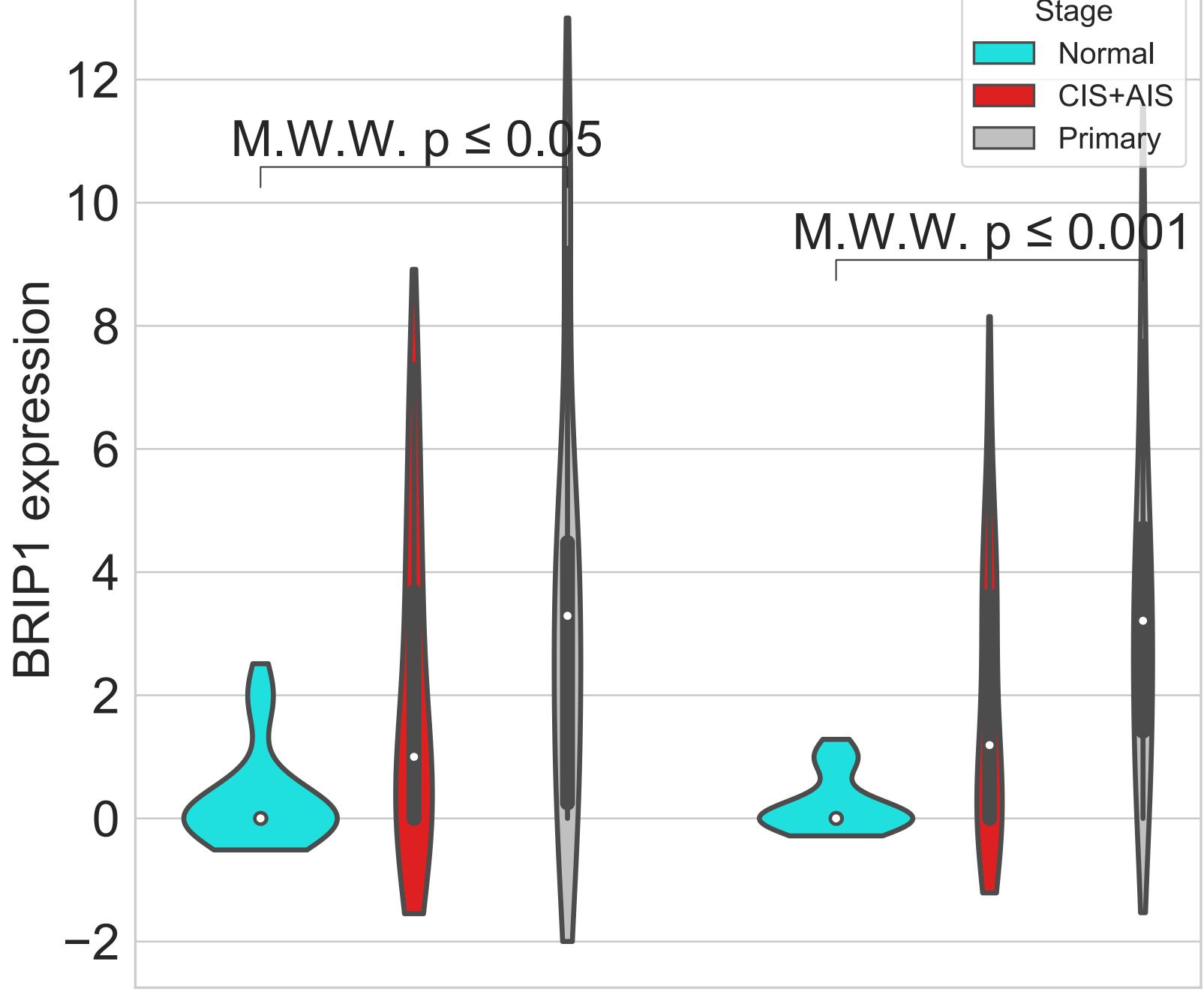
BRIP1: Kruskal-Wallis p=0.002 Stage Normal CIS+AIS



Higher Lower Mutation Shared Count (SYN) per TMB