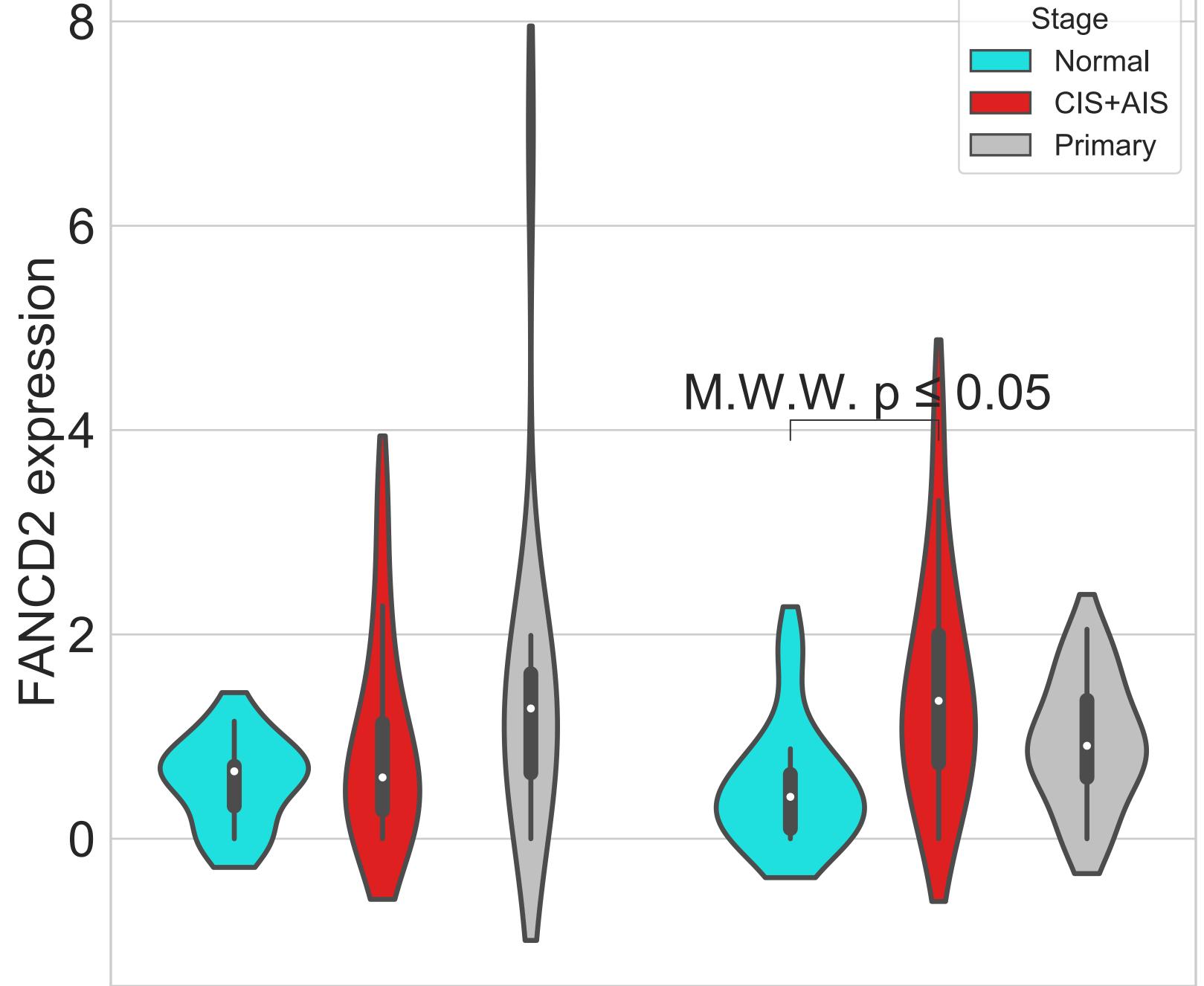
FANCD2: Kruskal-Wallis p=0.039 Stage Normal CIS+AIS Primary



Higher Lower Mutation Shared Count (SYN) per TMB