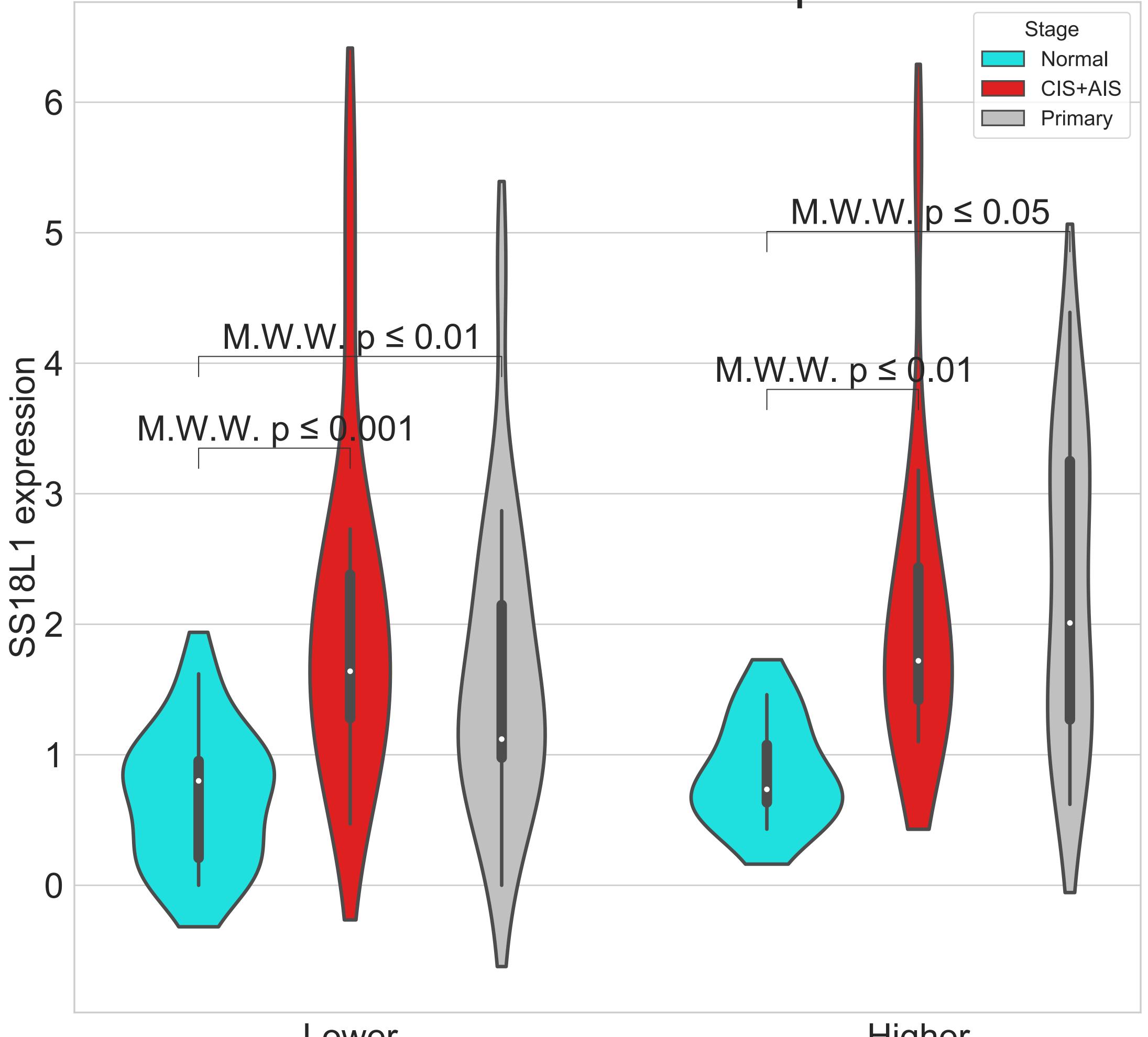
SS18L1: Kruskal-Wallis p=0.000



Higher Lower Mutation Shared Count (SYN) per TMB