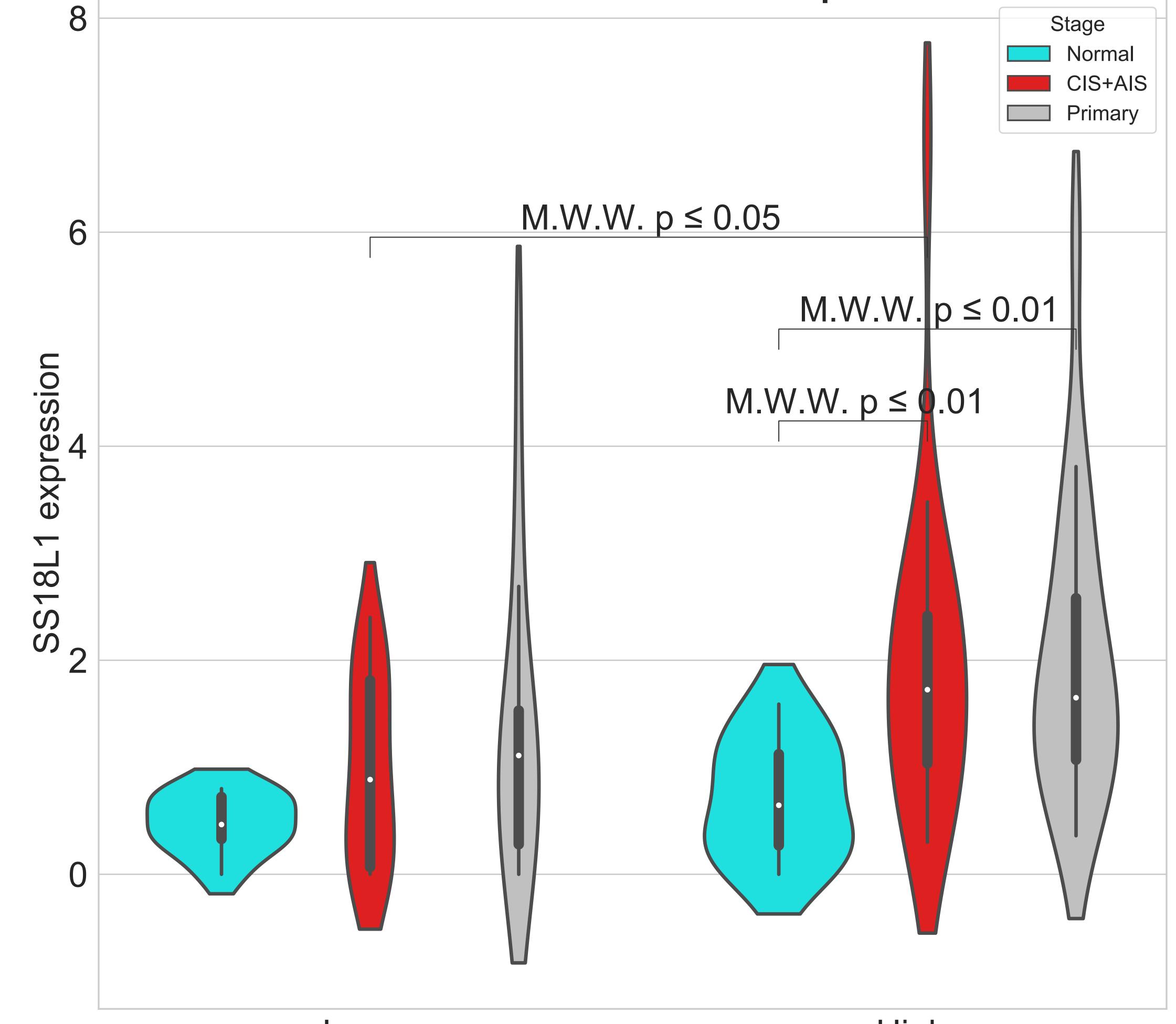
SS18L1: Kruskal-Wallis p=0.001



Higher Lower Mutation Shared Proportion (Union & SYN)