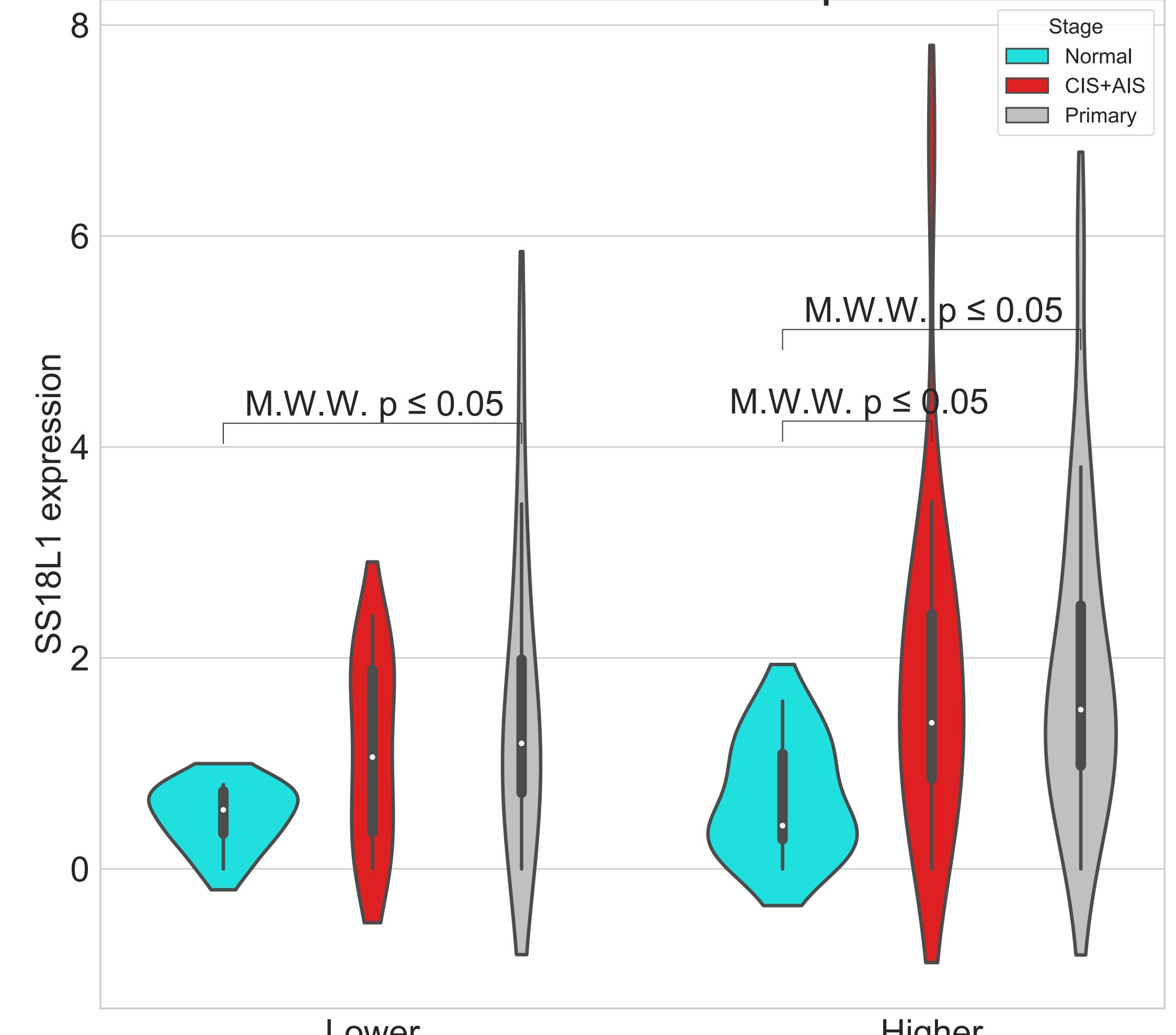
SS18L1: Kruskal-Wallis p=0.011



Higher Lower Mutation Shared Count per TMB