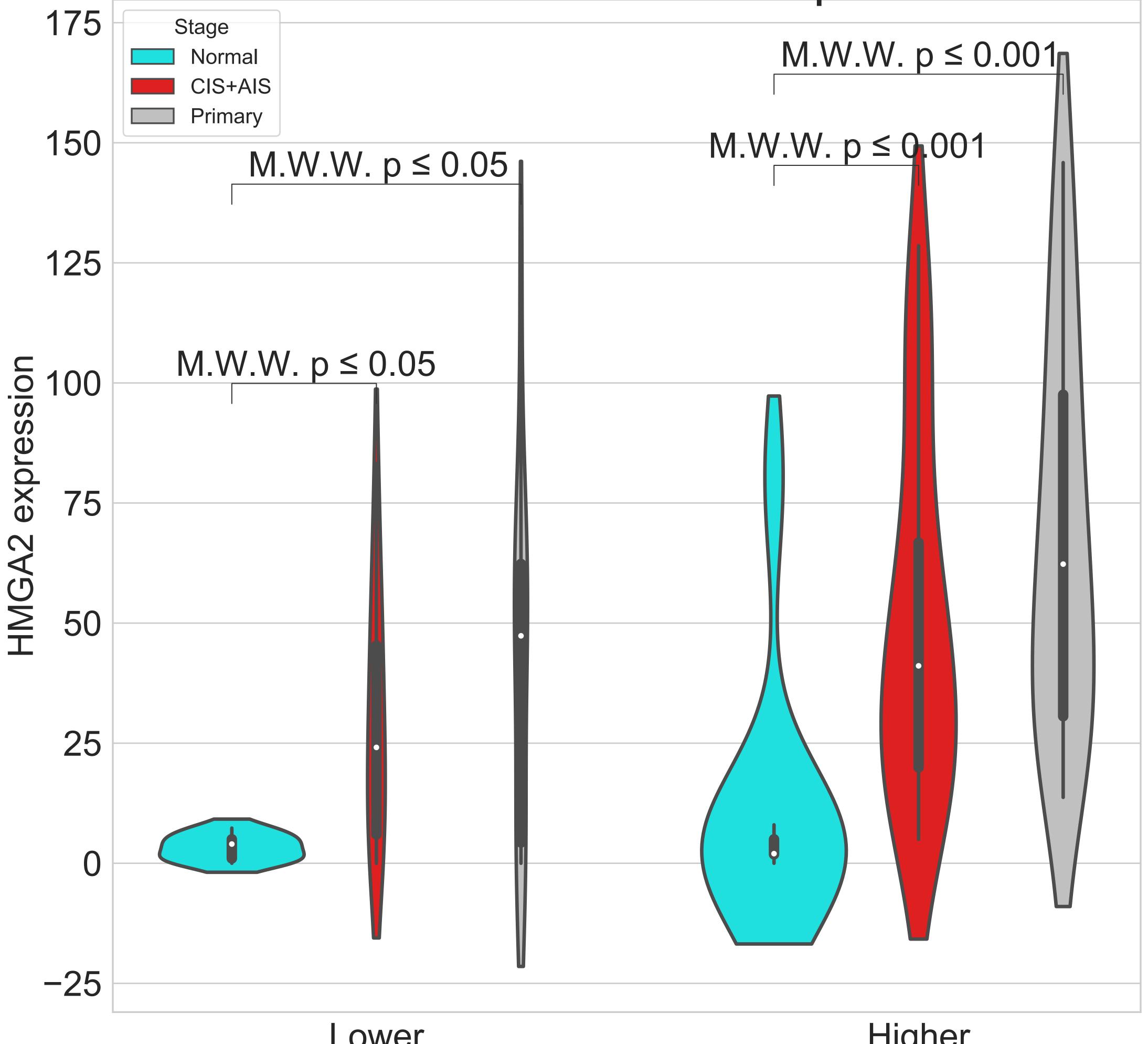
HMGA2: Kruskal-Wallis p=0.000



Higher Lower Mutation Shared Count (SYN) per TMB