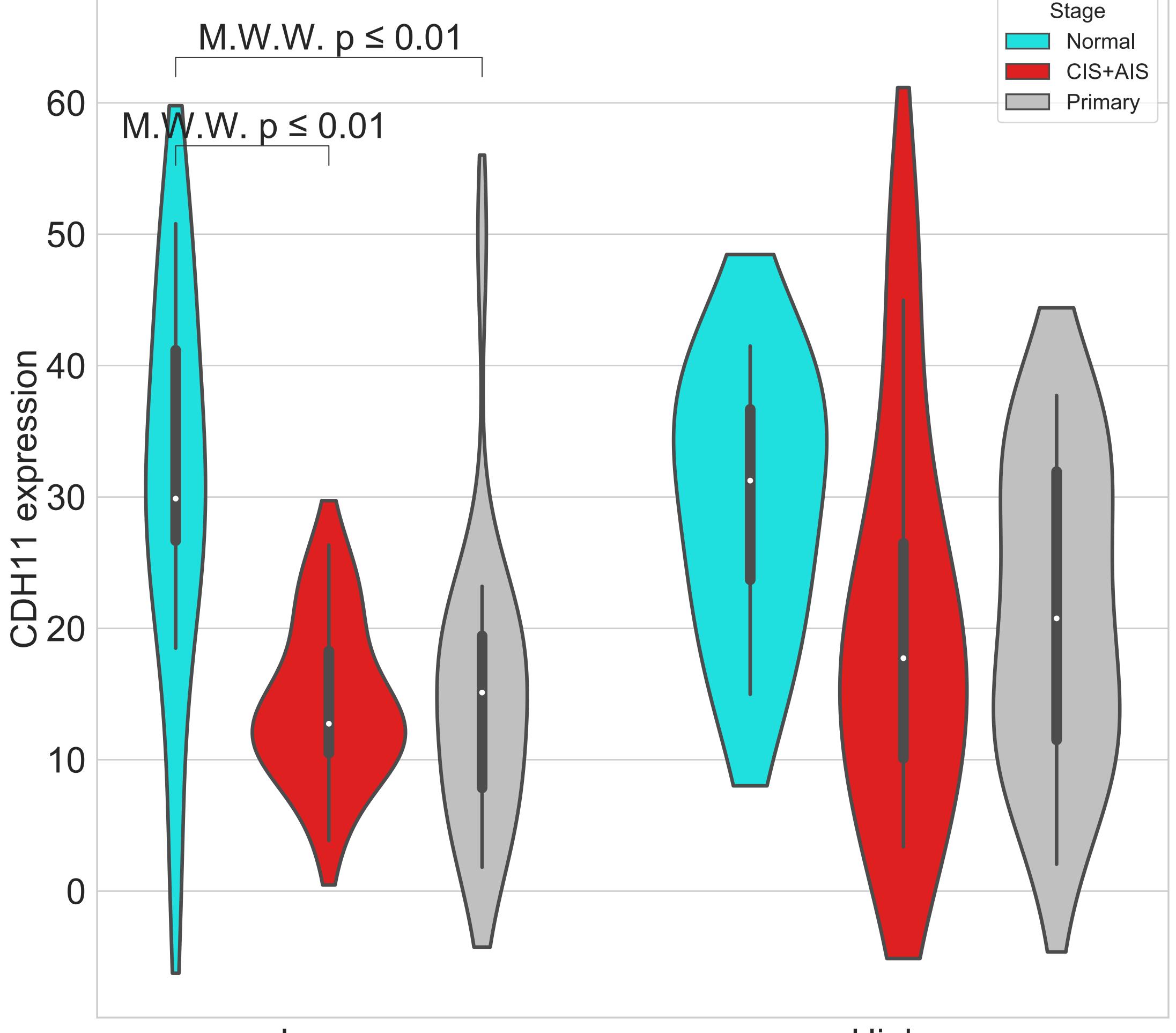
CDH11: Kruskal-Wallis p=0.004



Higher Lower Mutation Shared Count (SYN) per TMB