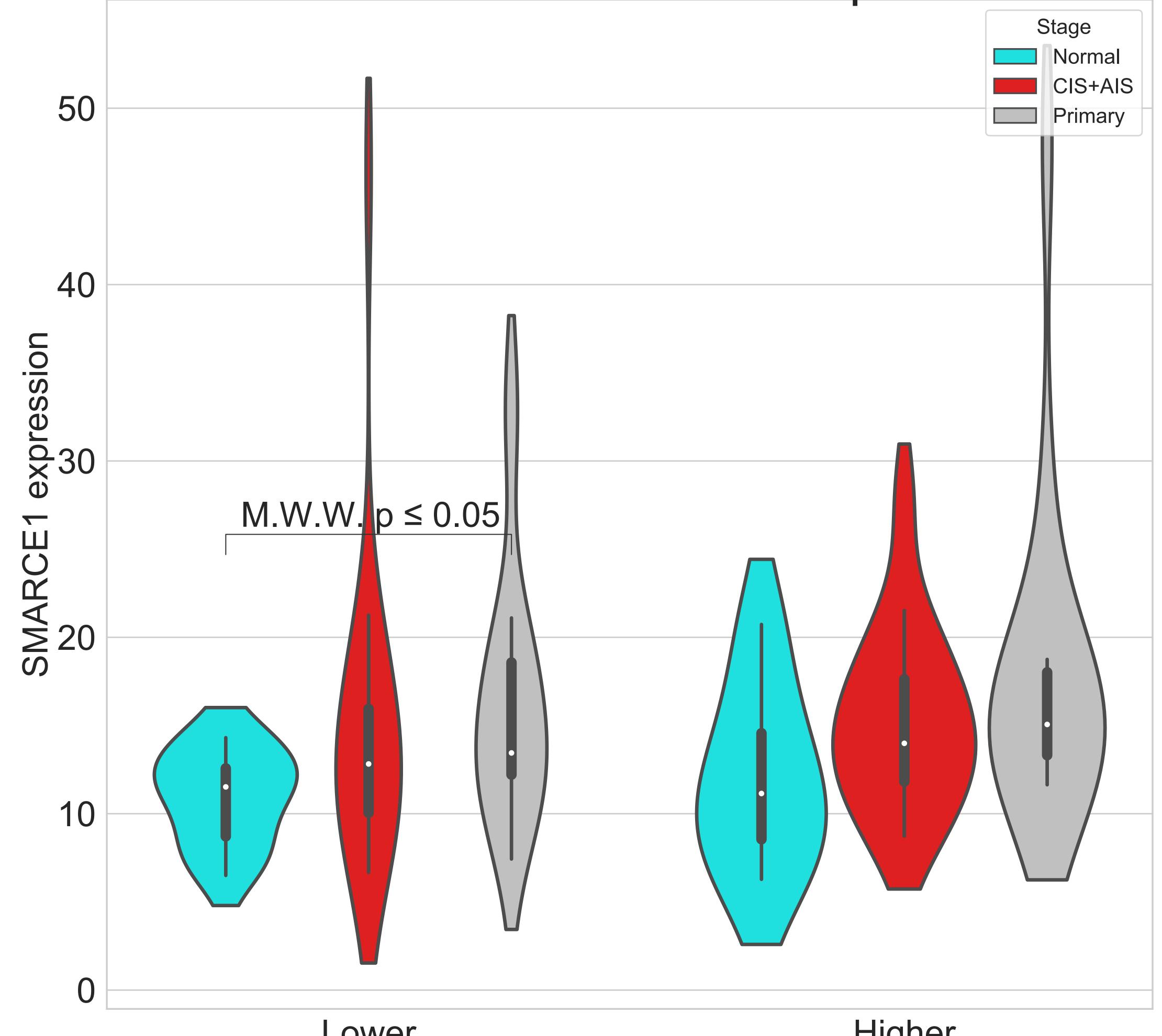
SMARCE1: Kruskal-Wallis p=0.041



Higher Lower Mutation Shared Count (SYN) per TMB