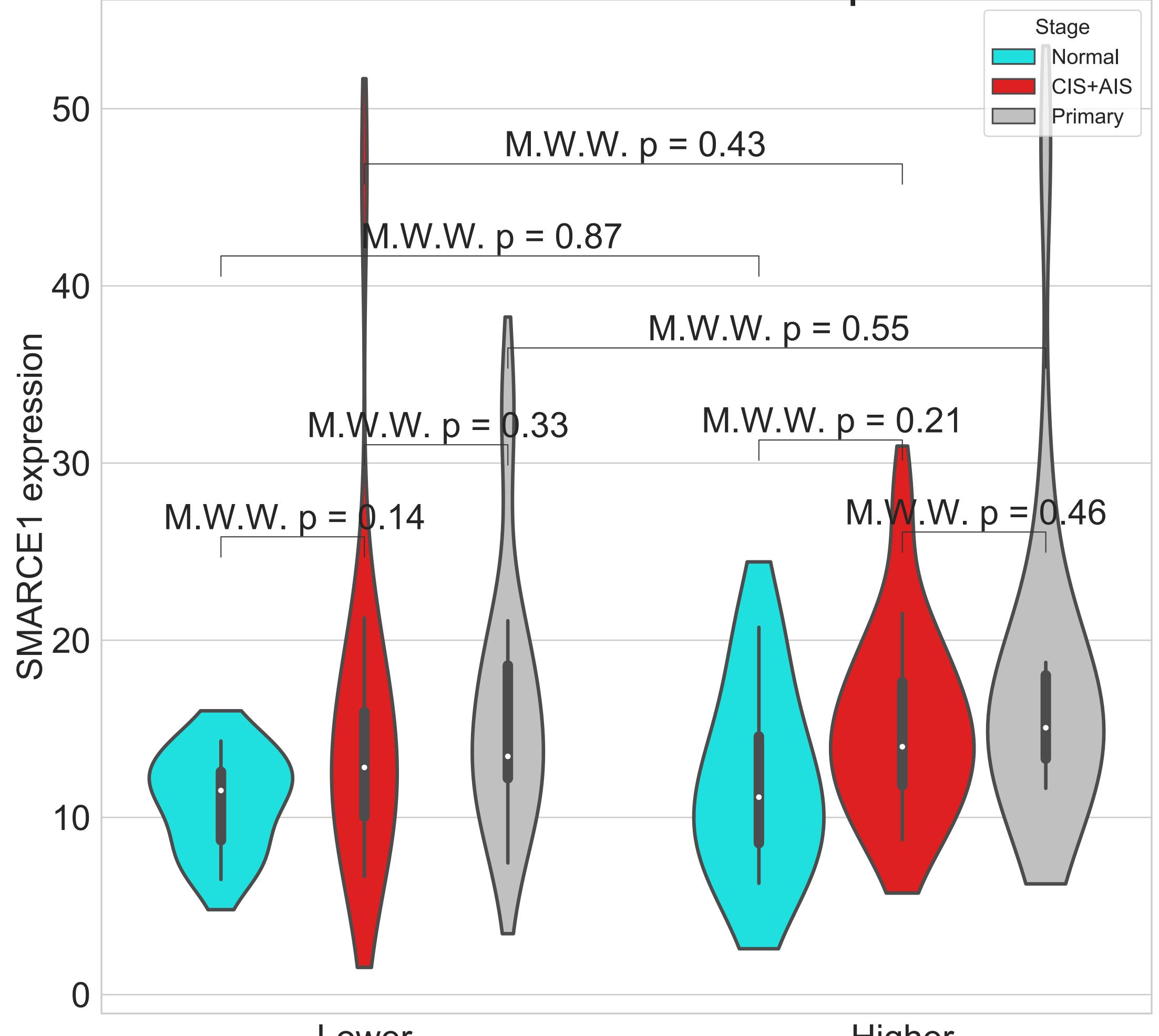
SMARCE1: Kruskal-Wallis p=0.041



Higher Lower Mutation Shared Proportion (Union & SYN)