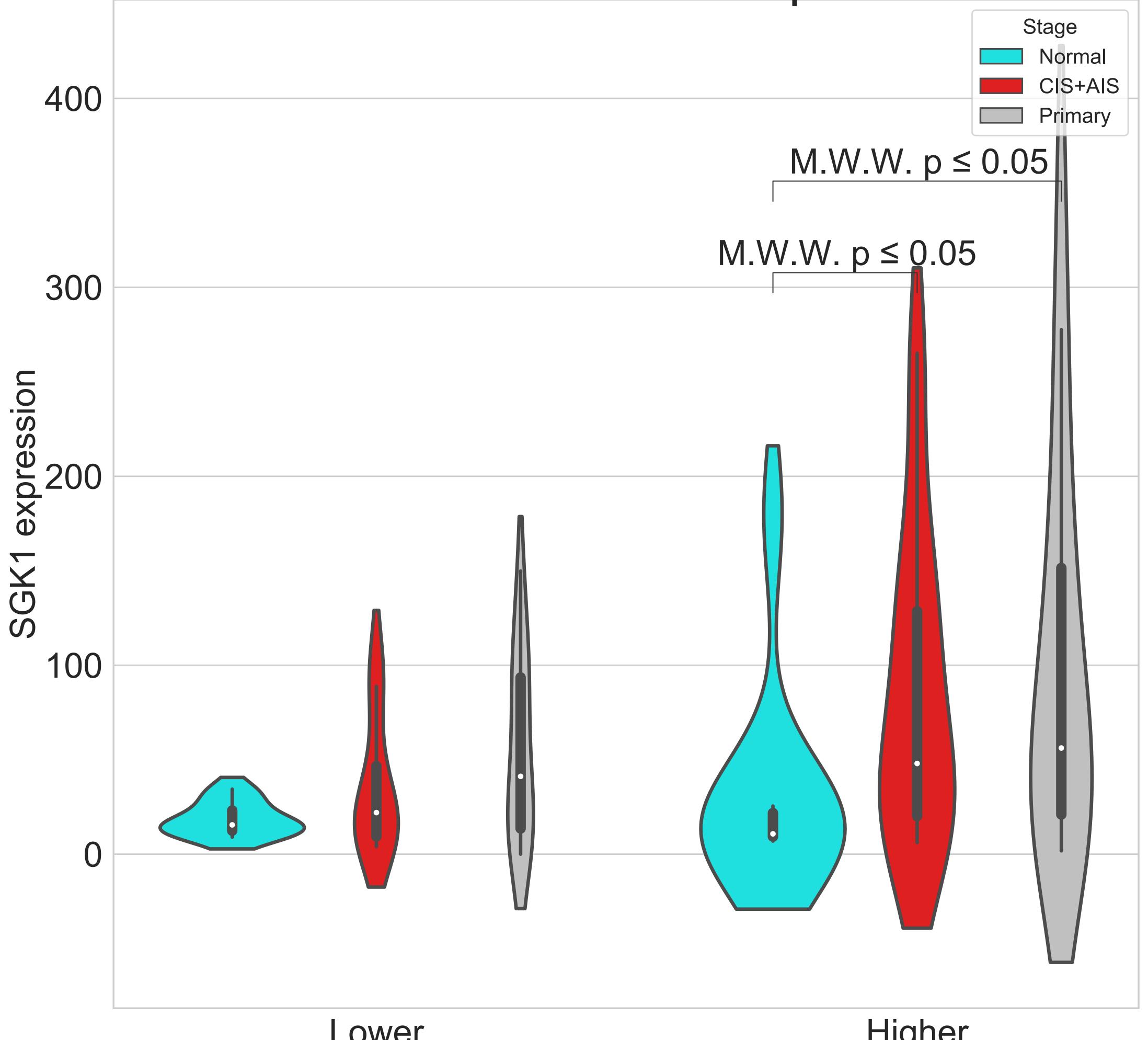
SGK1: Kruskal-Wallis p=0.048



Higher Lower Mutation Shared Proportion (SYN)