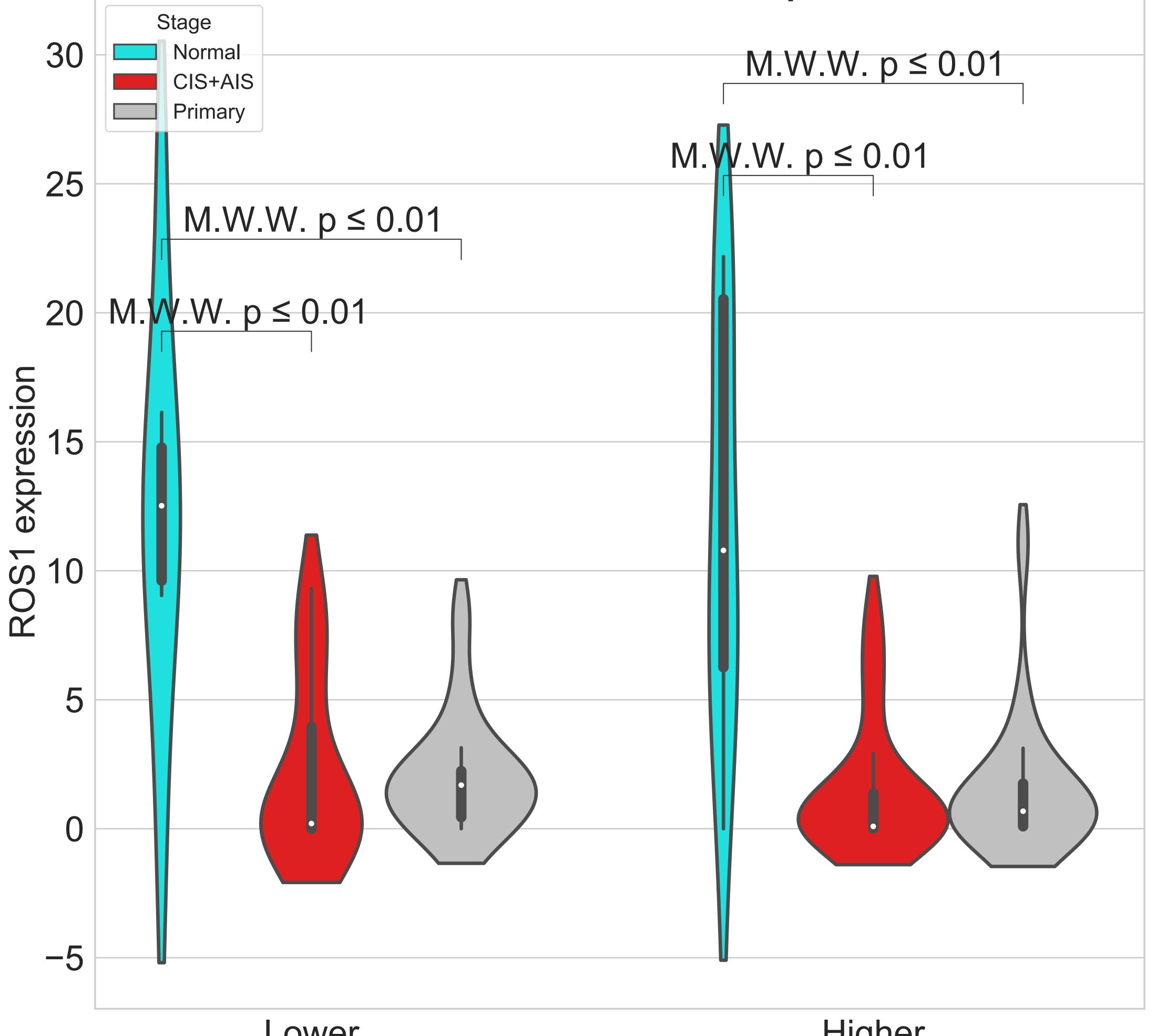
ROS1: Kruskal-Wallis p=0.000



Higher Lower Mutation Shared Proportion (SYN)