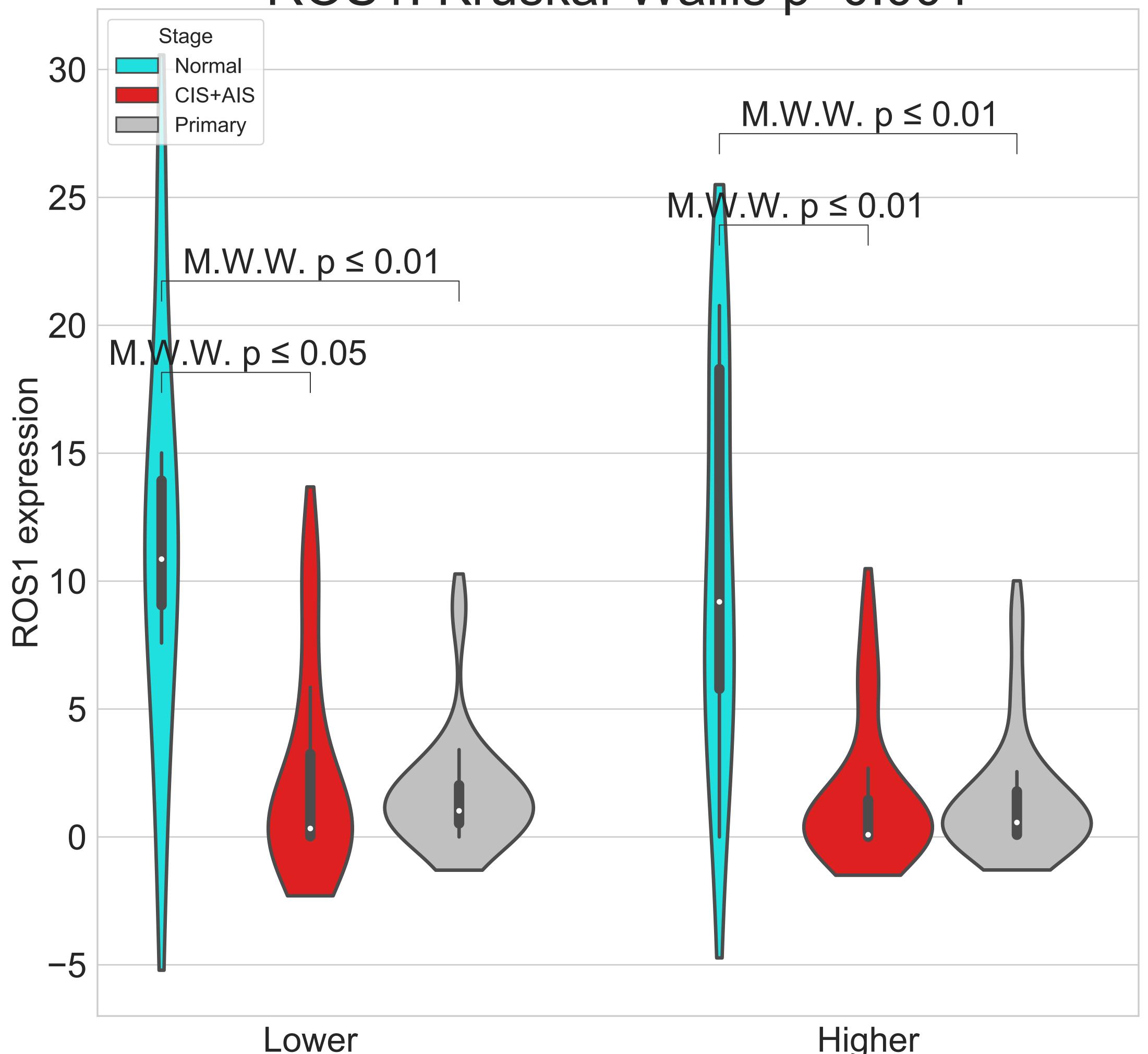
ROS1: Kruskal-Wallis p=0.001



r Mutation Shared Count per TMB