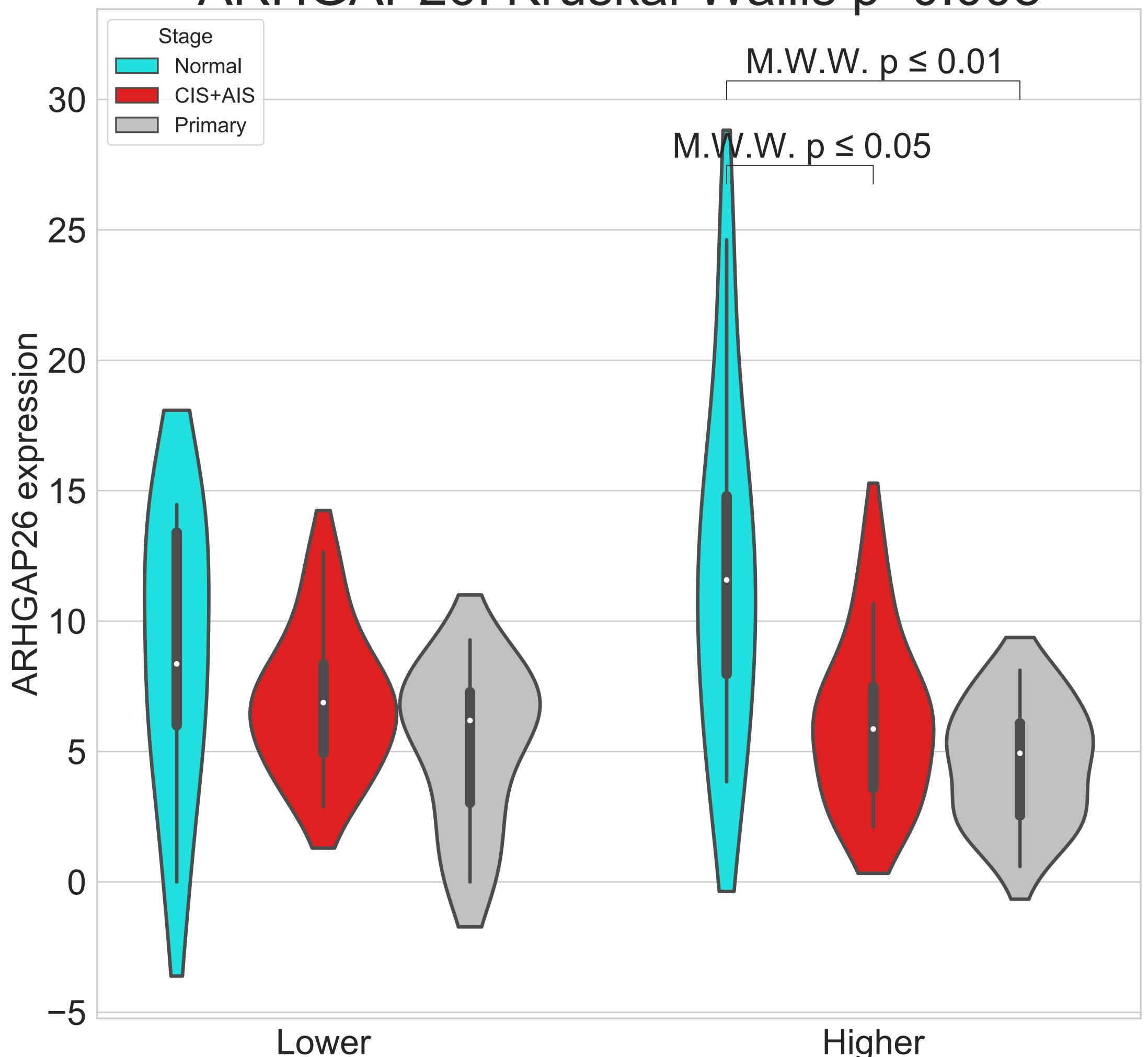
ARHGAP26: Kruskal-Wallis p=0.008



Higher Mutation Shared Count per TMB