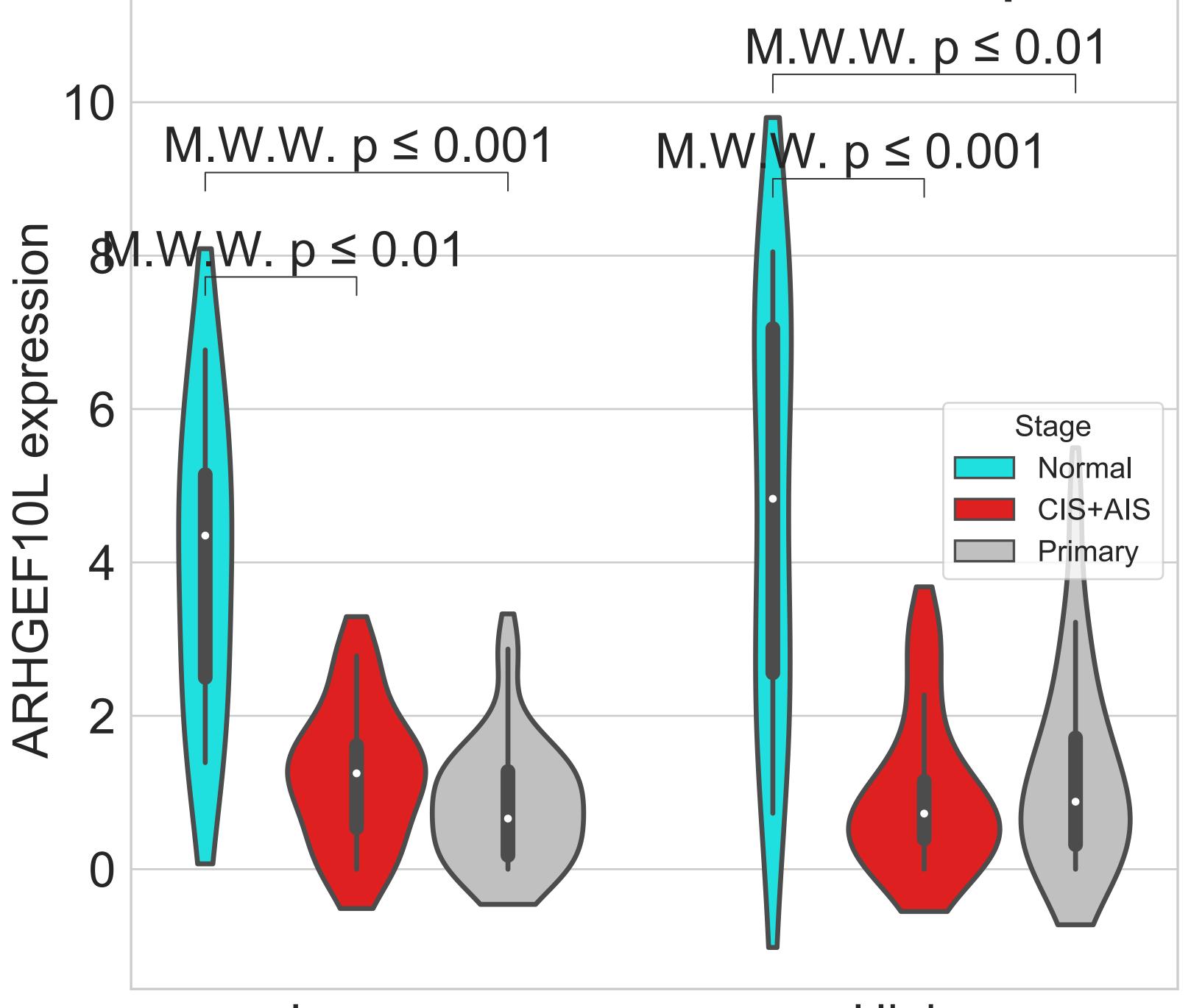
ARHGEF10L: Kruskal-Wallis p=0.00



Lower Higher Mutation Shared Count per TMB