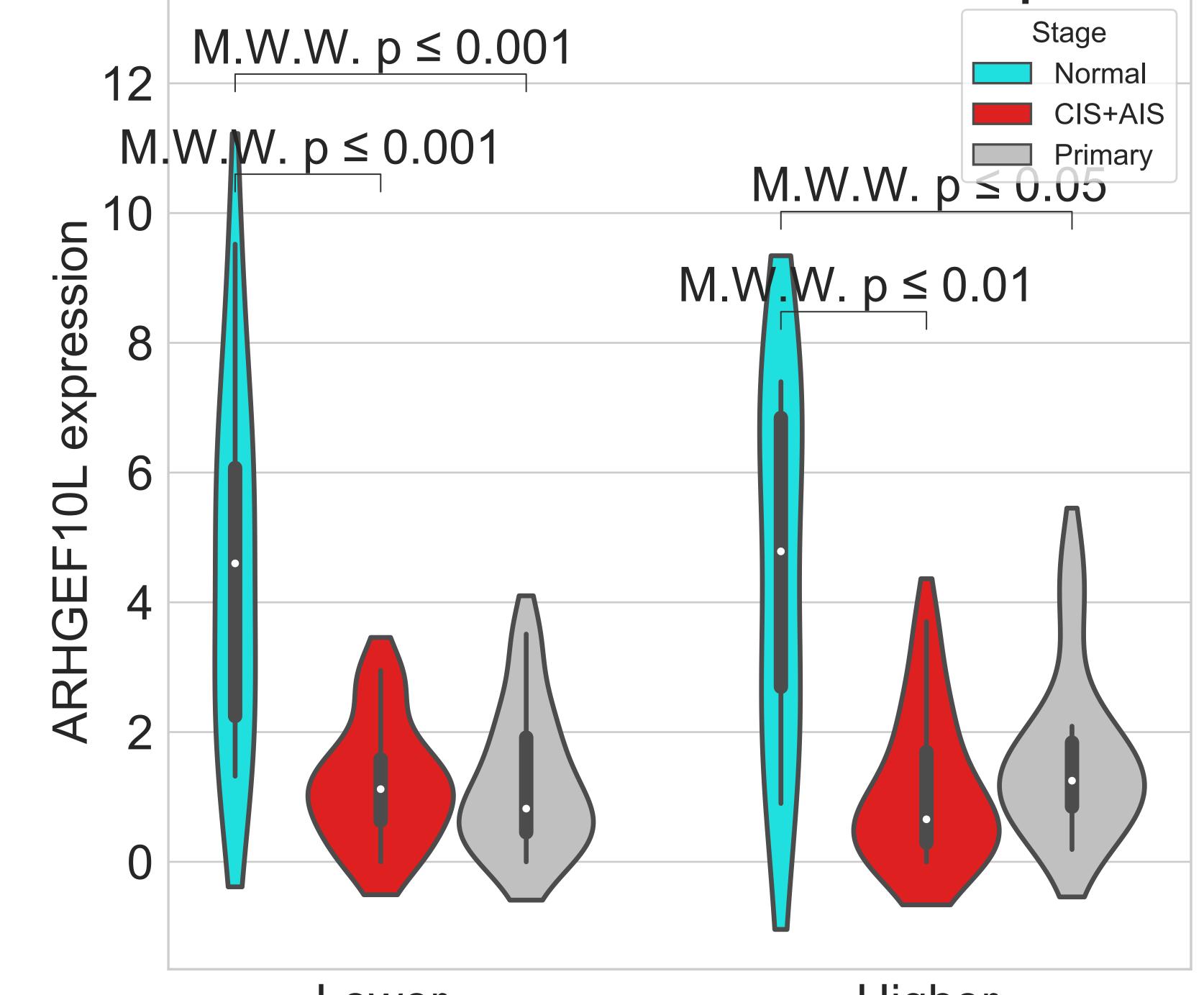
ARHGEF10L: Kruskal-Wallis p=0.00



Lower
Mutation Shared Count per TMB