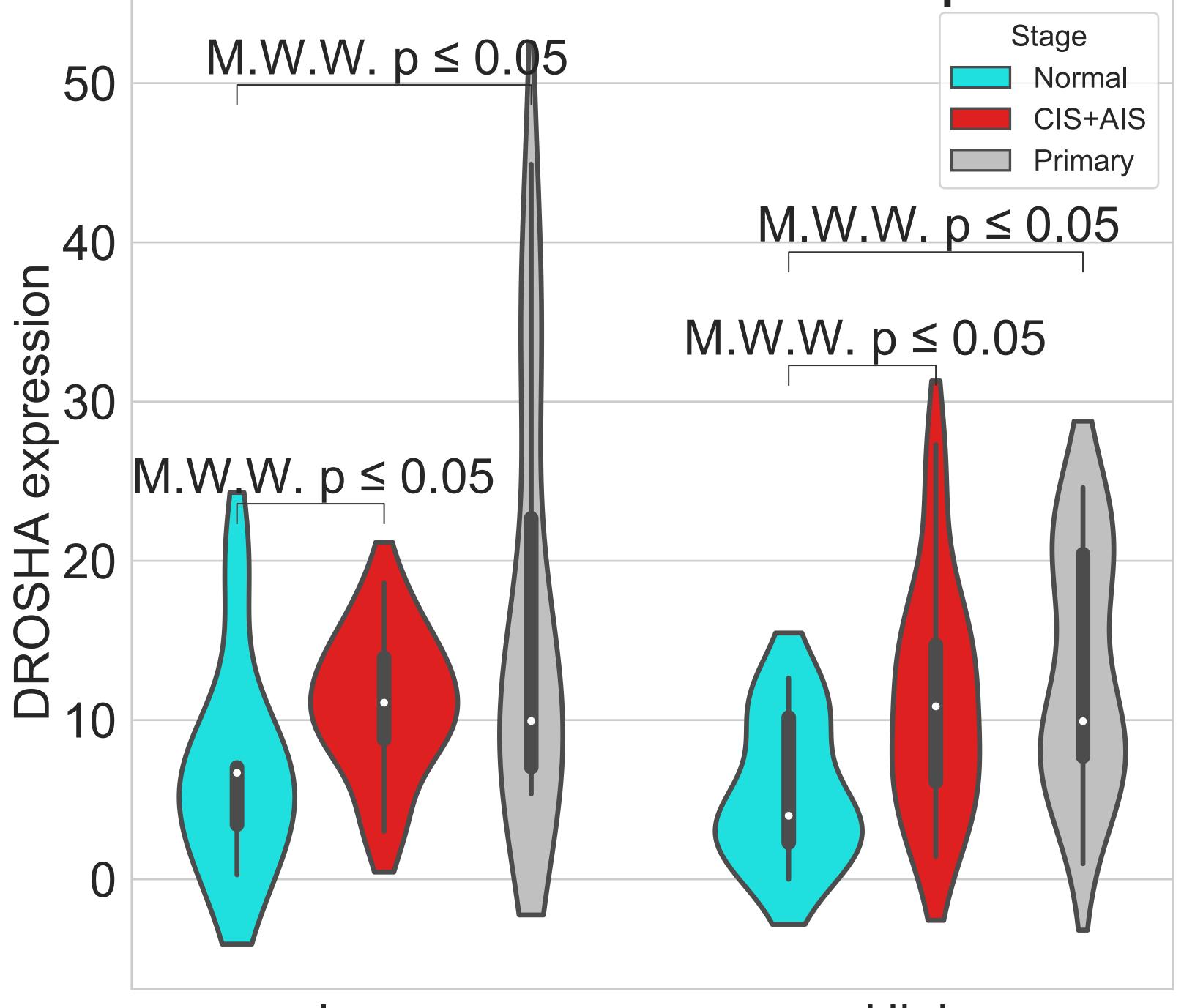
DROSHA: Kruskal-Wallis p=0.030



Lower Higher Mutation Shared Count per TMB