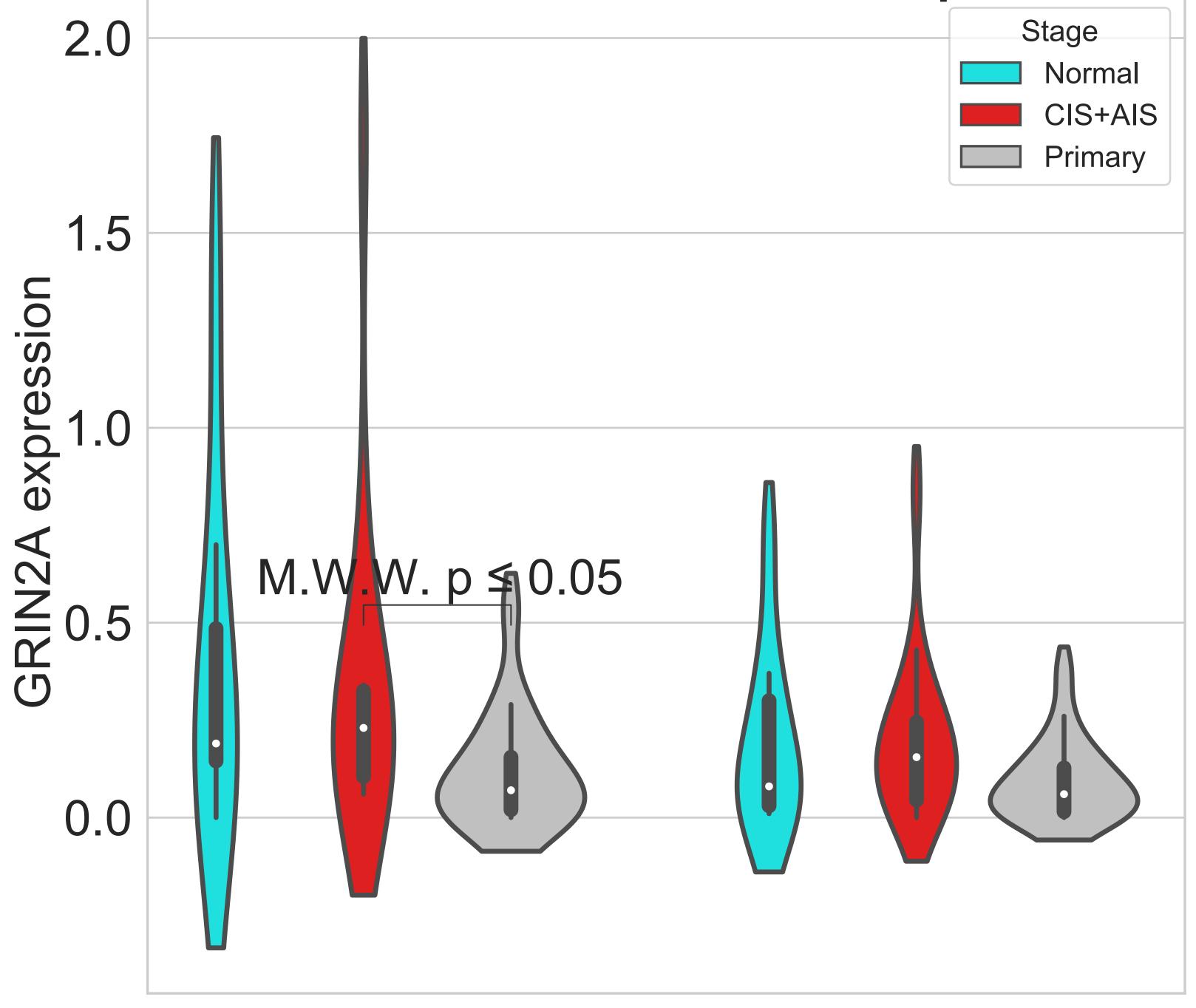
GRIN2A: Kruskal-Wallis p=0.030

Stage
Normal



Lower Higher Mutation Shared Count per TMB