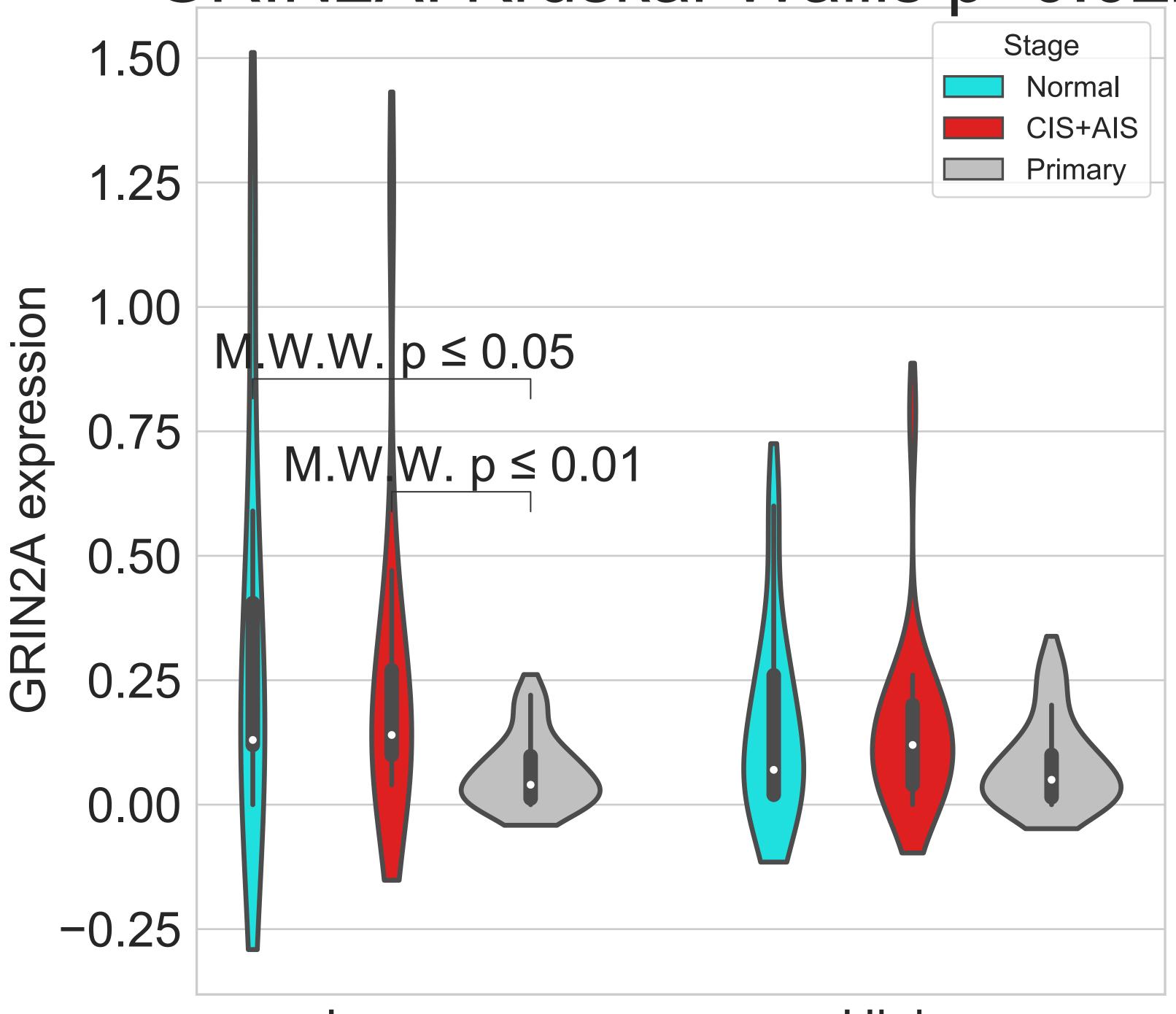
GRIN2A: Kruskal-Wallis p=0.022



Lower Higher Mutation Shared Count (SYN) per TMB