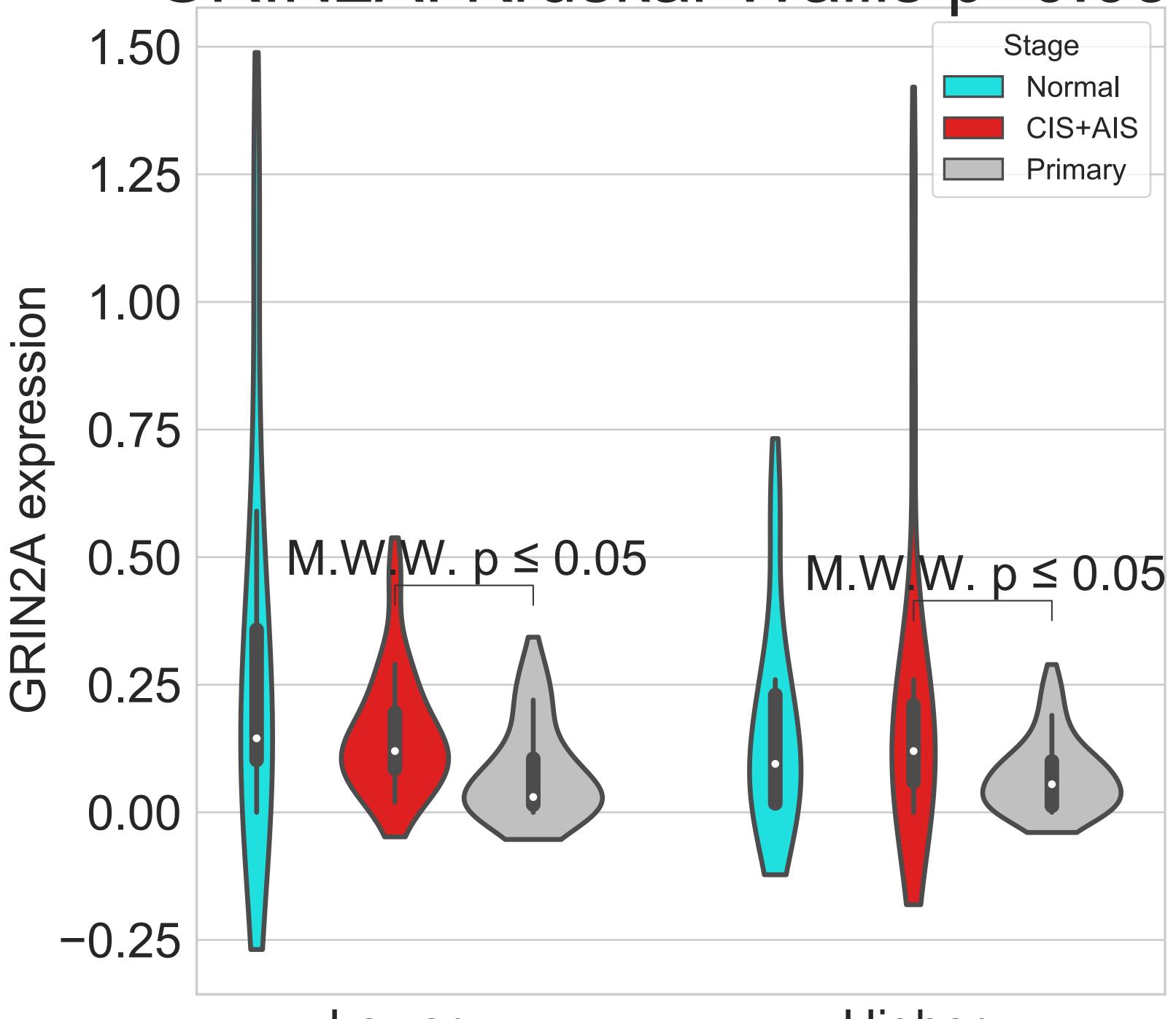
GRIN2A: Kruskal-Wallis p=0.038



Lower
Higher
Mutation Shared Count (SYN)