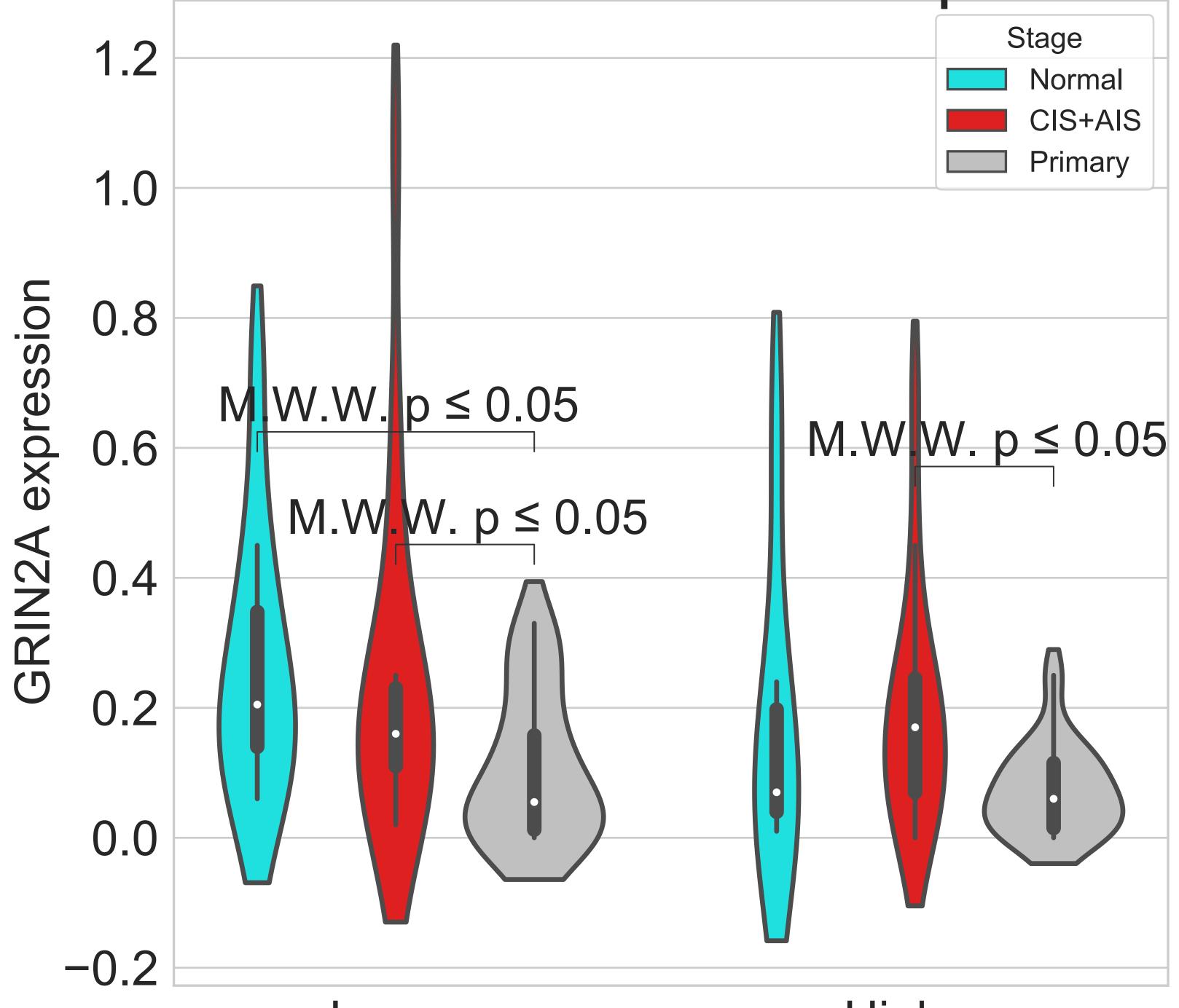
GRIN2A: Kruskal-Wallis p=0.015



Lower Higher Mutation Shared Proportion (Union & SYN)