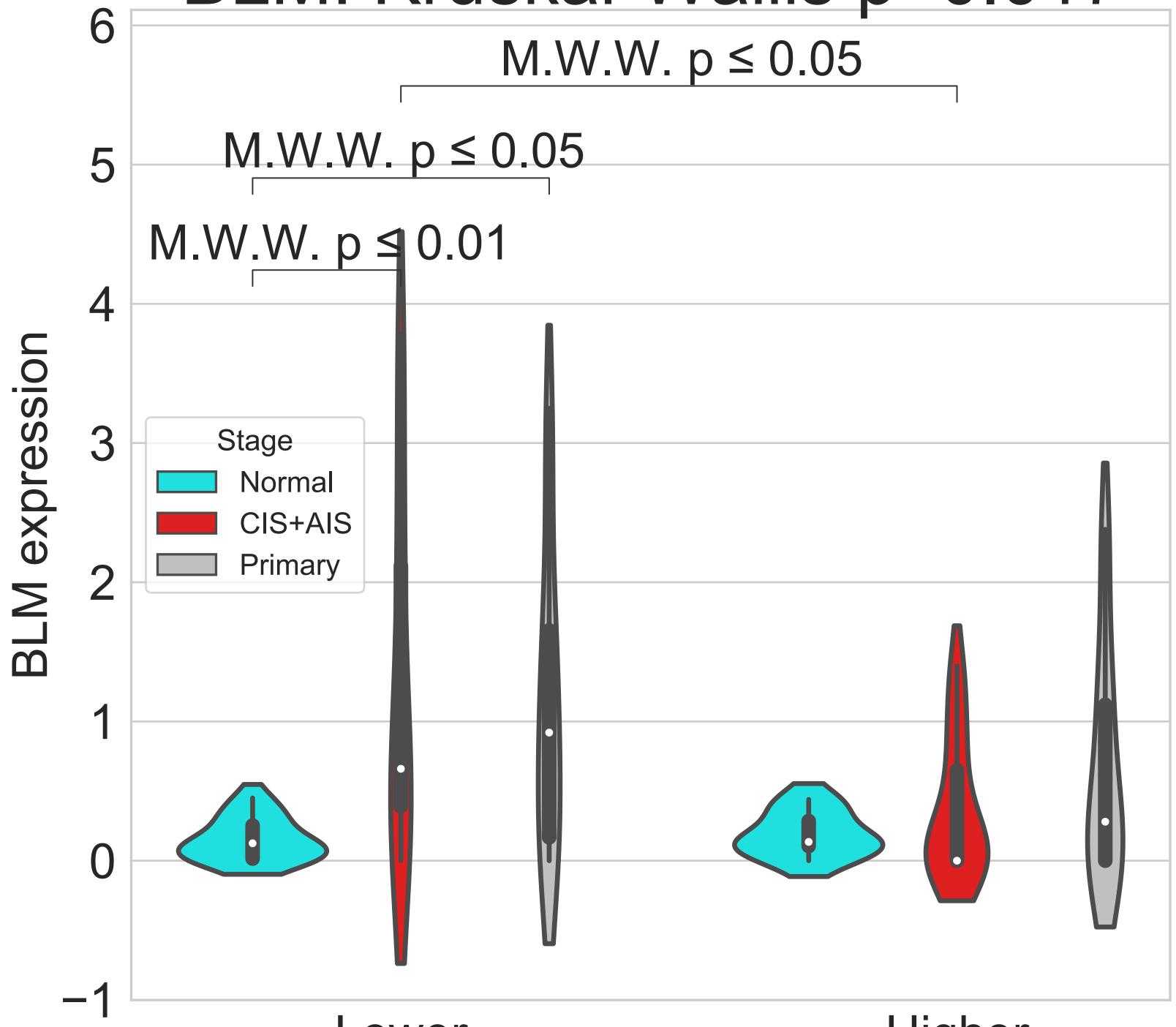
BLM: Kruskal-Wallis p=0.017



Lower Higher Mutation Shared Proportion (SYN)