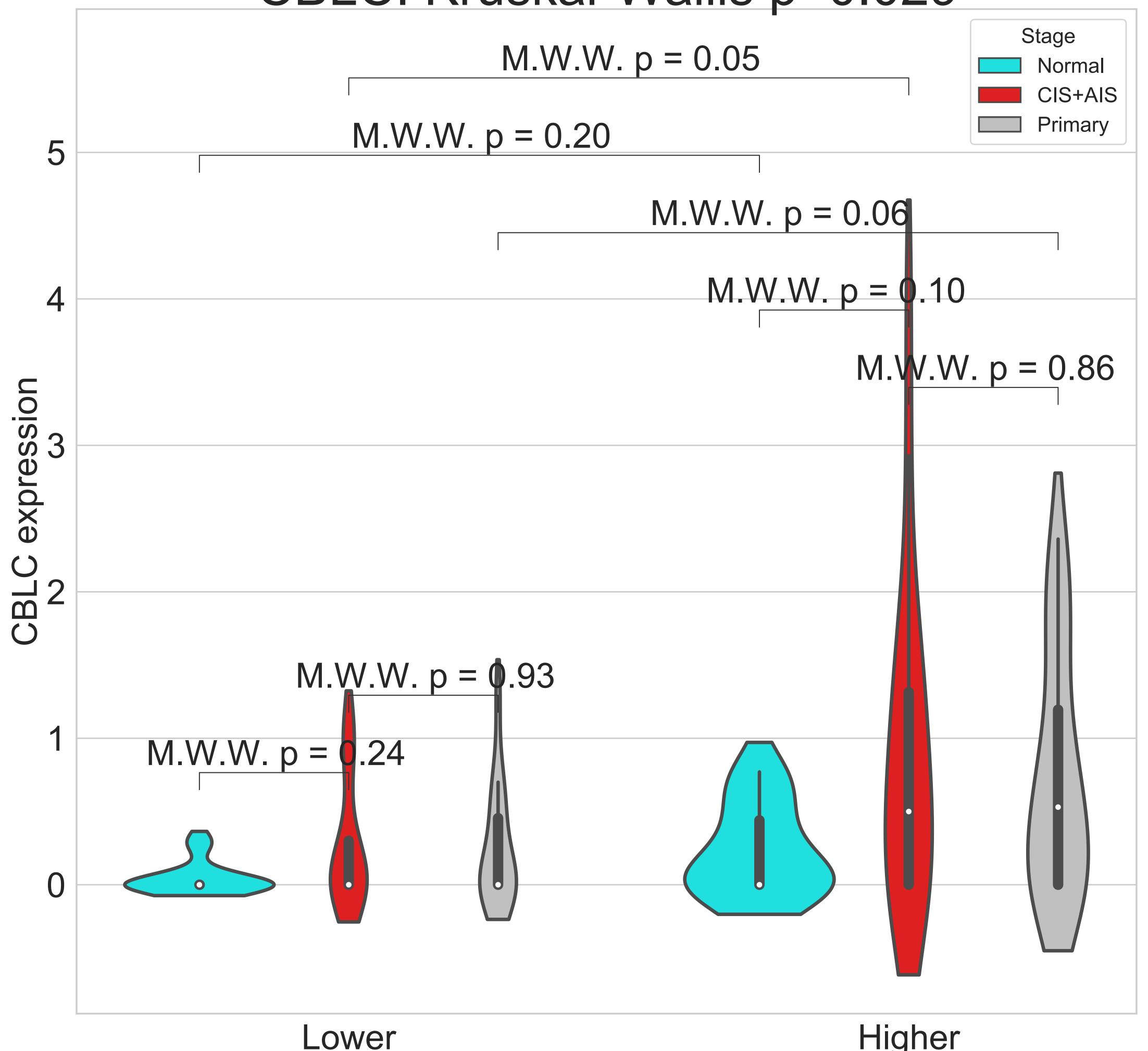
CBLC: Kruskal-Wallis p=0.029



Higher Mutation Shared Proportion (SYN)