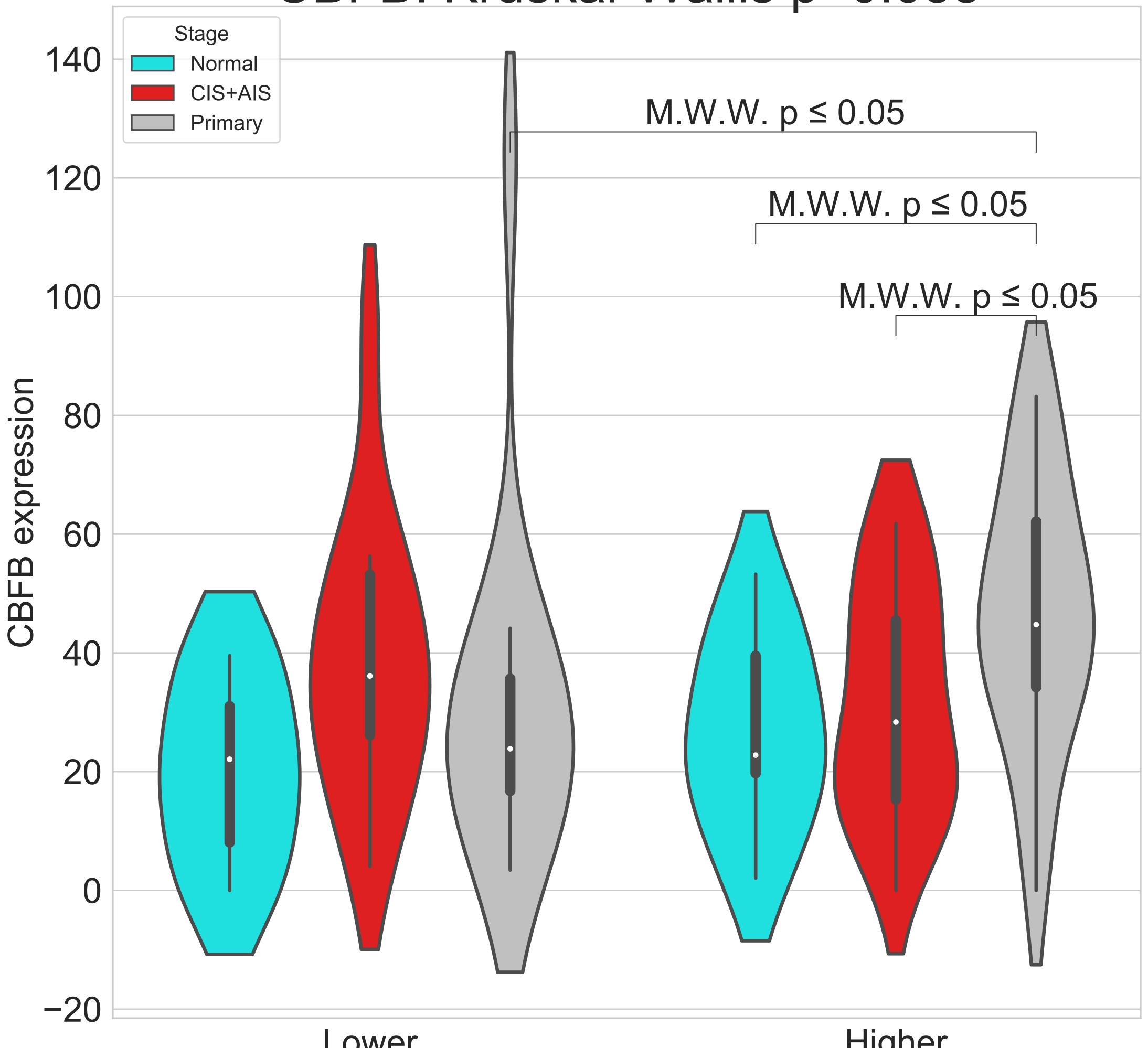
CBFB: Kruskal-Wallis p=0.038



Lower
Mutation Shared Proportion (SYN)