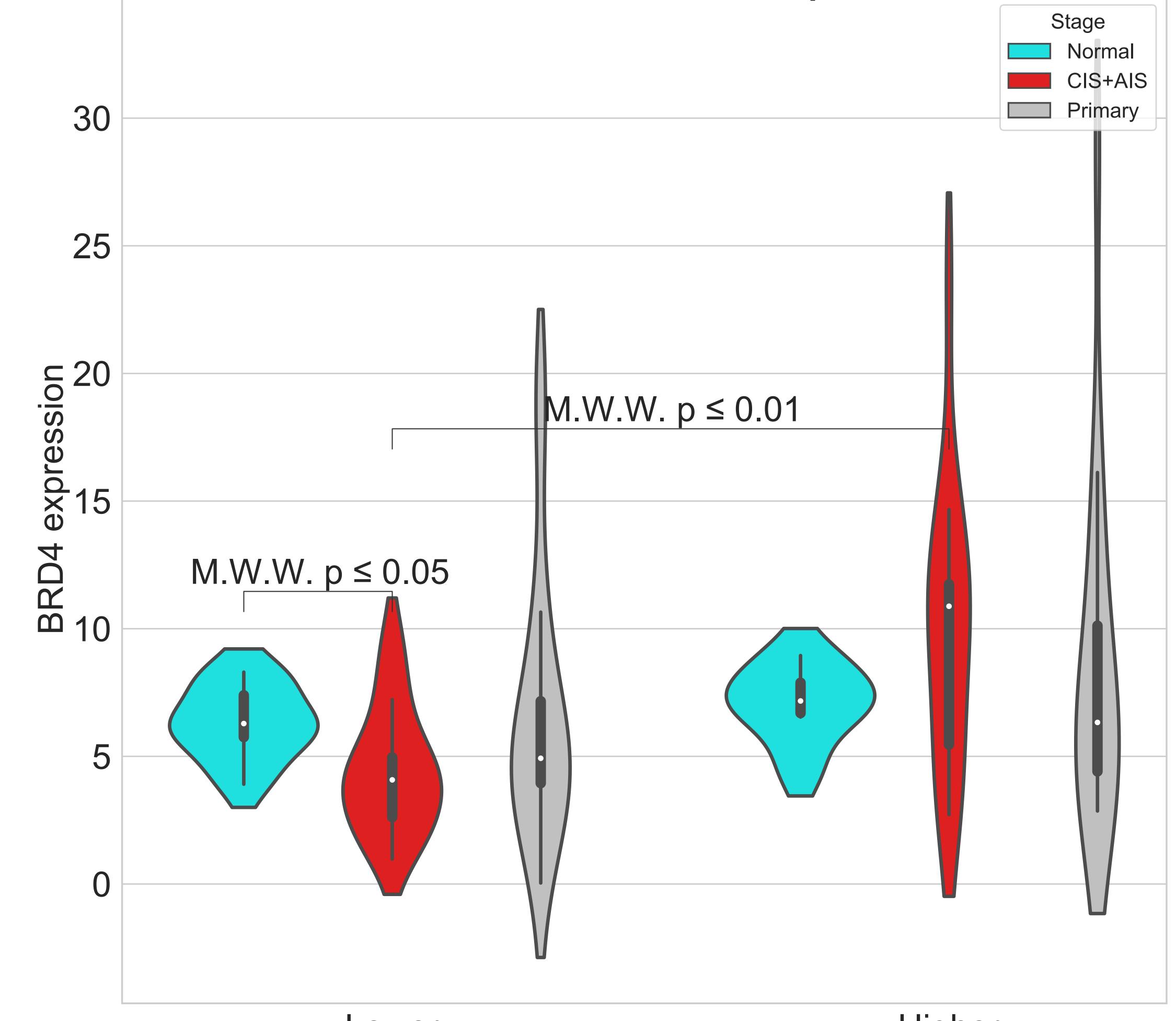
BRD4: Kruskal-Wallis p=0.012



Higher Lower Mutation Shared Proportion (Union & SYN)