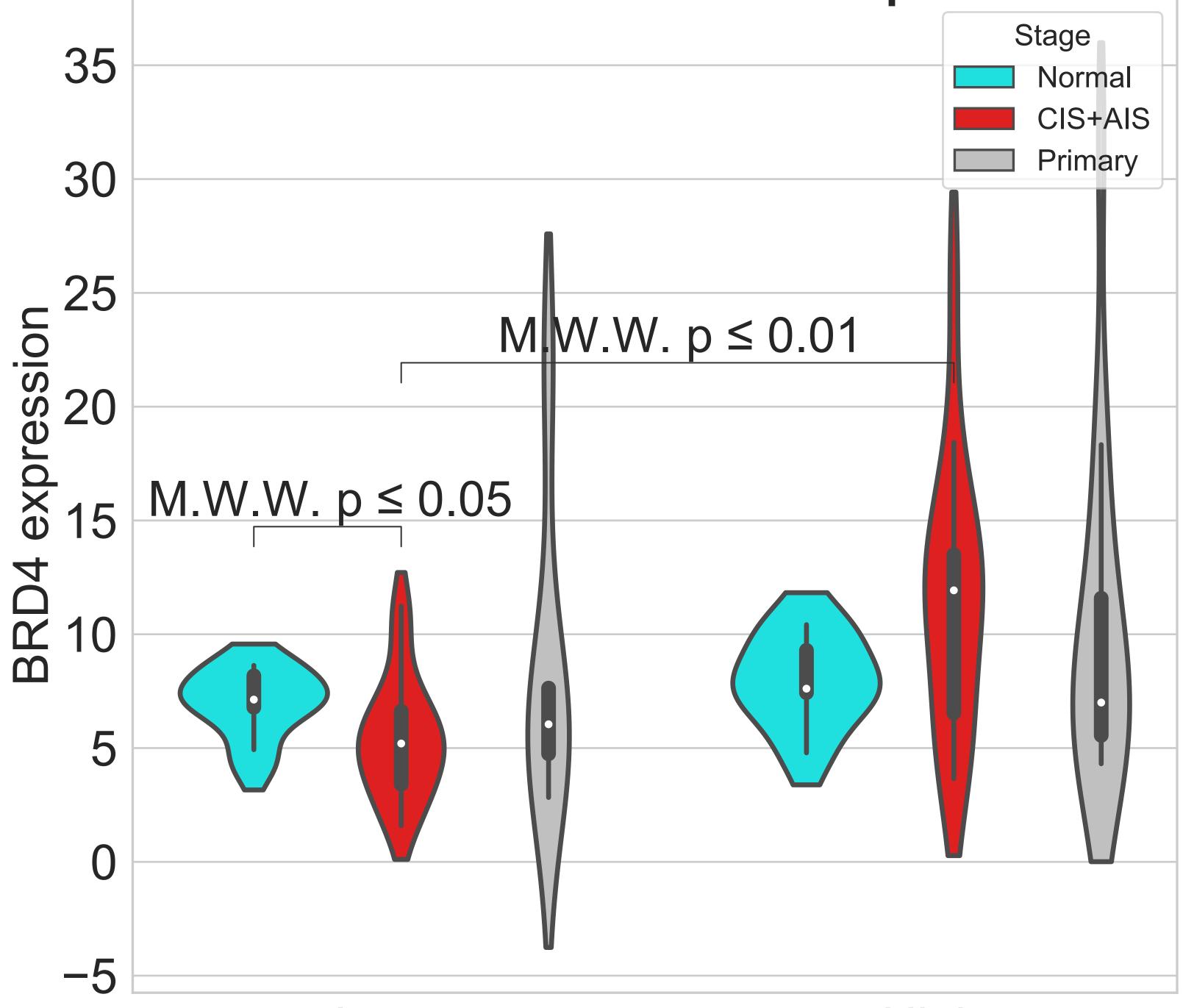
BRD4: Kruskal-Wallis p=0.013



Lower
Mutation Shared Proportion (SYN)