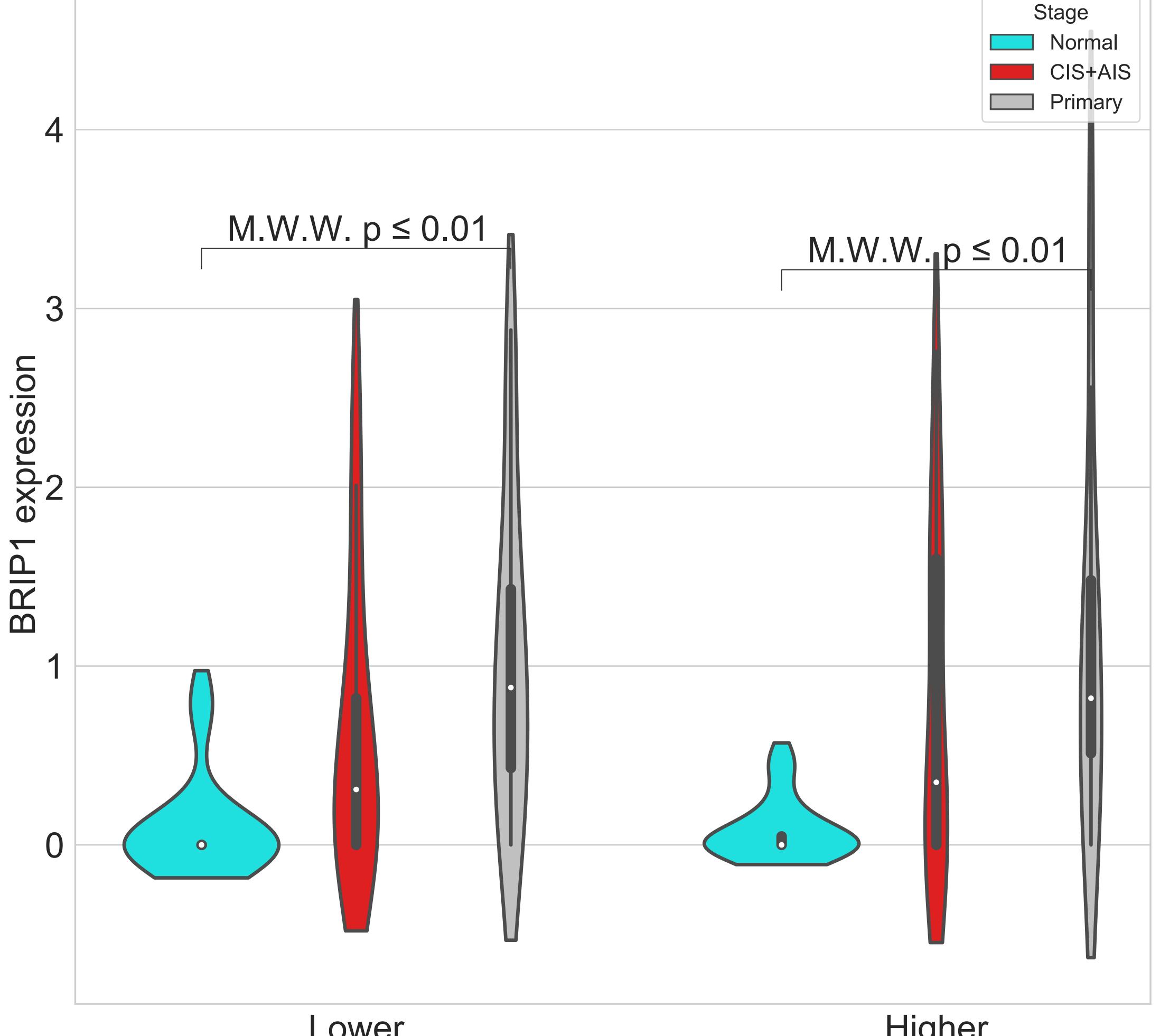
BRIP1: Kruskal-Wallis p=0.006



Higher Lower Mutation Shared Count (SYN)