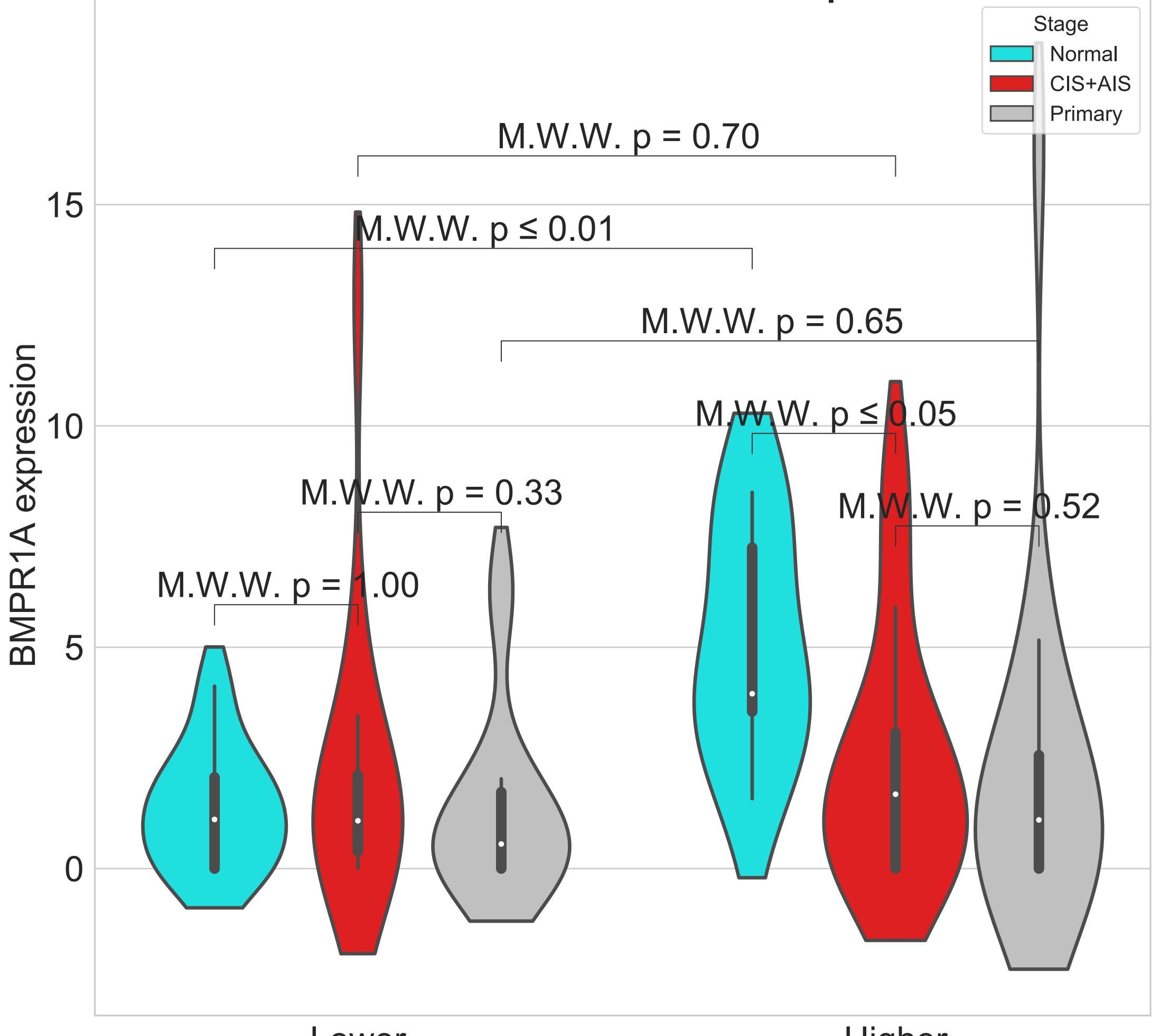
BMPR1A: Kruskal-Wallis p=0.038



Higher Lower Mutation Shared Count (SYN)