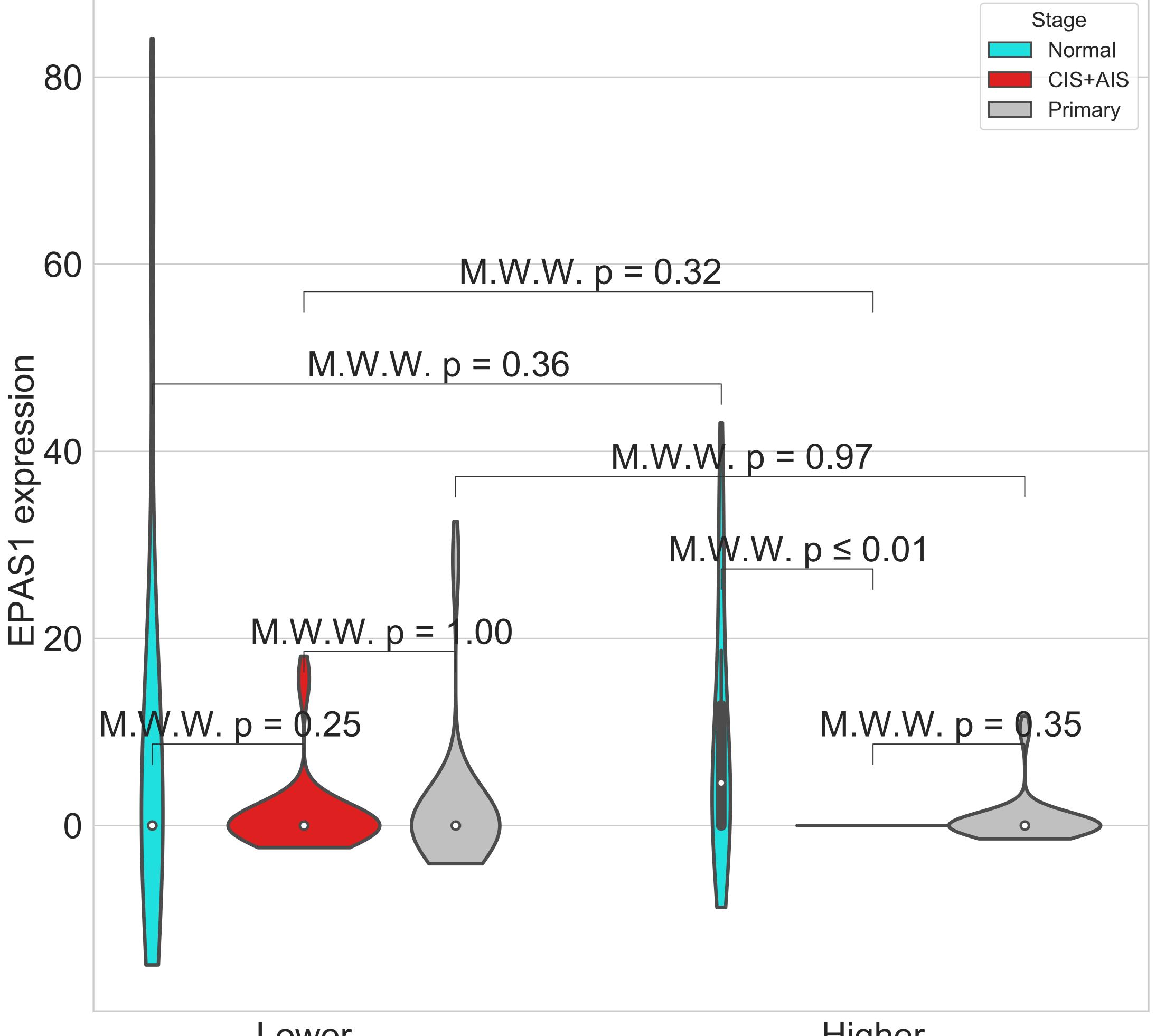
EPAS1: Kruskal-Wallis p=0.002



Higher Lower Mutation Shared Count (SYN)