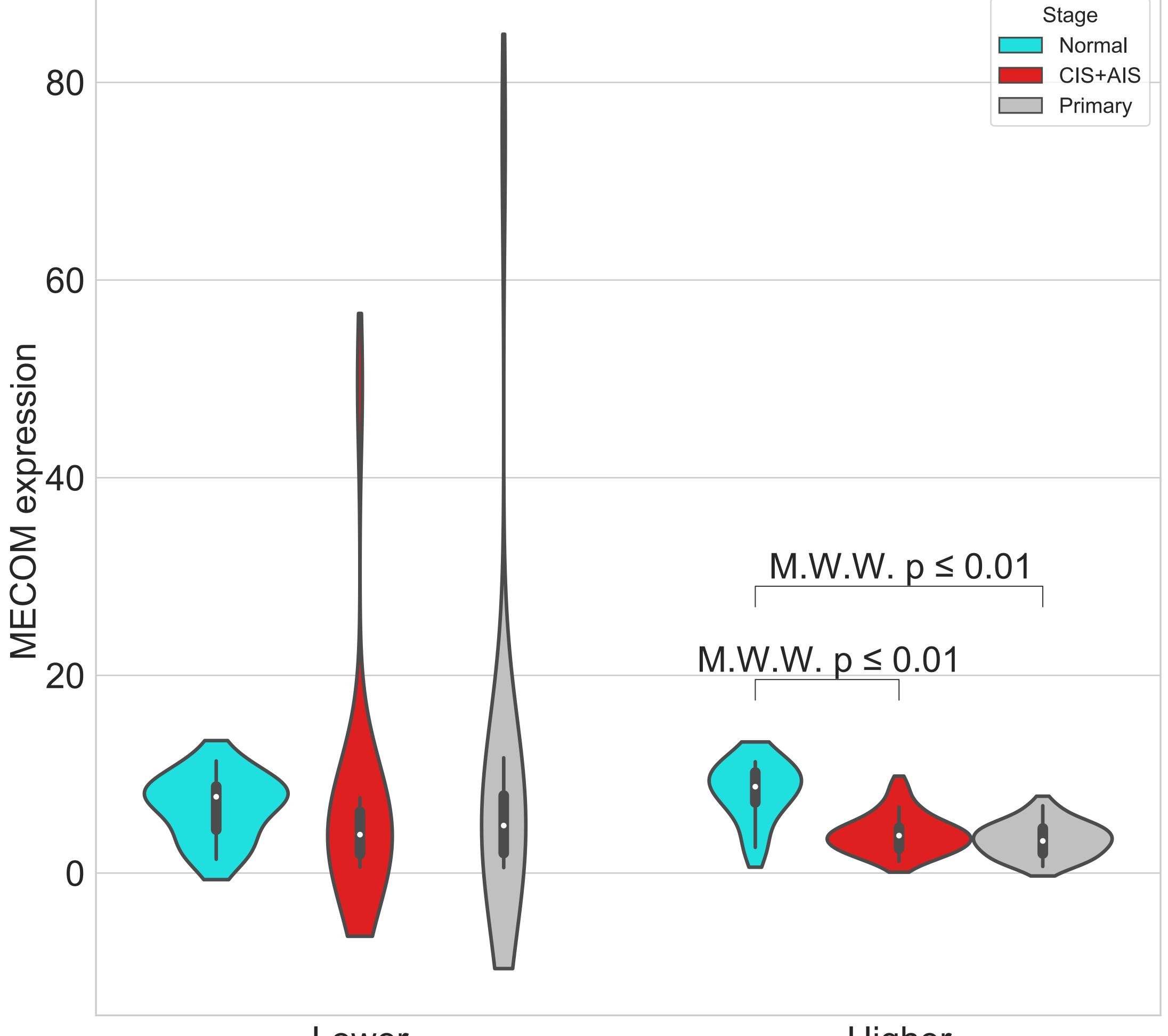
MECOM: Kruskal-Wallis p=0.019



Higher Lower Mutation Shared Count (SYN)