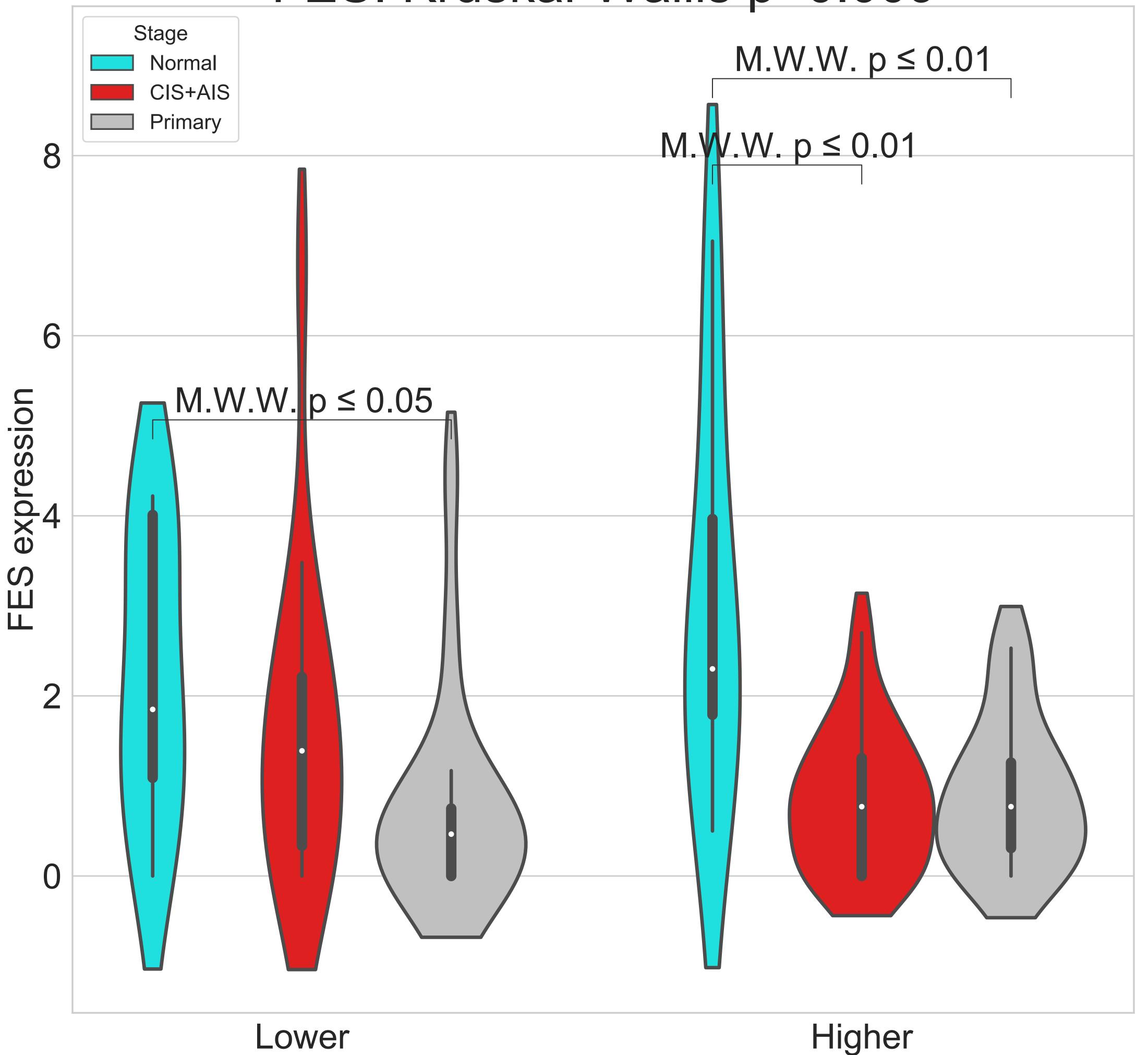
FES: Kruskal-Wallis p=0.009



Higher Mutation Shared Count (SYN)