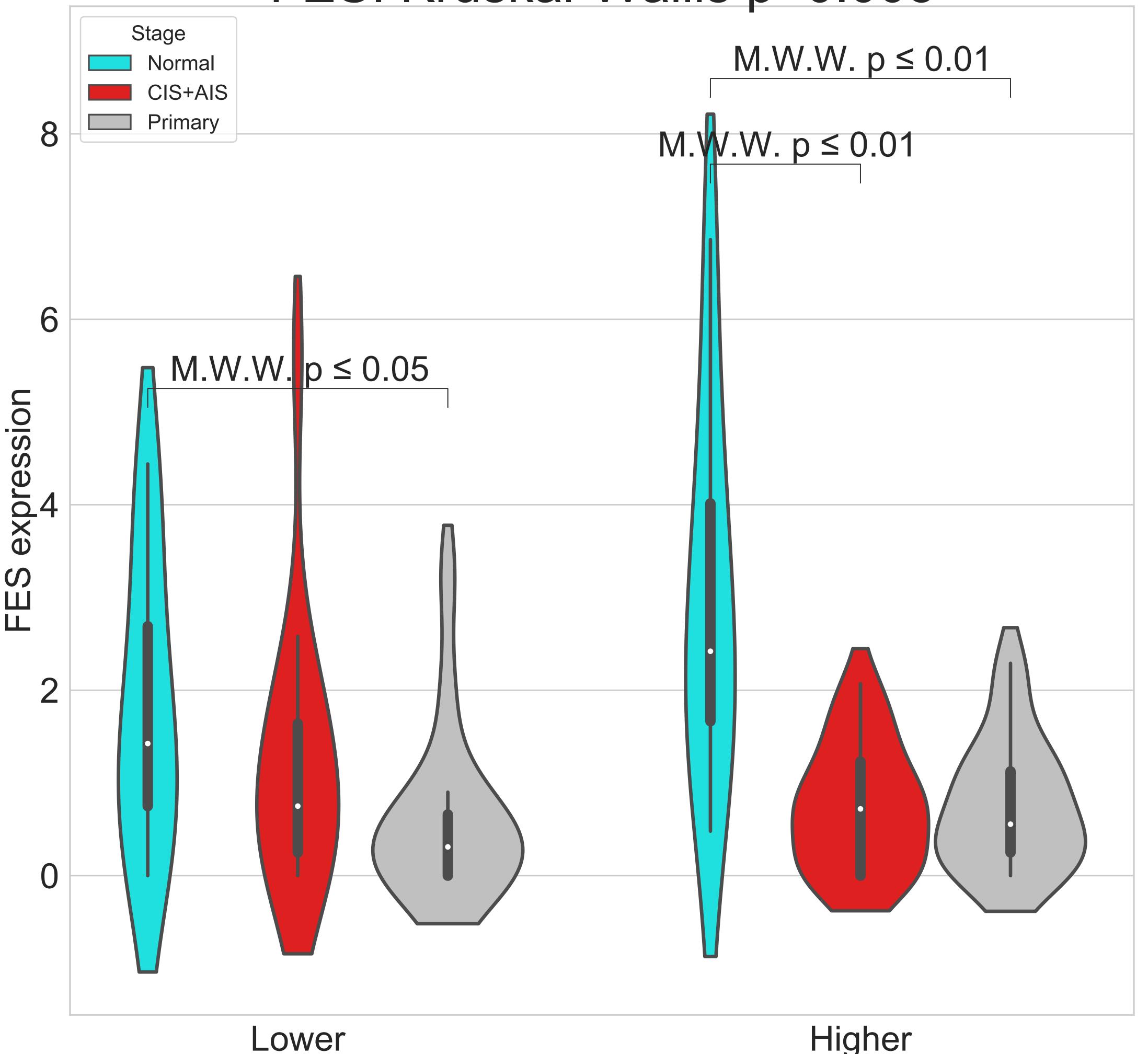
FES: Kruskal-Wallis p=0.005



Higher Mutation Shared Count (SYN)