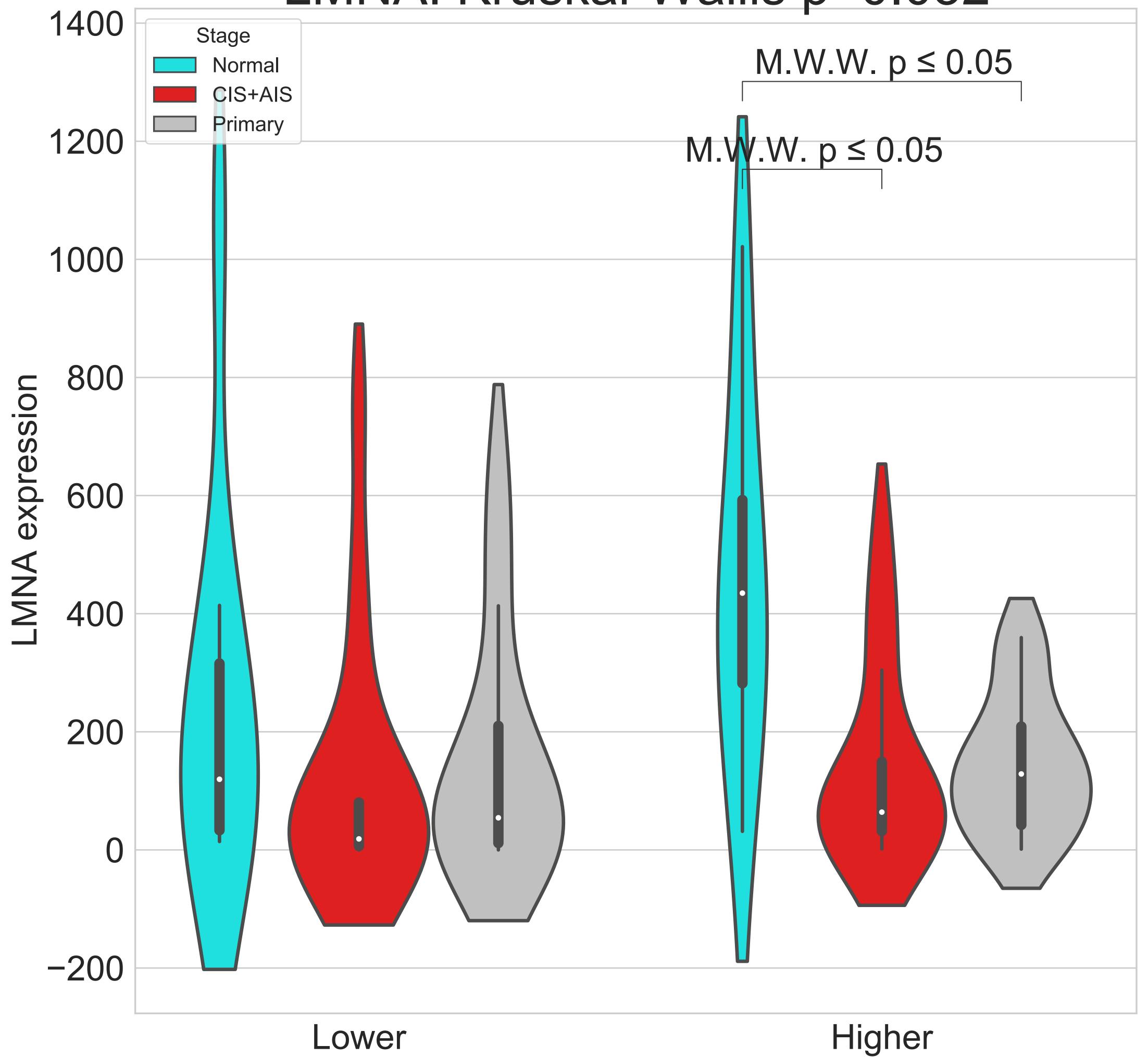
LMNA: Kruskal-Wallis p=0.032



Mutation Shared Count (SYN)