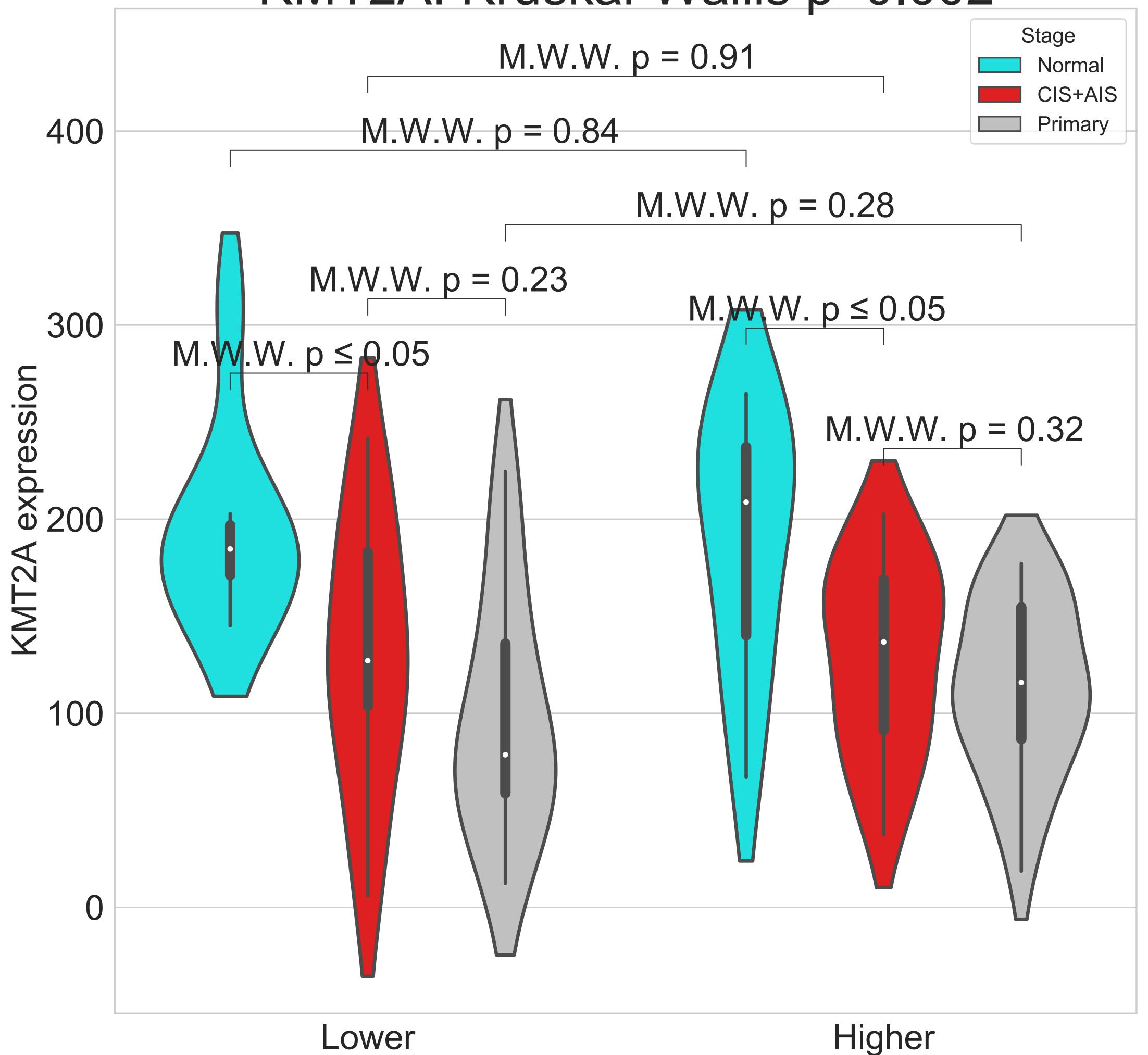
KMT2A: Kruskal-Wallis p=0.002



Mutation Shared Proportion (SYN)