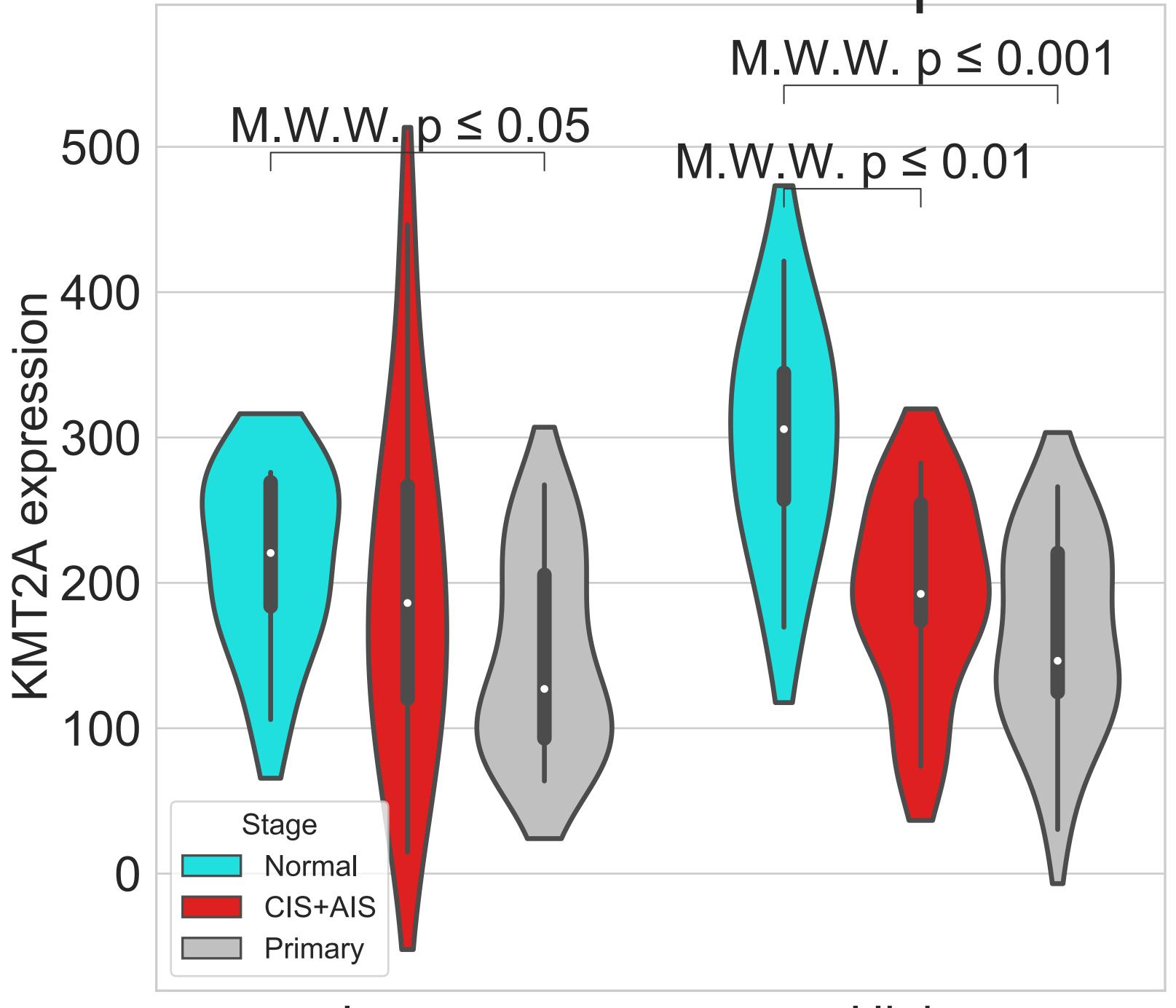
KMT2A: Kruskal-Wallis p=0.008



Lower
Mutation Shared Count (SYN)