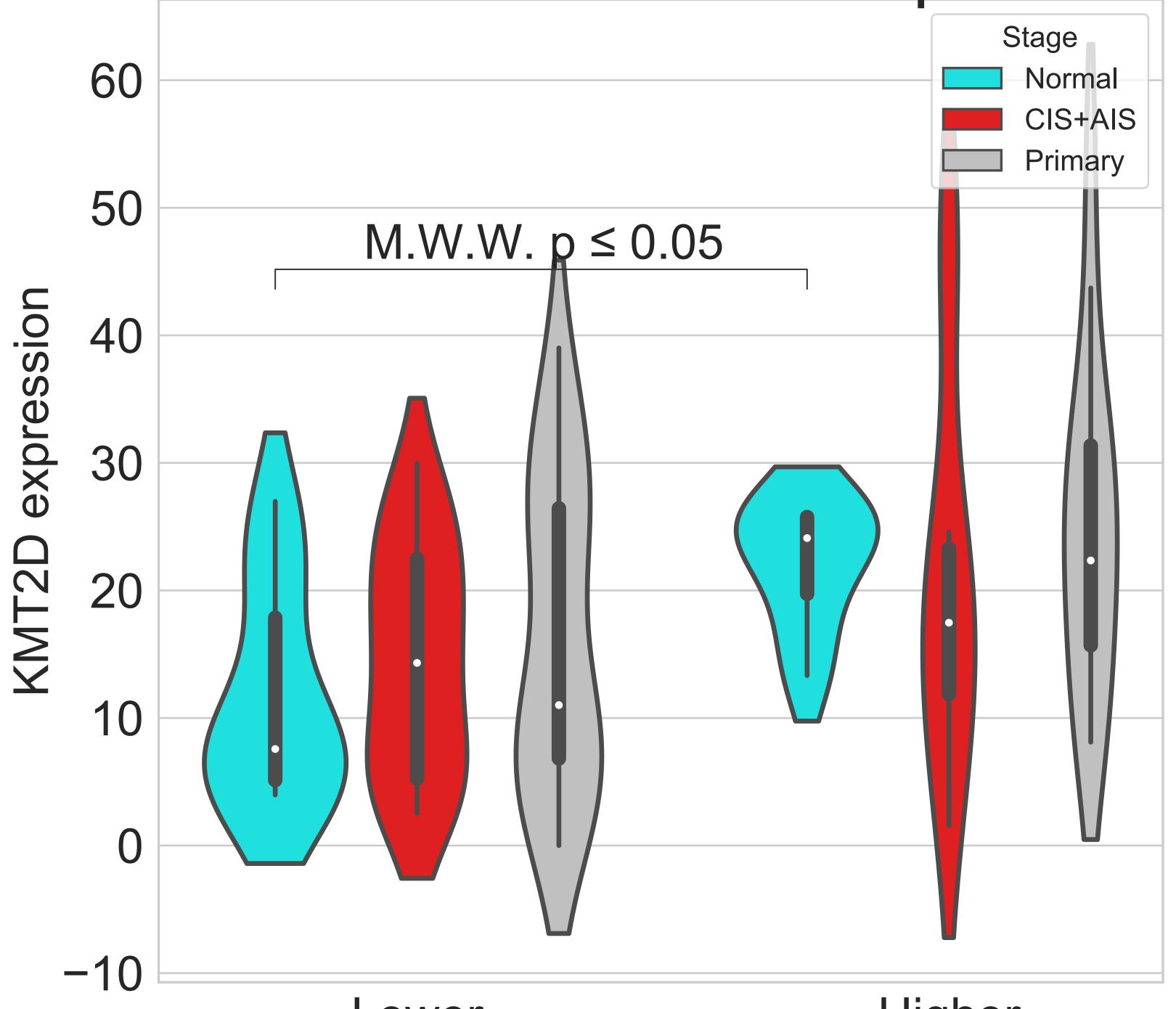
KMT2D: Kruskal-Wallis p=0.049



Lower
Mutation Shared Proportion (SYN)