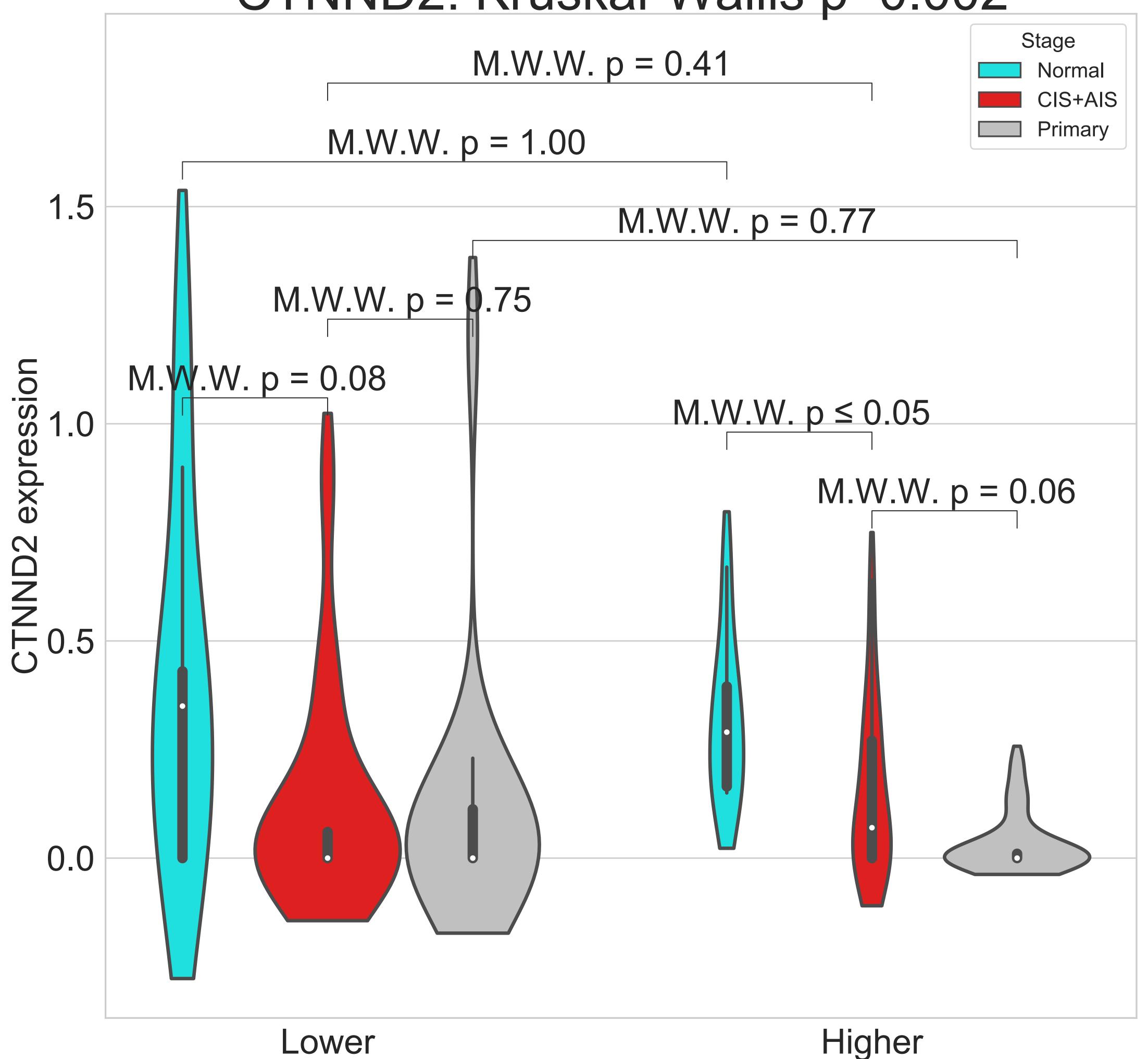
CTNND2: Kruskal-Wallis p=0.002



r Higher Mutation Shared Count (SYN)