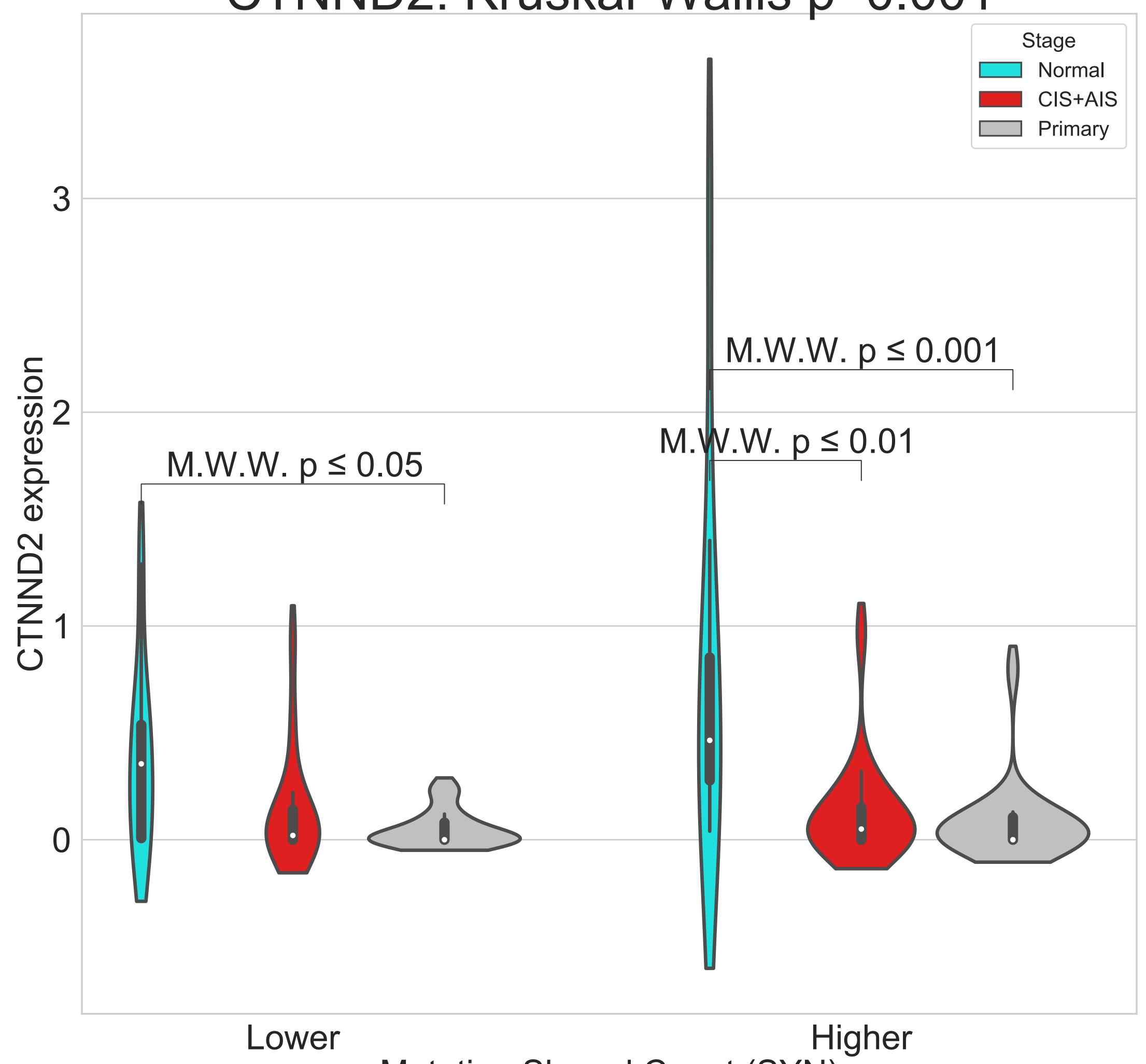
CTNND2: Kruskal-Wallis p=0.001



Mutation Shared Count (SYN)