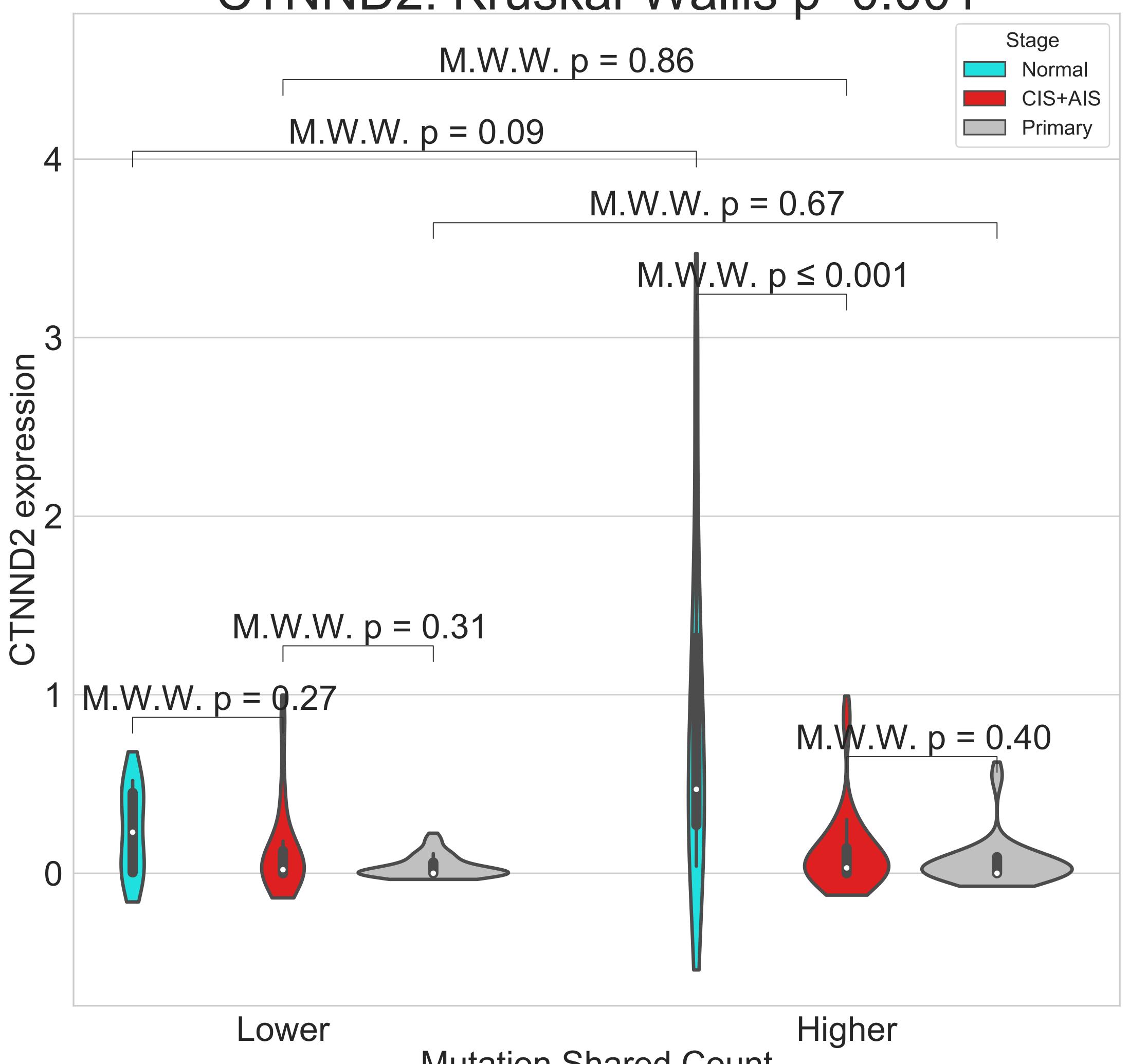
CTNND2: Kruskal-Wallis p=0.001



**Mutation Shared Count**