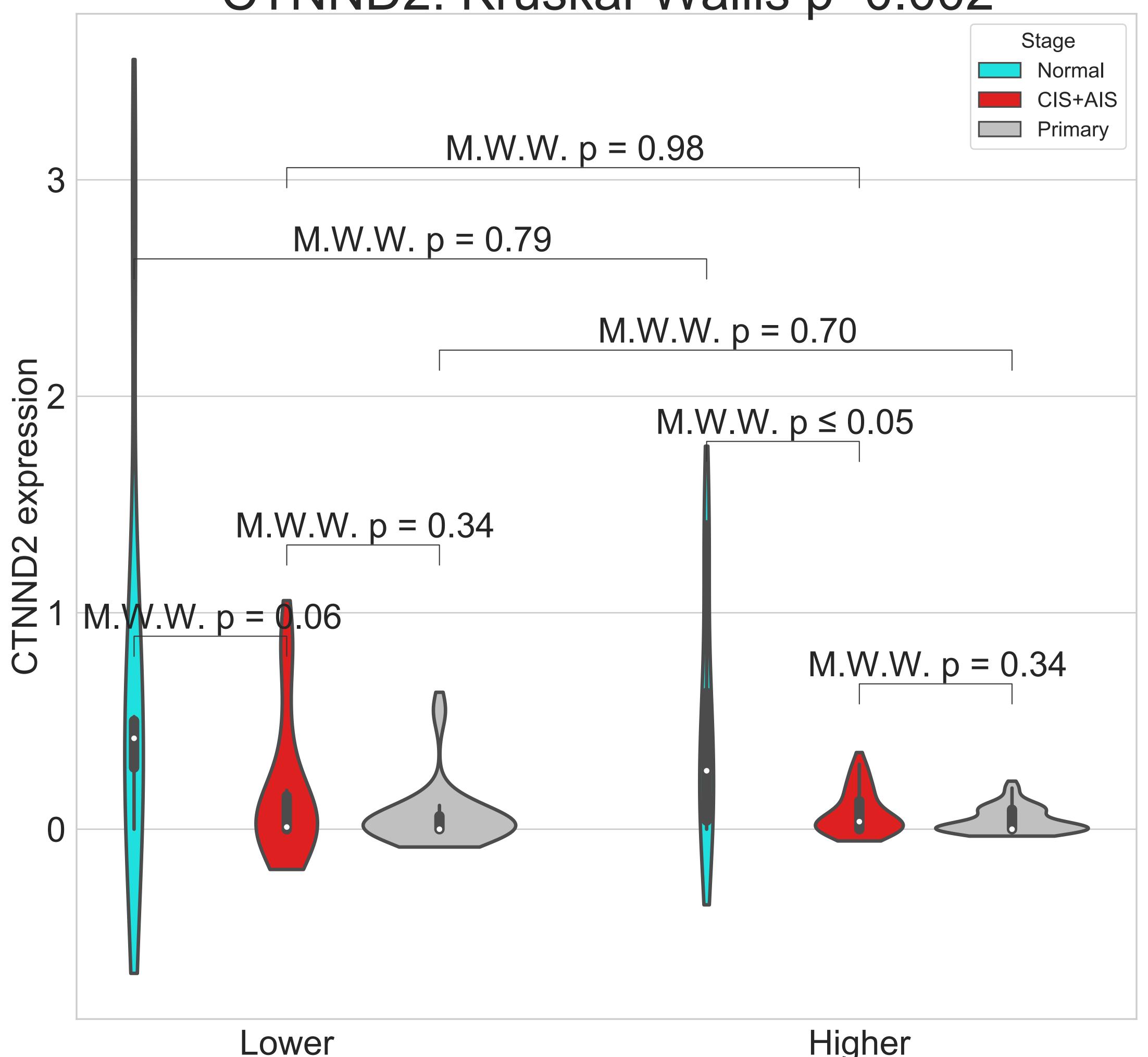
CTNND2: Kruskal-Wallis p=0.002



Higher Mutation Shared Proportion