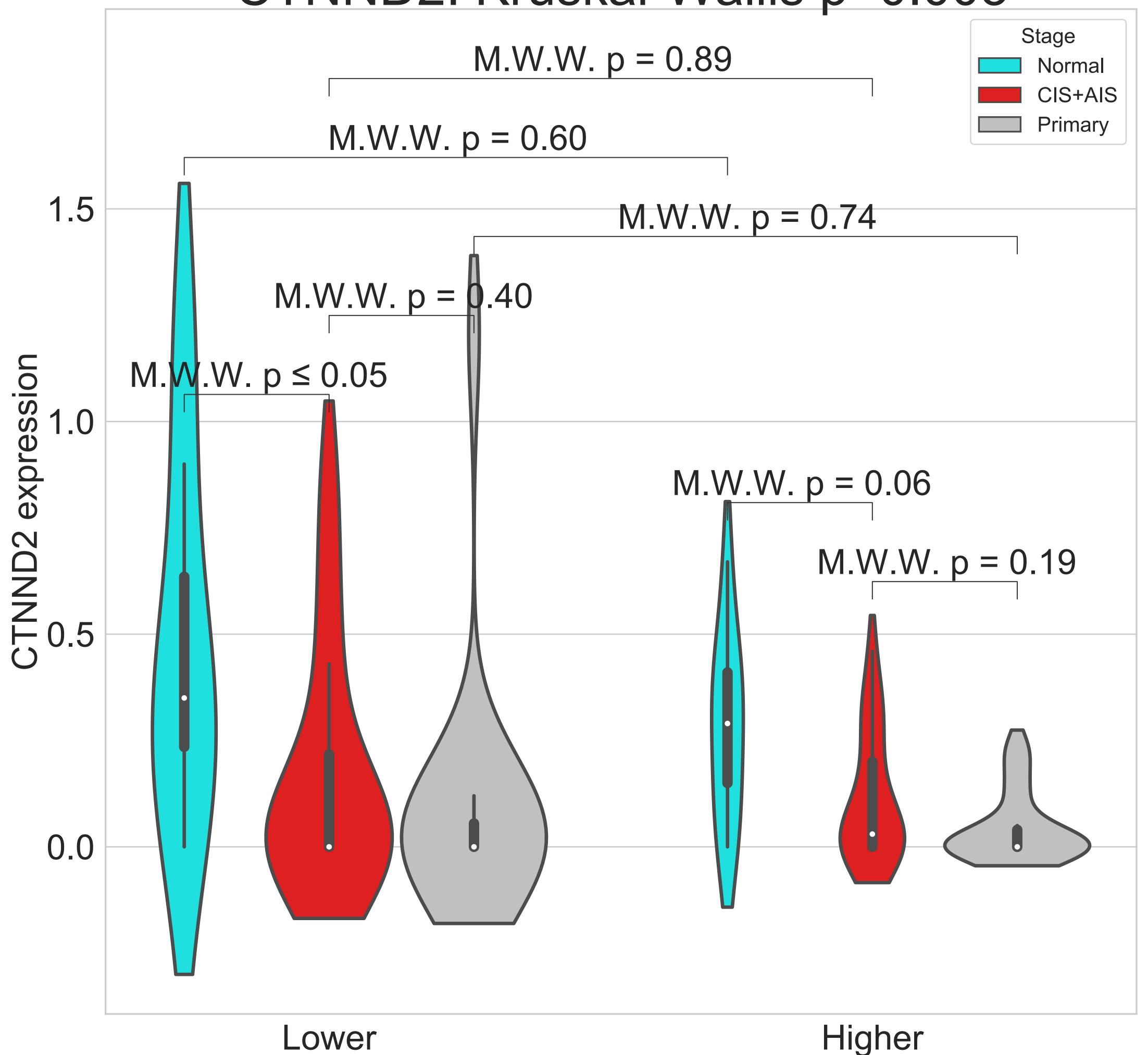
CTNND2: Kruskal-Wallis p=0.003



Higher Mutation Shared Proportion