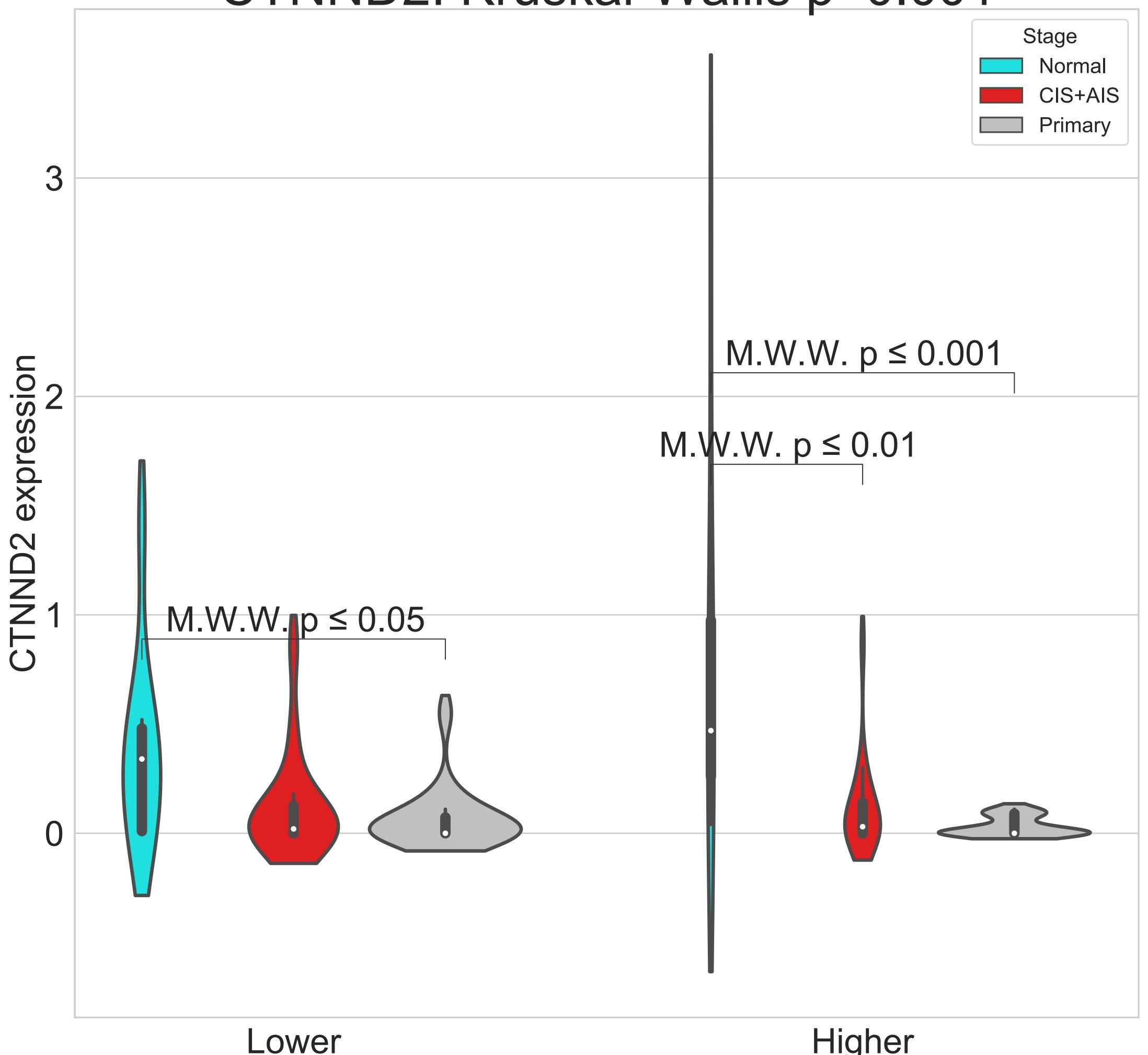
CTNND2: Kruskal-Wallis p=0.001



ver
Mutation Shared Count (SYN)