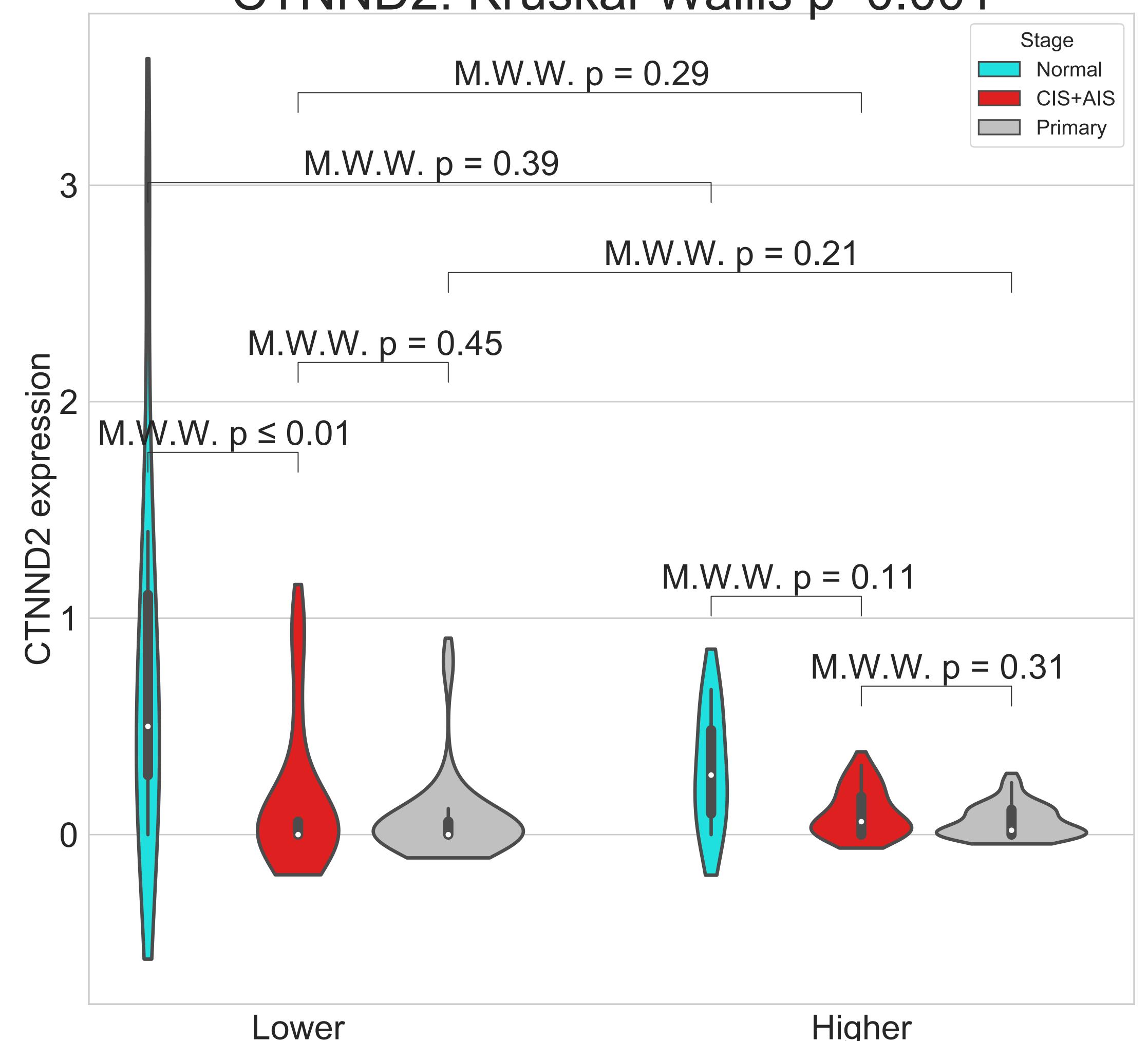
CTNND2: Kruskal-Wallis p=0.001



wer
Mutation Shared Proportion (SYN)