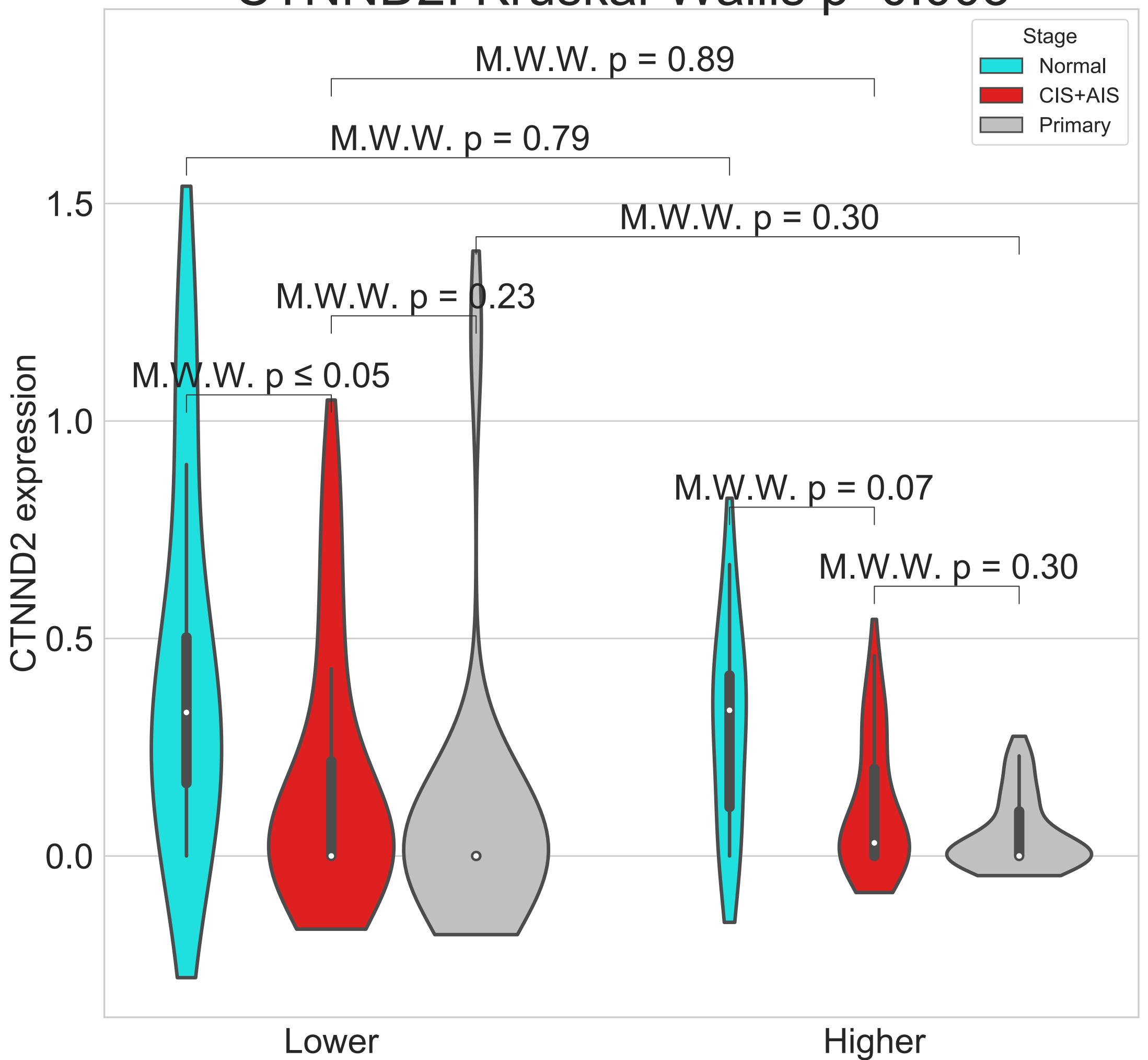
CTNND2: Kruskal-Wallis p=0.003



Mutation Shared Proportion (Union & SYN)