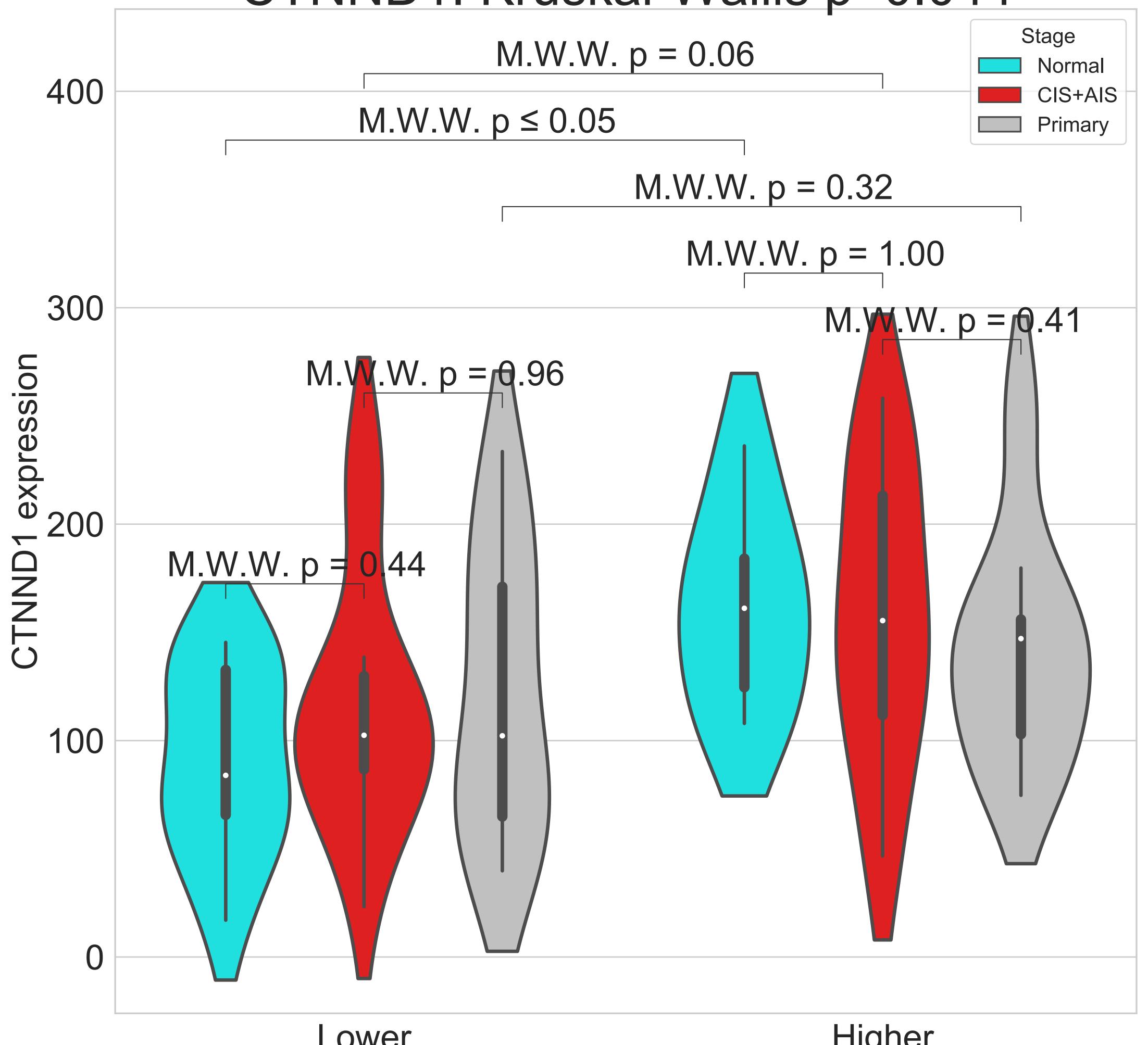
CTNND1: Kruskal-Wallis p=0.044



Lower
Mutation Shared Proportion (SYN)