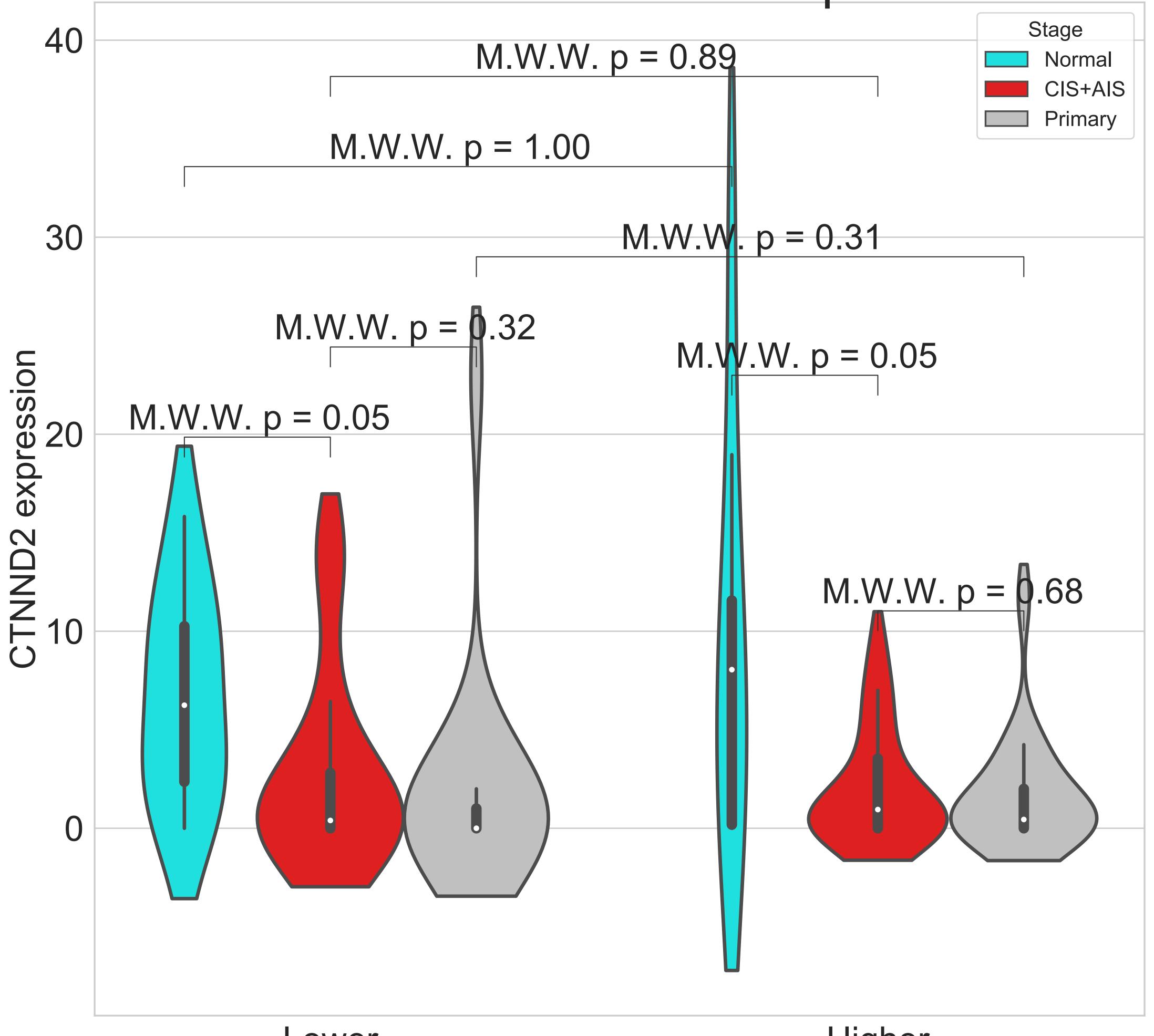
CTNND2: Kruskal-Wallis p=0.011



Lower
Mutation Shared Proportion (Union)