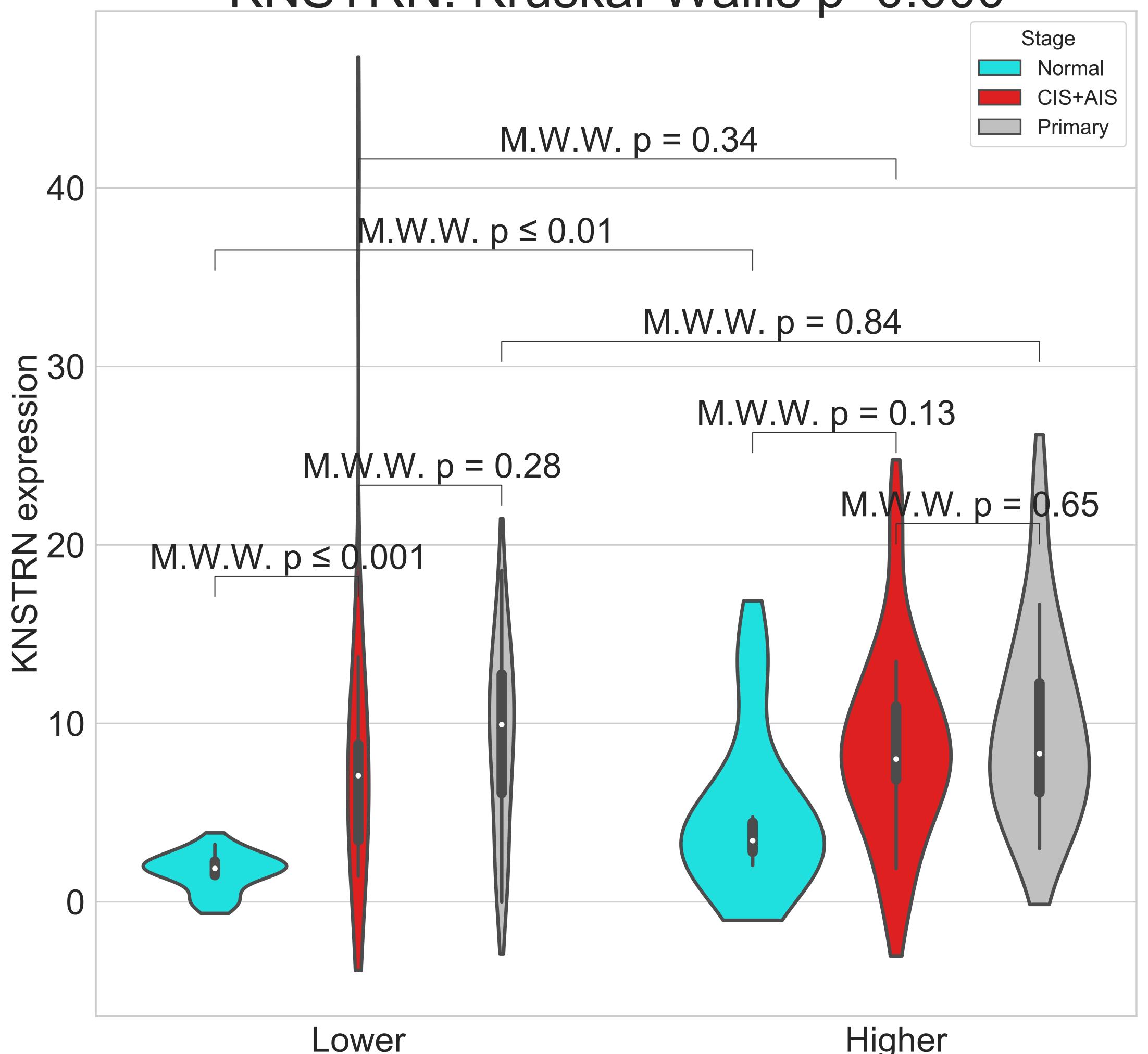
KNSTRN: Kruskal-Wallis p=0.000



Wer Higher Mutation Shared Proportion (SYN)