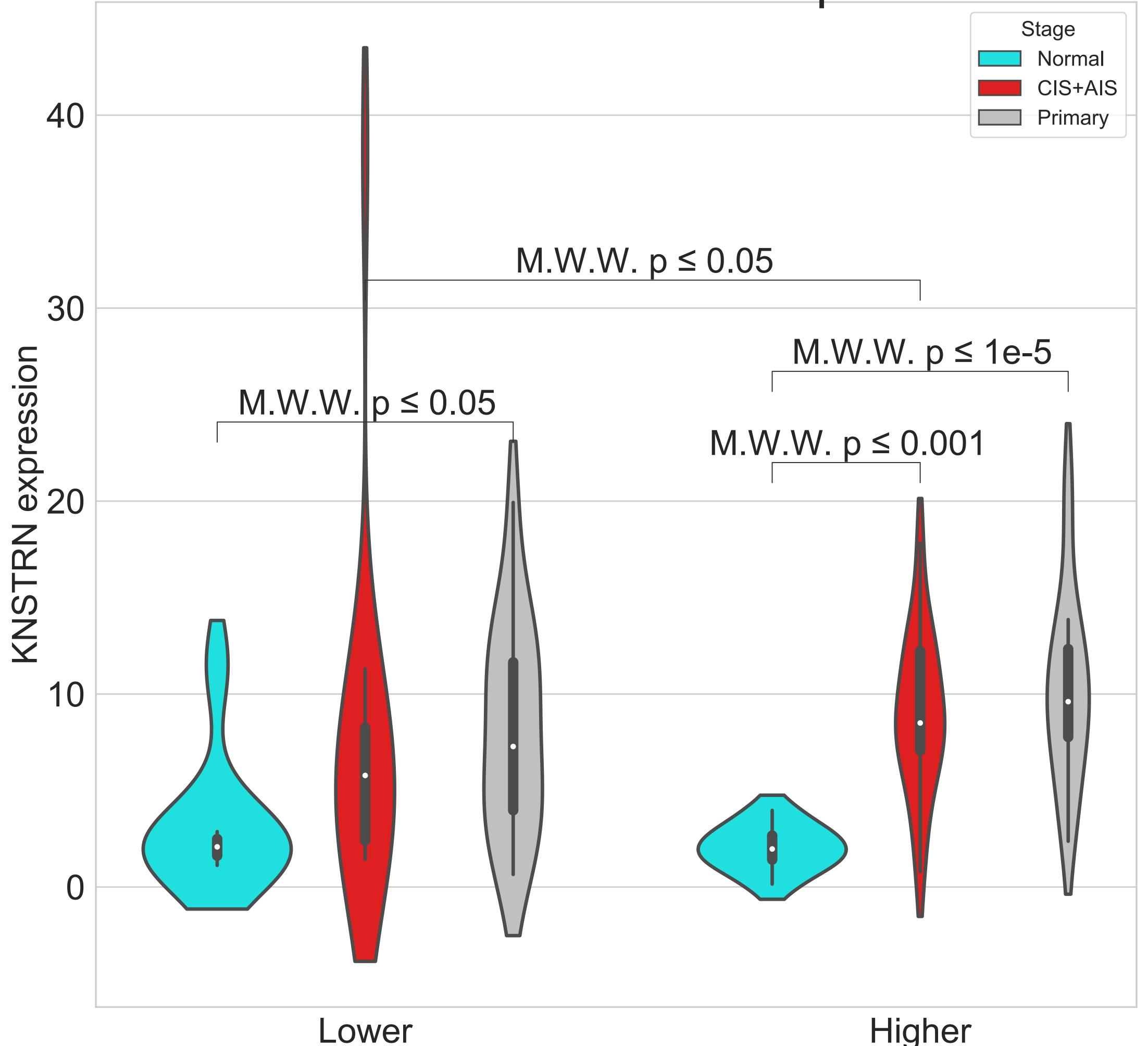
KNSTRN: Kruskal-Wallis p=0.000



Higher Mutation Shared Count (SYN)