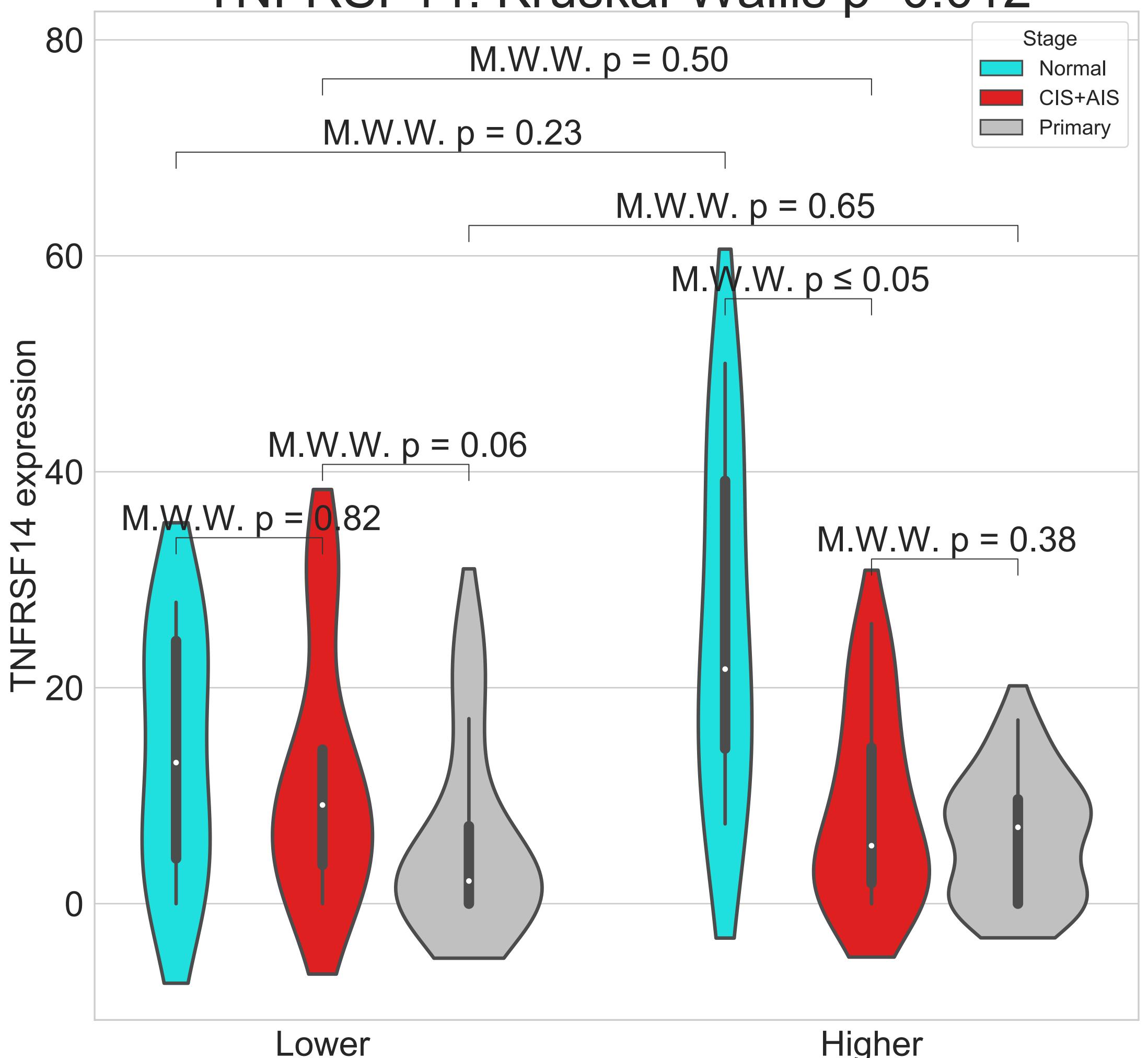
TNFRSF14: Kruskal-Wallis p=0.012



Higher Mutation Shared Count (SYN)