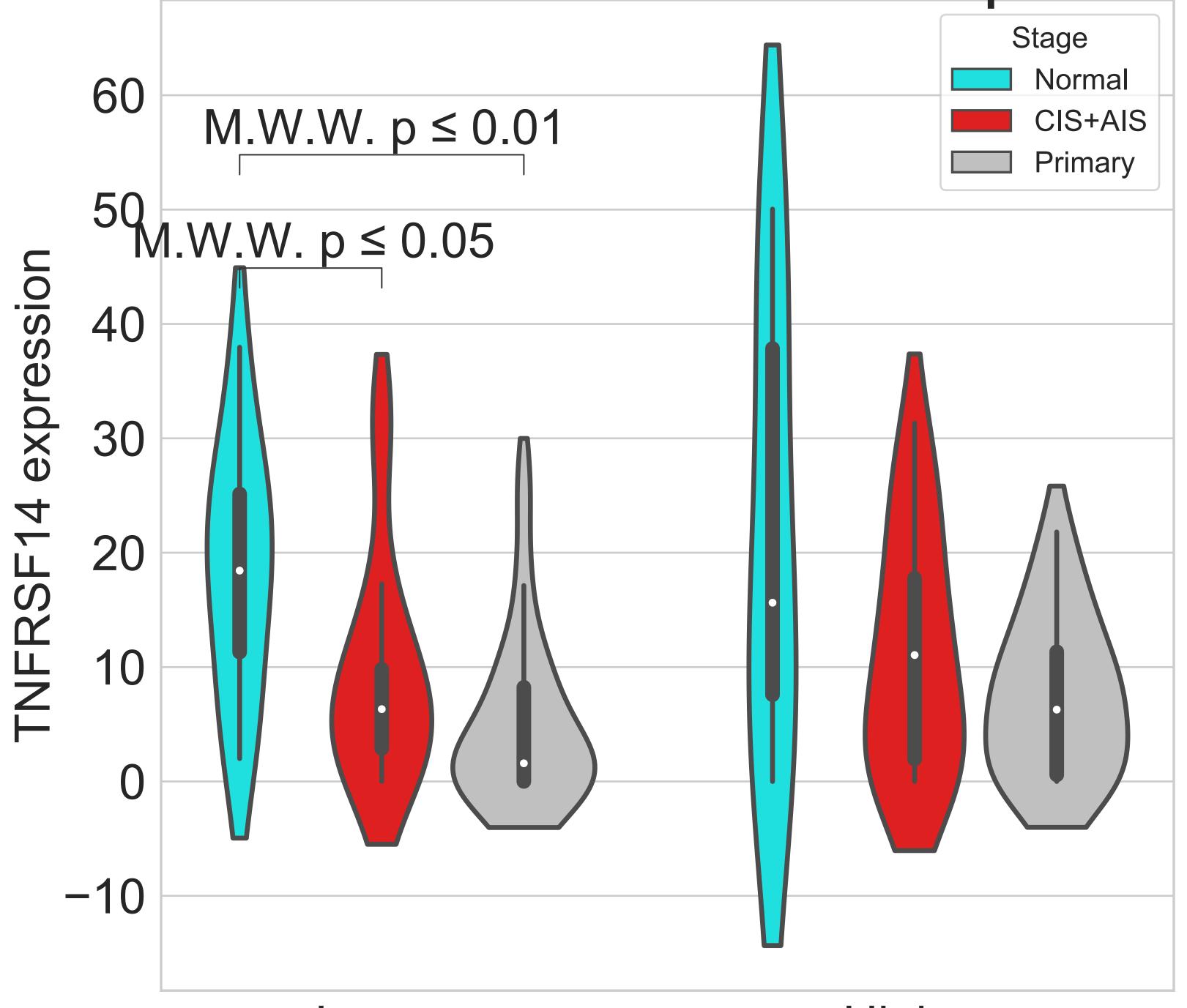
TNFRSF14: Kruskal-Wallis p=0.017



Lower Higher Mutation Shared Proportion (Union & SYN)