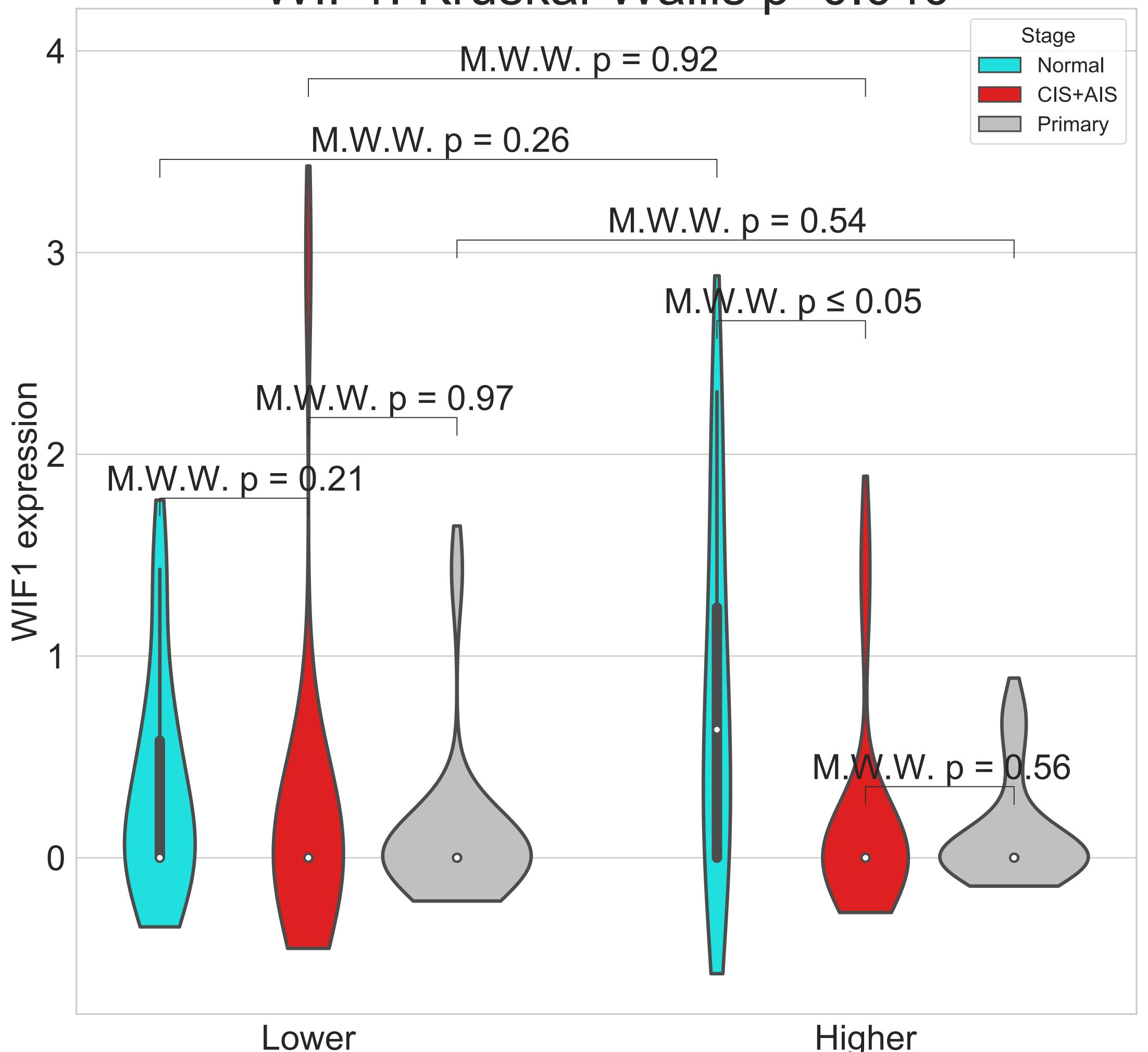
WIF1: Kruskal-Wallis p=0.040



Higher Mutation Shared Count (SYN)