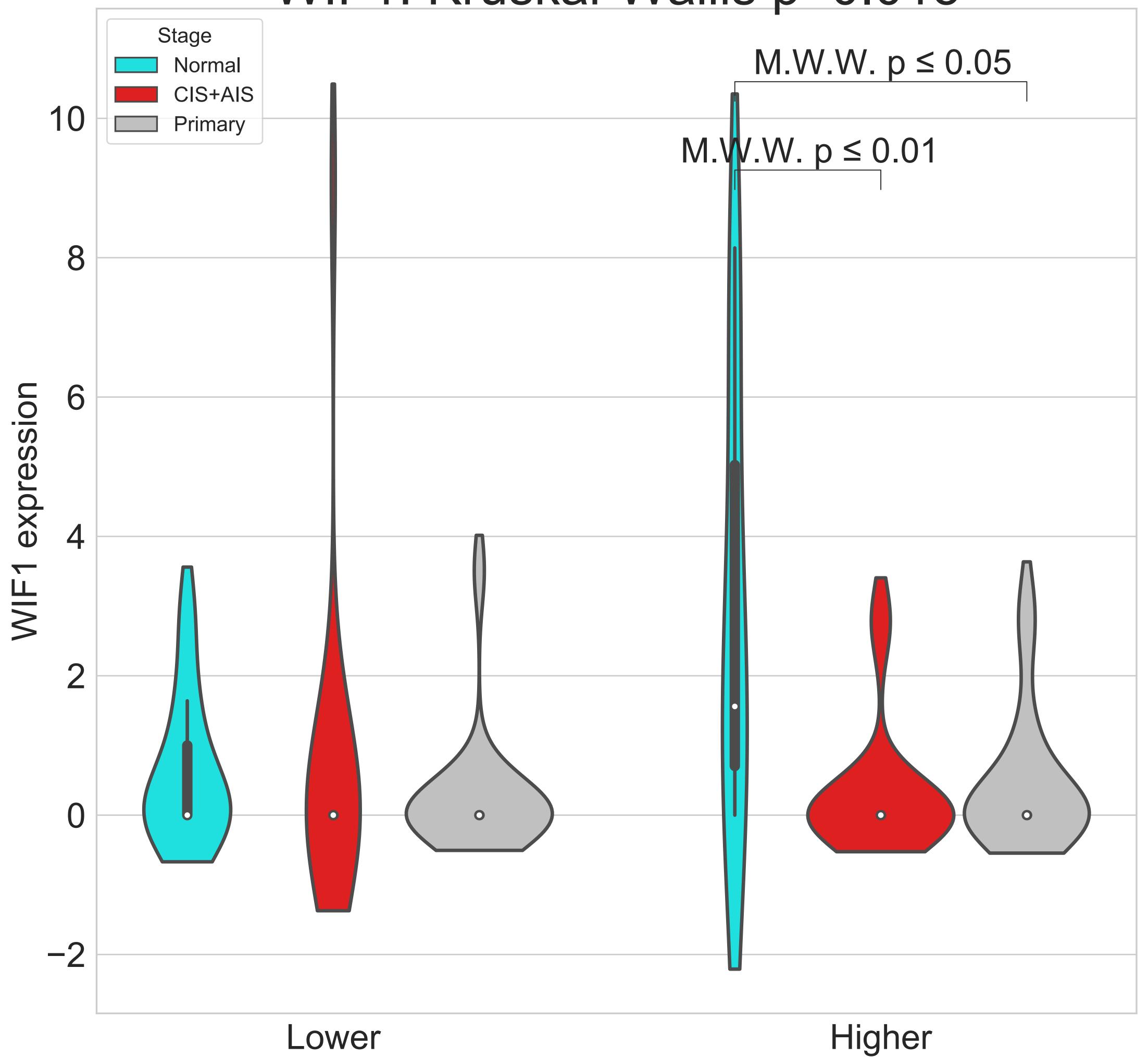
WIF1: Kruskal-Wallis p=0.015



Mutation Shared Count (SYN)