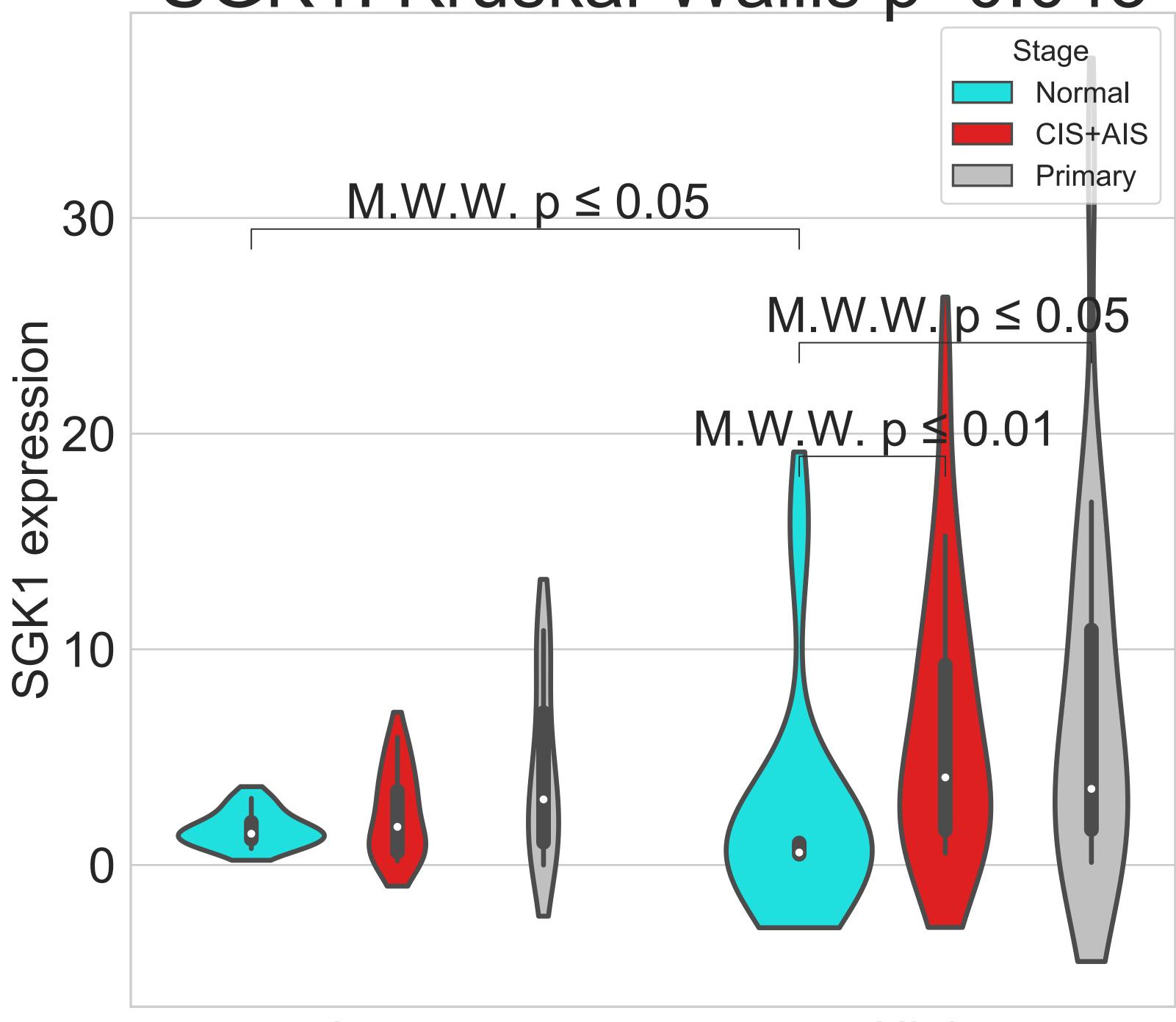
SGK1: Kruskal-Wallis p=0.043



Lower Higher Mutation Shared Proportion (SYN)