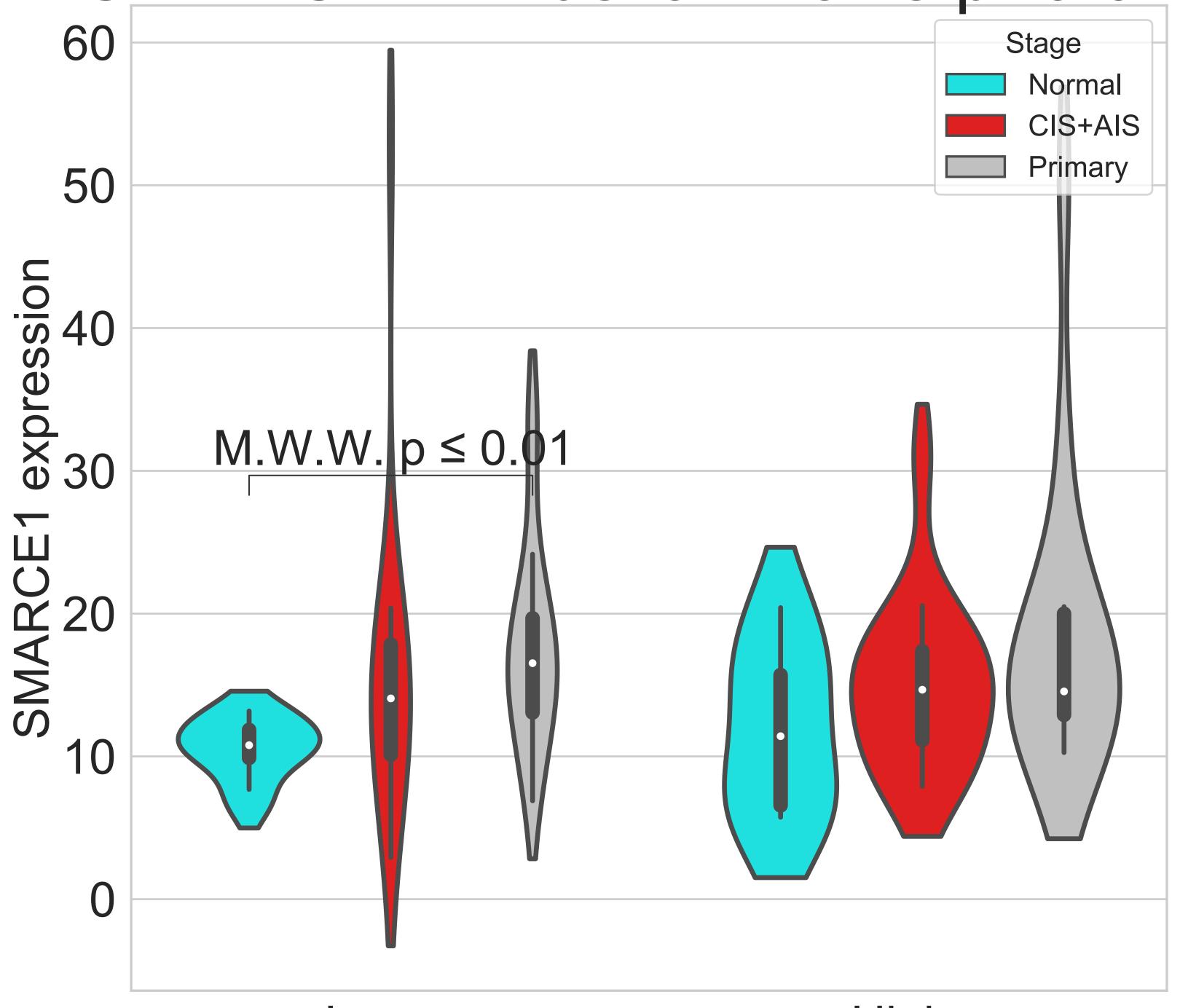
SMARCE1: Kruskal-Wallis p=0.026



Lower Higher Mutation Shared Proportion (SYN)