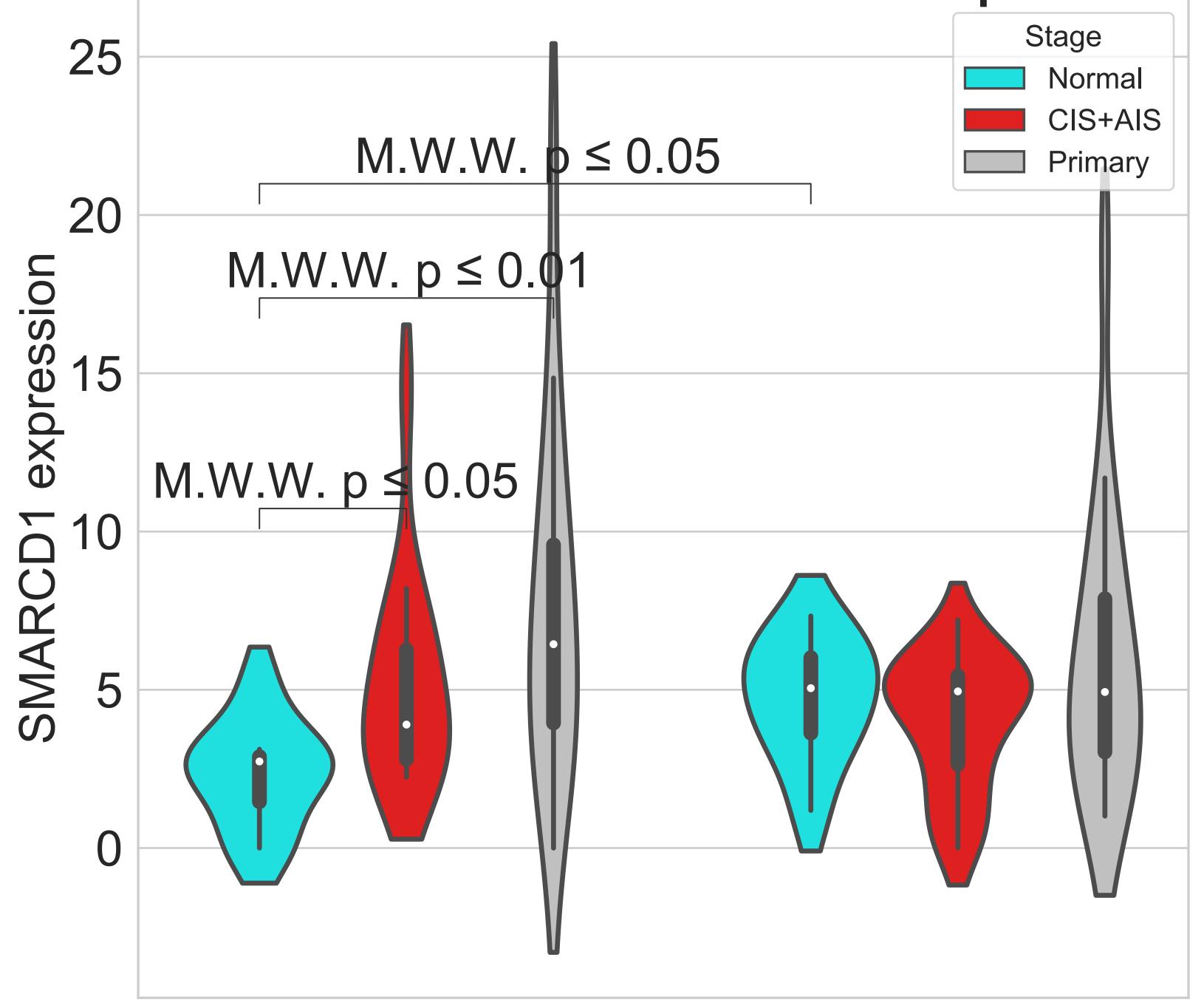
SMARCD1: Kruskal-Wallis p=0.047



Lower Higher Mutation Shared Proportion (Union & SYN)