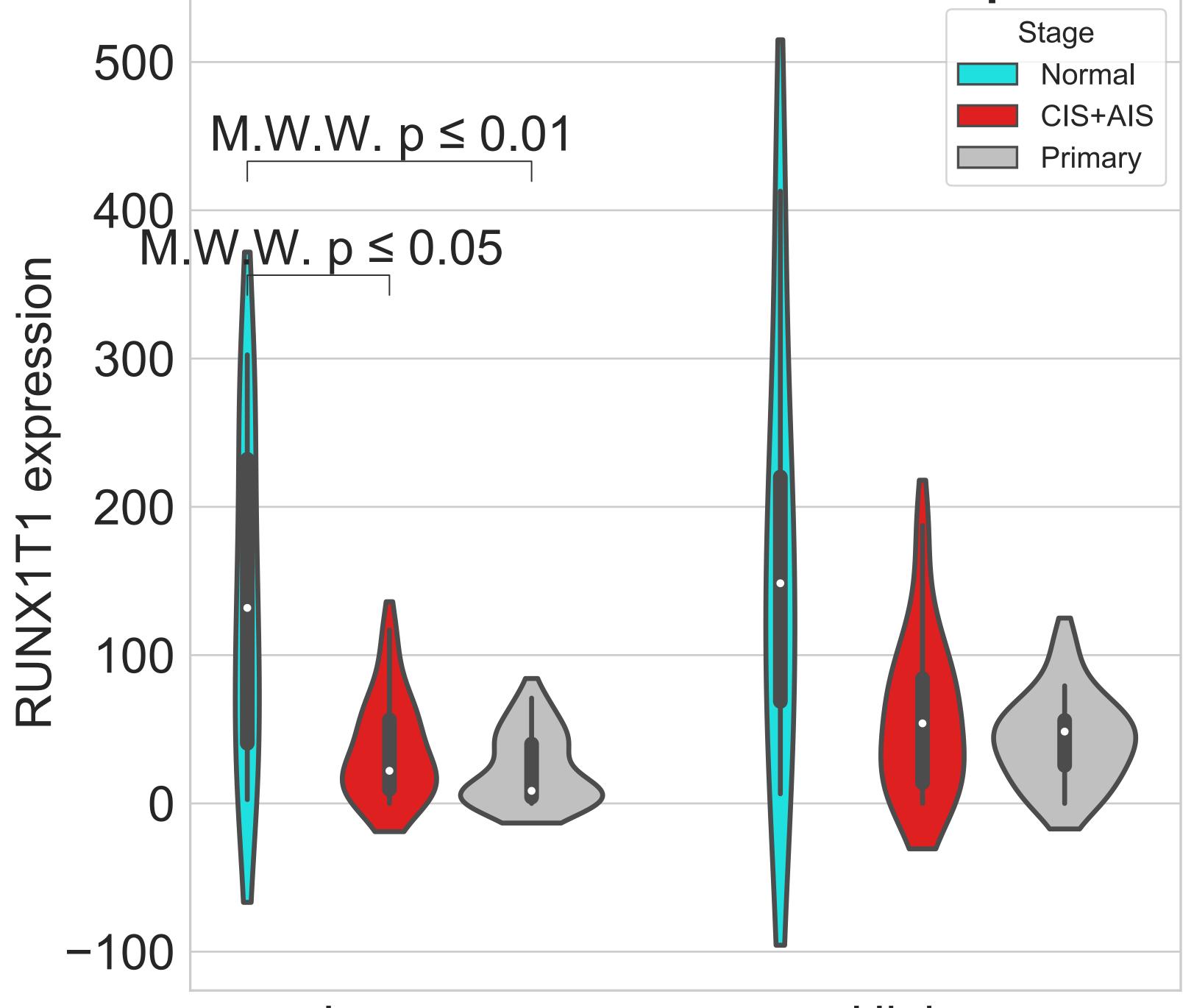
RUNX1T1: Kruskal-Wallis p=0.003



Lower Higher Mutation Shared Proportion (Union & SYN)