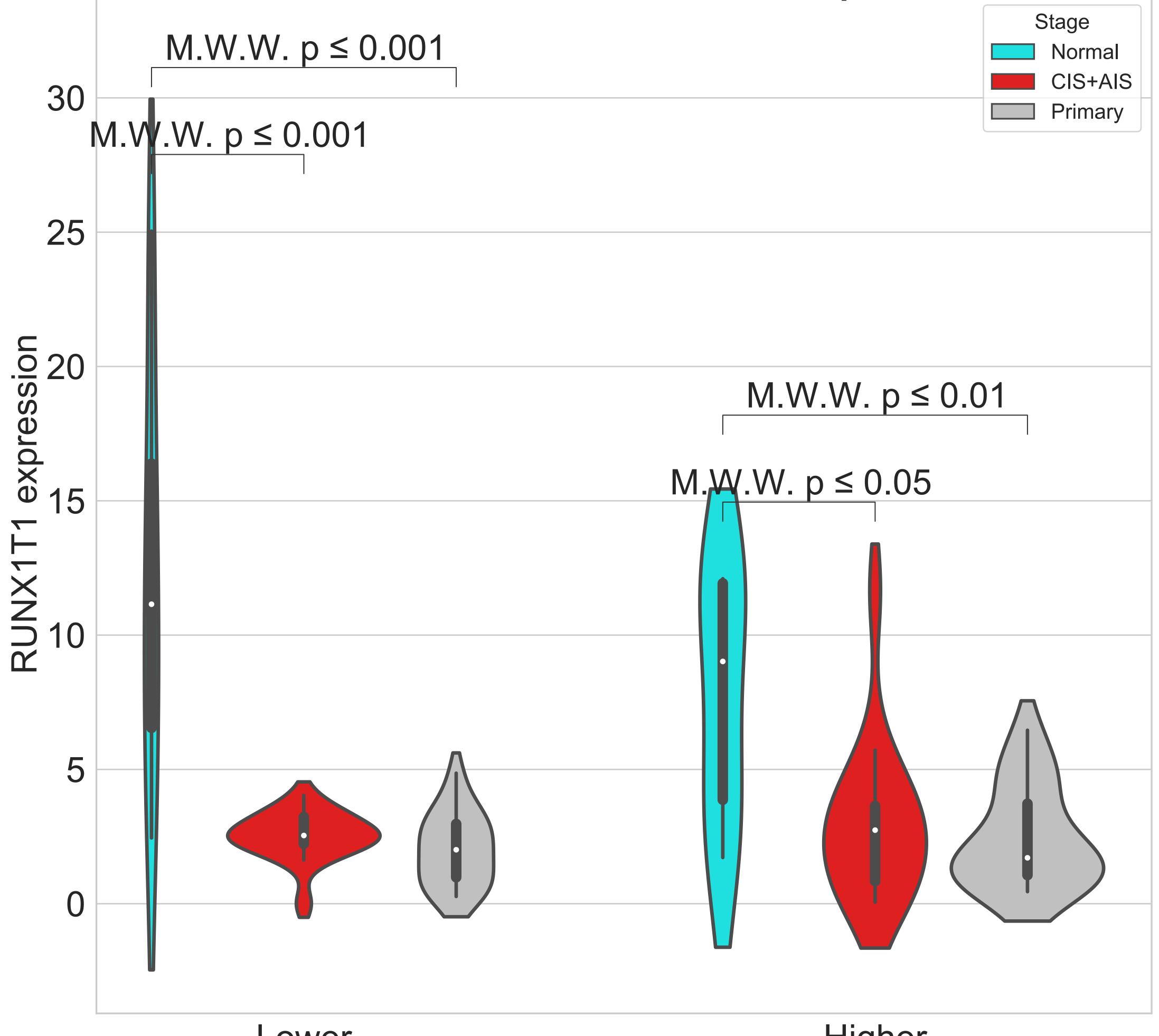
RUNX1T1: Kruskal-Wallis p=0.000



Higher Lower Mutation Shared Proportion (Union & SYN)