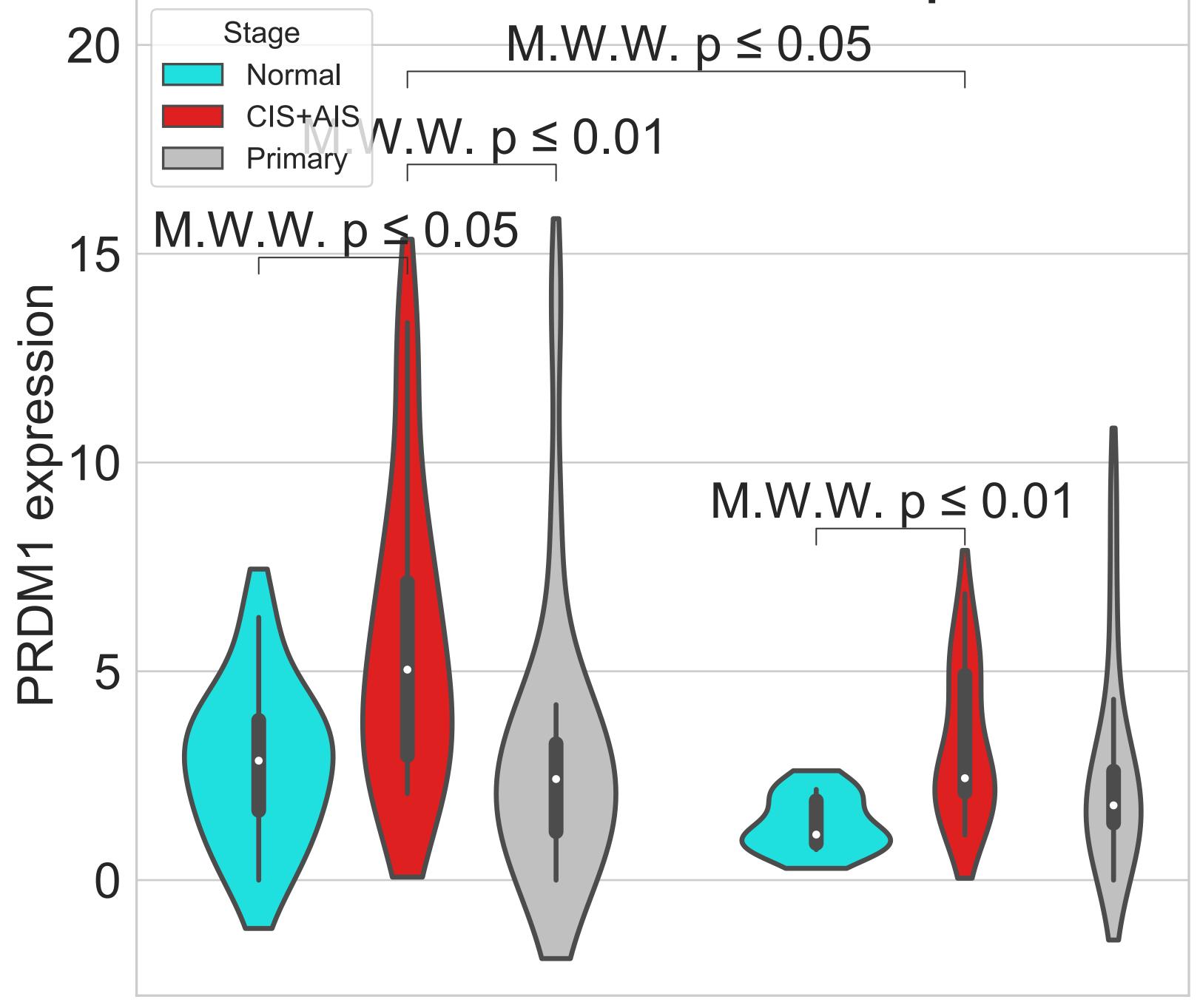
PRDM1: Kruskal-Wallis p=0.001



Lower
Mutation Shared Proportion (Union & SYN)