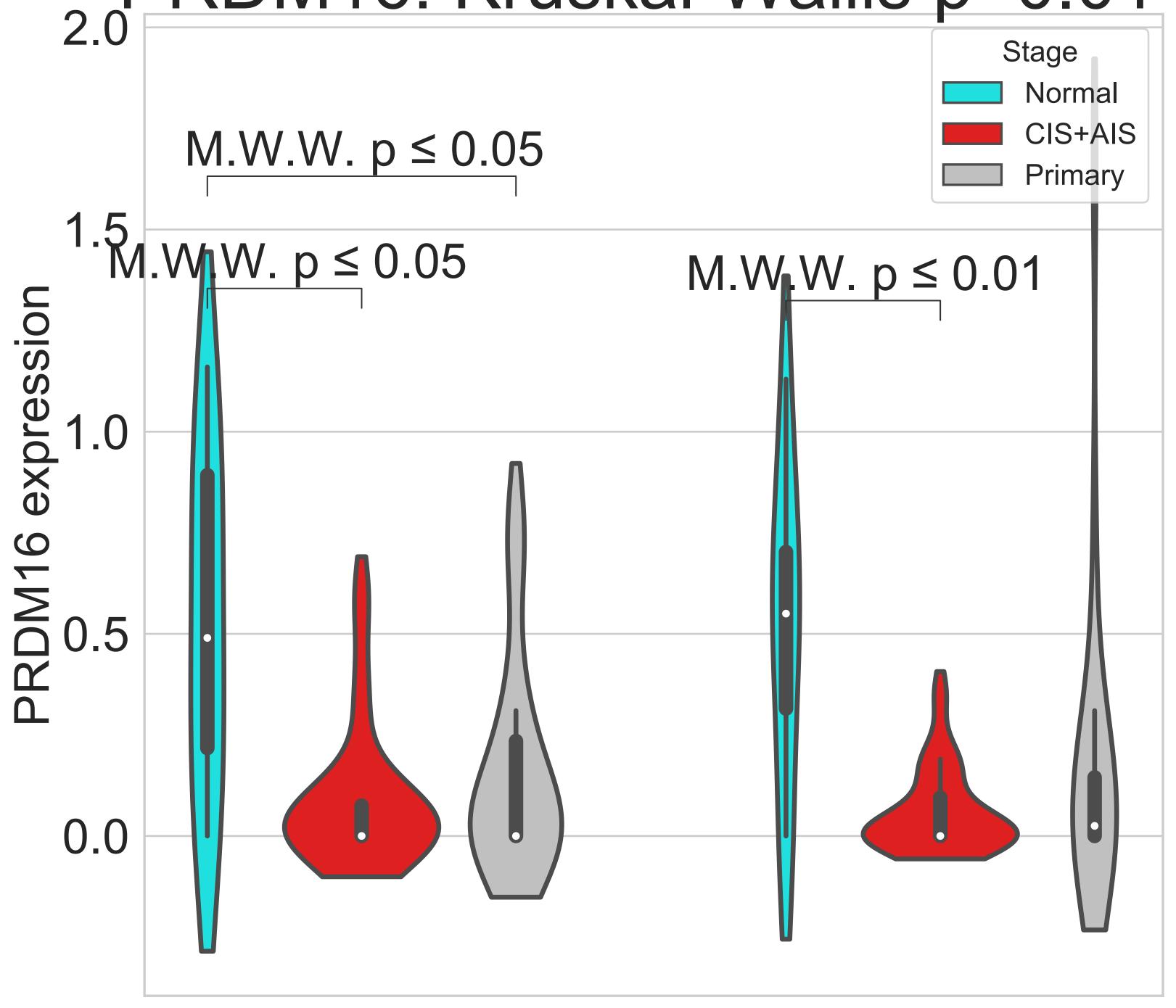
PRDM16: Kruskal-Wallis p=0.011



Lower
Mutation Shared Proportion (Union & SYN)