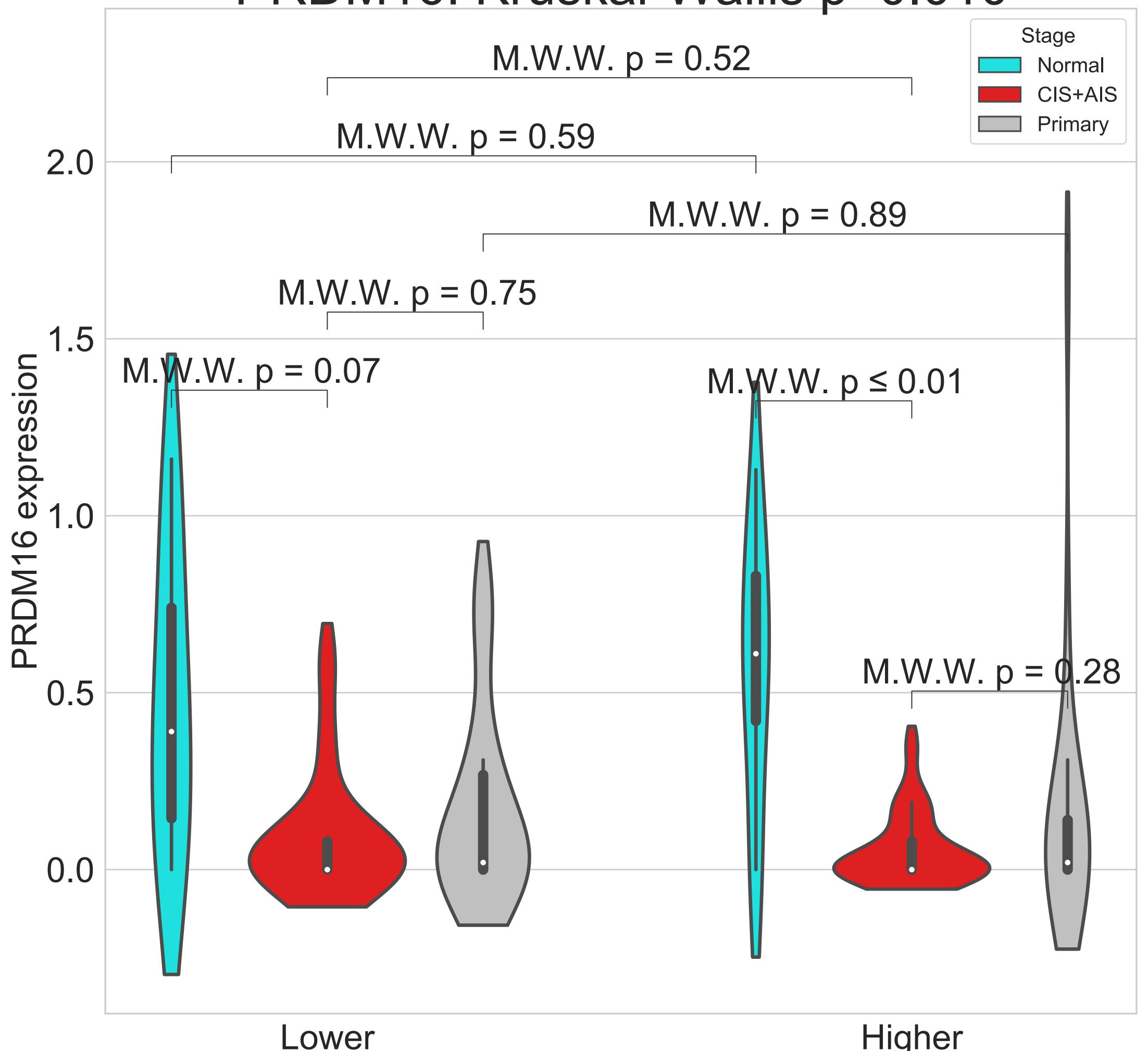
PRDM16: Kruskal-Wallis p=0.010



wer
Mutation Shared Proportion (SYN)