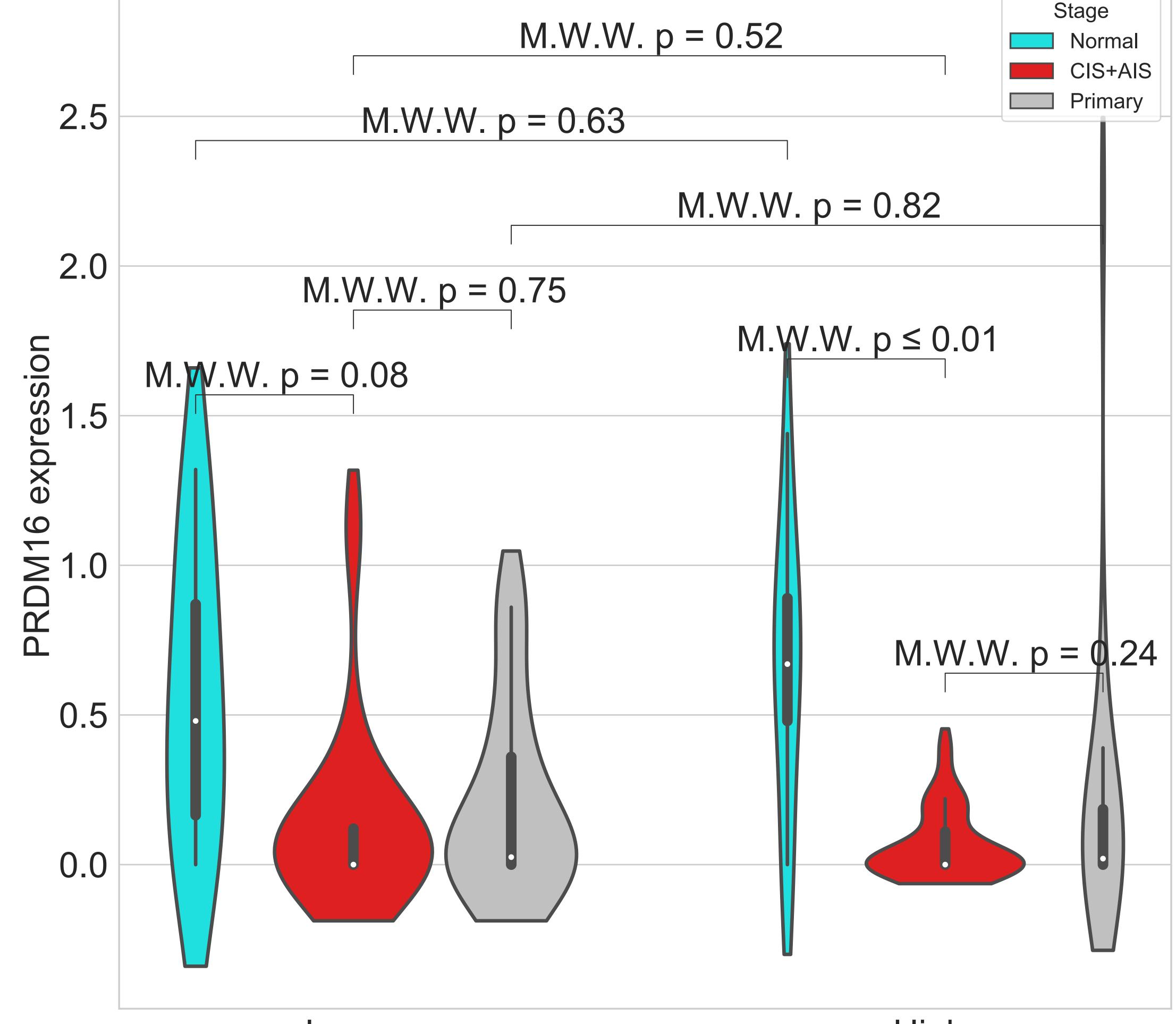
PRDM16: Kruskal-Wallis p=0.011 M.W.W. p = 0.52



Higher Lower Mutation Shared Proportion (SYN)