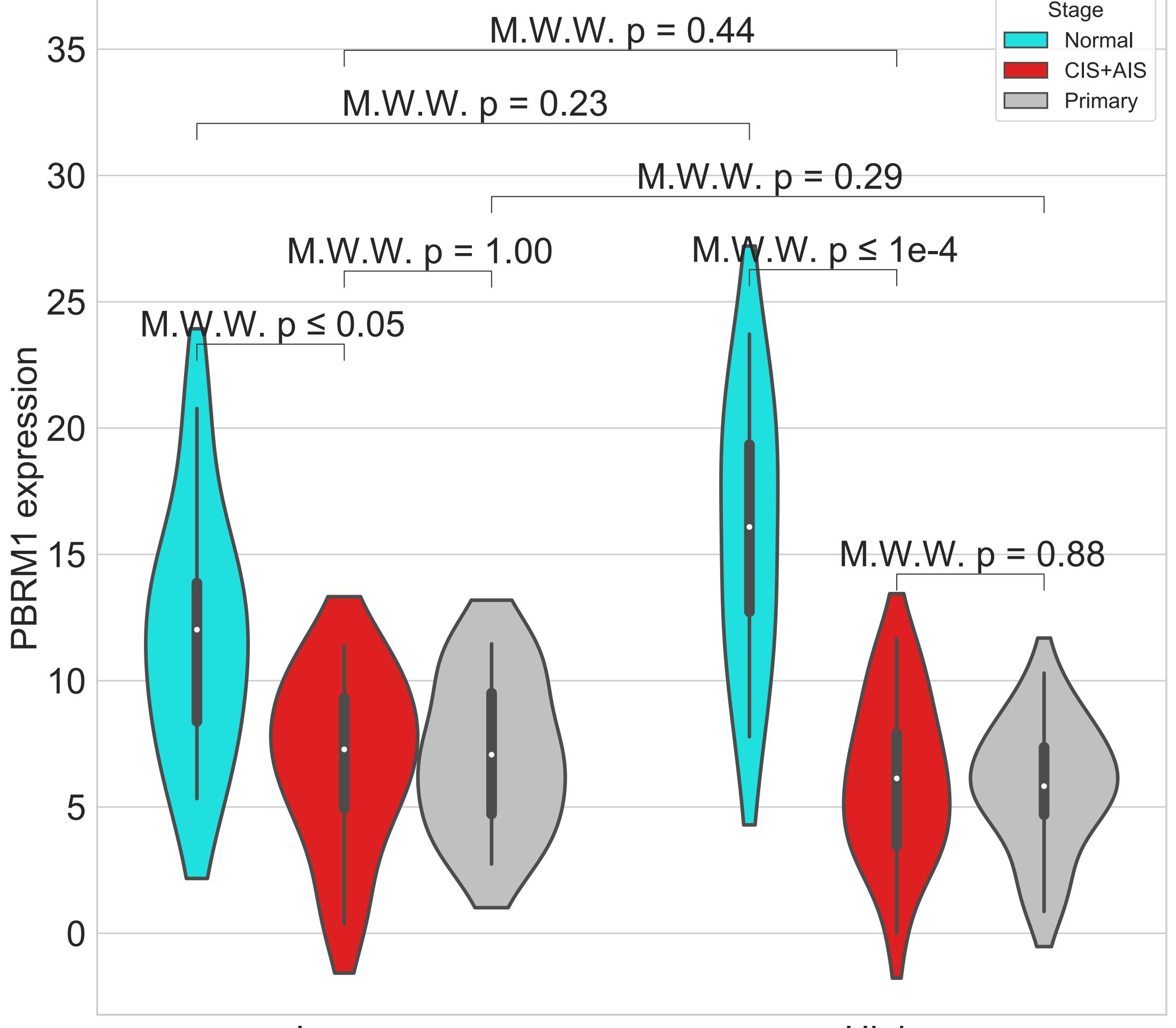
PBRM1: Kruskal-Wallis p=0.000 M.W.W. p = 0.44M.W.W. p = 0.23



Higher Lower Mutation Shared Count (SYN)