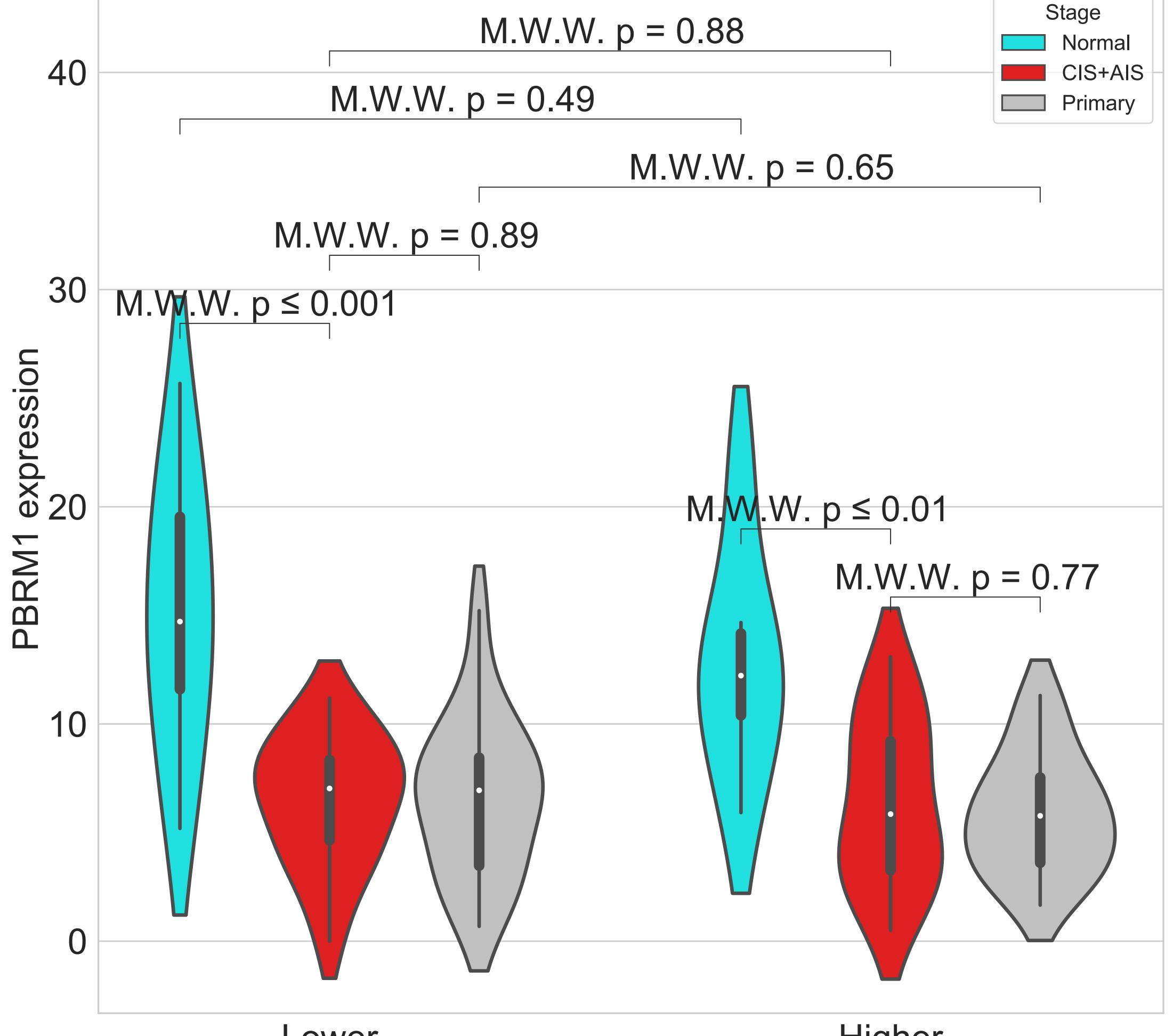
PBRM1: Kruskal-Wallis p=0.000 M.W. p = 0.88



Higher Lower Mutation Shared Proportion (SYN)