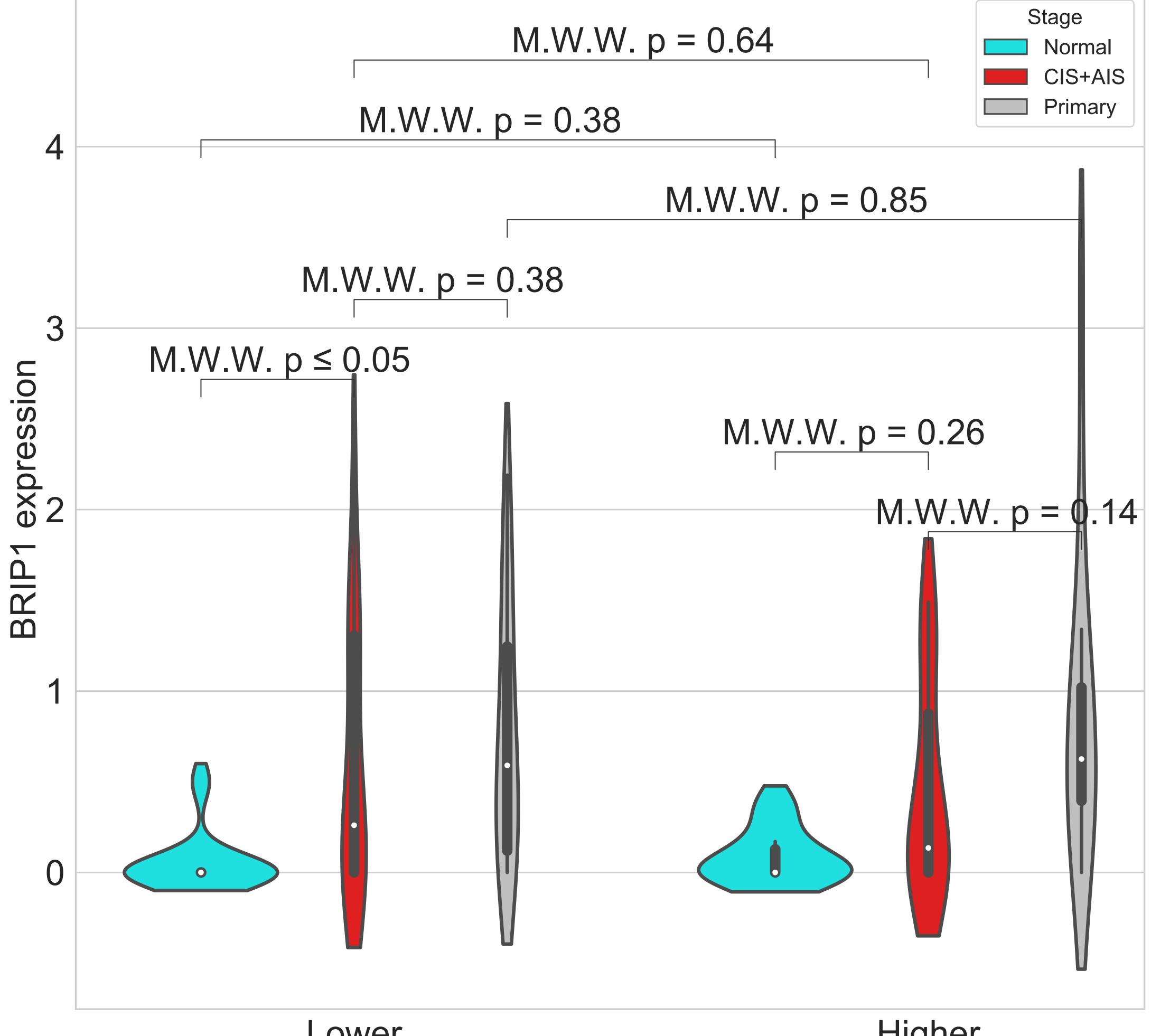
BRIP1: Kruskal-Wallis p=0.004 M.W.W. p = 0.64



Higher Lower Mutation Shared Count per TMB