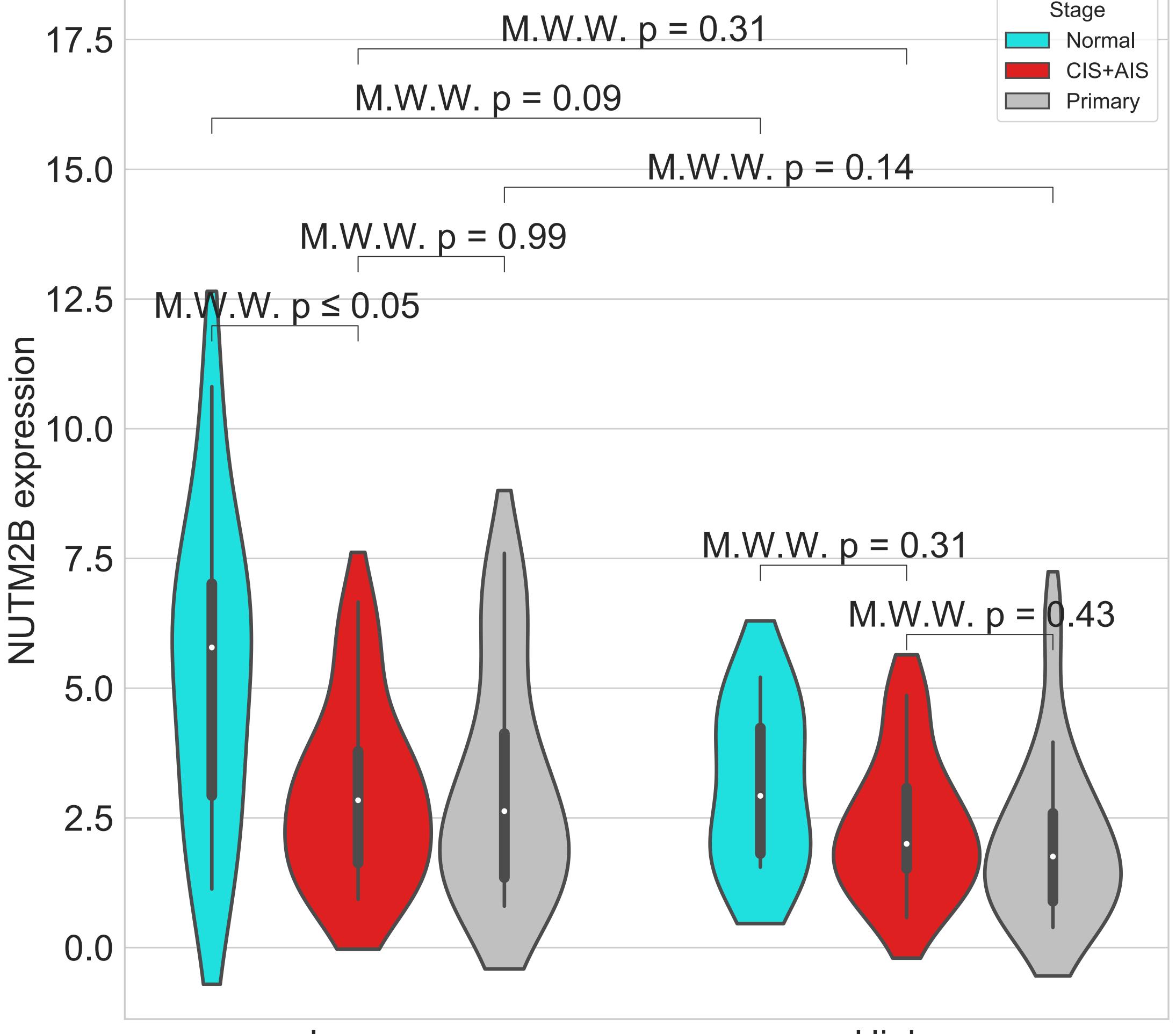
NUTM2B: Kruskal-Wallis p=0.034 M.W.W. p = 0.31



Higher Lower Mutation Shared Count per TMB