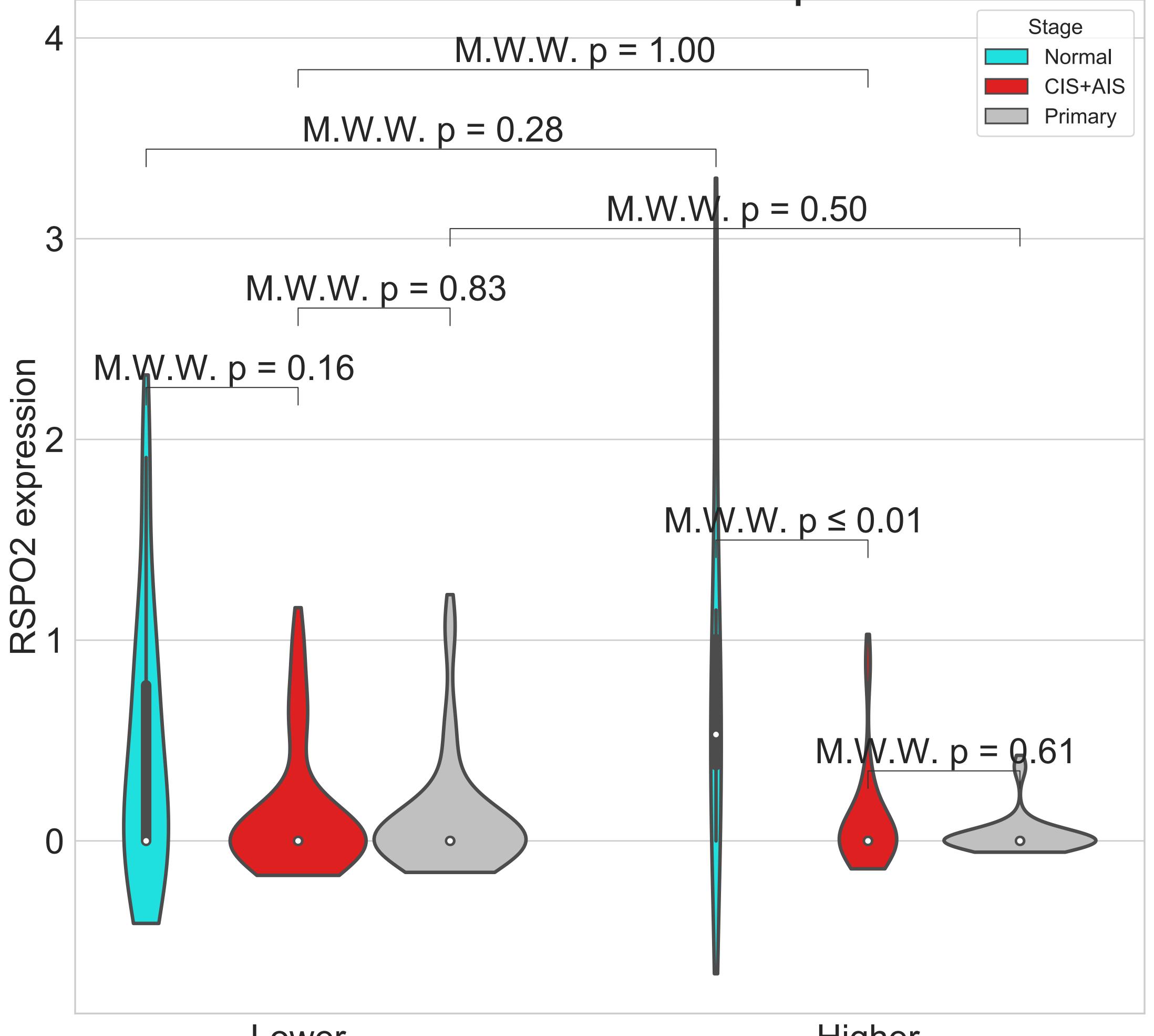
RSP02: Kruskal-Wallis p=0.007 M.W.W. p = 1.00



Higher Lower Mutation Shared Count (SYN) per TMB