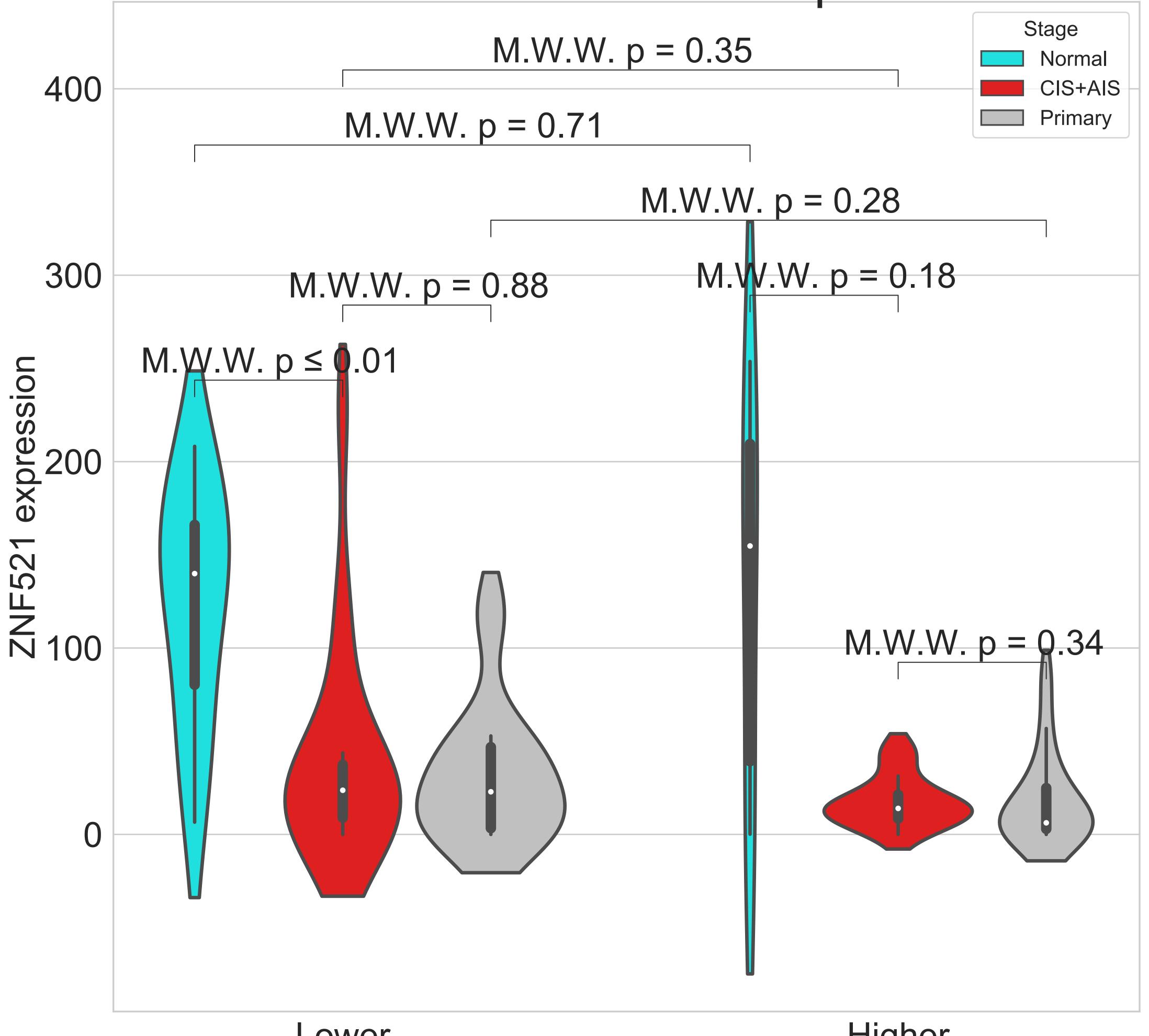
ZNF521: Kruskal-Wallis p=0.001 M.W.W. p = 0.35



Higher Lower Mutation Shared Count (SYN) per TMB