RECQL4: Kruskal-Wallis p=0.000 008 Stage M.W.W. p = 0.16Normal CIS+AIS Primary M.W.W. p = 0.84M.W.W. p = 0.41600 M.W.W. $p \le 0.001$ L4 expression M.W.W. p = 0.08M.W.W. p \leq 0.05 200 M.W.W. $p \le 0.001$ Higher Lower Mutation Shared Count (SYN) per TMB