RECQL4: Kruskal-Wallis p=0.000 800 Stage M.W.W. p = 0.06Normal CIS+AIS Primary M.W.W. p = 0.84M.W.W. p = 0.18600 M.W.W. $p \le 0.001$ -4 expression M.W.W. p = 0.07M.W.W. p \leq 0.05 200 M.W.W. $p \le 0.001$ Higher Lower Mutation Shared Count per TMB