HRAS: Kruskal-Wallis p=0.000 Stage M.W.W. p = 0.25Normal CIS+AIS 400 M.W.W. p = 0.67Primary M.W.W. p = 0.25M.W.W. $p \le 0.01$ 300 M.W.W. p = 0.13expression 5 0 M.W. p = 0.48M.W.W. $p \leq 0.05$ 100 Higher Lower Mutation Shared Count (SYN) per TMB