HRAS: Kruskal-Wallis p=0.000 Stage M.W.W. p = 0.17Normal CIS+AIS 500 M.W.W. p = 0.07**Primary** M.W.W. p = 0.11M.W.W.  $p \le 0.05$ 400 M.W.W. p = 0.16M.W.W. p = 0.40M.W.W. p  $\leq$  0.01 100 Higher Lower Mutation Shared Count (SYN) per TMB