HMGA2: Kruskal-Wallis p=0.000 400 Stage M.W.W. p = 0.91Normal CIS+AIS Primary 350 M.W.W. p = 0.22M.W.W. p = 0.73300 M.W.W. p ≤ 0.01 M.W.W. p = 0.20M.W.W. p = 0.18M.W.W. $p \le 1e-4$ 150 100 50 Higher Lower Mutation Shared Count (SYN) per TMB