HRAS: Kruskal-Wallis p=0.000 Stage M.W.W. p = 0.19Normal CIS+AIS 400 M.W.W. p = 0.23Primary M.W.W. p = 0.28M.W.W. p ≤ 0.01 300 M.W.W. p = 0.19expression 00 00 M.W.W. p = 0.32M.W.W. p ≤ 0.01 100 Higher Lower Mutation Shared Count per TMB