ARHGAP26: Kruskal-Wallis p=0.008 Stage Normal CIS+AIS 500 Primary M.W.W. p = 0.63M.W. p = 0.06400 M.W.V p = 0.11ARHGAP26 expression 300 M.W. p = 0.08M.W.W. p = 0.12M.W. p = 0.87M.W.W.  $p \le 0.05$ 200 100 -100Higher Lower Mutation Shared Count per TMB