ARHGAP5: Kruskal-Wallis p=0.028 Stage 300 M.W.W. p = 0.12Normal CIS+AIS **Primary** M.W.W. $p \le 0.01$ 250 M.W.W. p = 0.62M.W.W. $p \le 0.05$ expression 500 M.W.W. p = 0.90M.W.W. p = 0.72M.W.W. p = 0.95100 50 Higher Lower Mutation Shared Count per TMB