RAP1GDS1: Kruskal-Wallis p=0.014 Stage M.W.W. p = 0.05Normal 20 CIS+AIS Primary M.W.W. p = 0.47M.W.W. p = 0.70M.W.W. p = 0.1815 S1 expression M.W. p = 1.00M.W.W.  $p \le 0.05$ M.W.W. p = 0.30RAP1GD Higher Lower Mutation Shared Count (SYN) per TMB