RAP1GDS1: Kruskal-Wallis p=0.005 Stage M.W.W. p = 0.12Normal CIS+AIS 15.0 Primary M.W.W. p = 0.60M.W.W. p = 0.8712.5 M.W.W. p = 0.23S1 expression 10.0 M.W. p = 0.88M.W.W. p ≤ 0.05 M.W.W. p = 0.297.5 5.0 2.5 0.0 Higher Lower Mutation Shared Count (SYN) per TMB