RAP1GDS1: Kruskal-Wallis p=0.010 Stage M.W.W. p = 0.28Normal CIS+AIS 15.0 Primary M.W.W. p = 0.60M.W.W. p = 0.9012.5 M.W.W. p = 0.25S1 expression 10.0 M.W. p = 0.69M.W.W. p ≤ 0.05 M.W.W. p = 0.277.5 5.0 2.5 0.0 Higher Lower Mutation Shared Count per TMB