RAP1GDS1: Kruskal-Wallis p=0.015 Stage M.W. p = 0.41Normal CIS+AIS Primary M.W.W. p = 1.00M.W.W. p = 0.68M.W.W. p = 0.17M.W. p = 0.24M.W.W. p = 0.09M.W.W. p = 0.29

15.0

12.5

RAP1GD

5.0

2.5

0.0

Lower
Mutation Shared Count (SYN) per TMB